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XANTHONES FROM GUTTIFERAE

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Key Word Index—Guttiferae, xanthonenes; benzophenones, chemotaxonomy; biosynthesis.

Abstract—Since the last review in 1980, over eighty new xanthonenes have been isolated from the Guttiferae. These are listed with reference to structure elucidation and synthesis. The distribution of xanthonenes is examined in relation to the taxonomic divisions of the Guttiferae. Xanthone biosynthesis is discussed in the light of new biosynthetic results and the various pharmacological properties of xanthonenes are summarized.

INTRODUCTION

As the quest for new natural products continues, it becomes increasingly clear that xanthonenes are very restricted in occurrence. The majority of natural xanthonenes have been found in just two families of higher plants—Guttiferae and Gentianaceae [1]. Simple, oxygenated xanthonenes occur in both families and are generally more highly oxygenated in the Gentianaceae. Prenylated xanthonenes are widely distributed in the Guttiferae but not known in the Gentianaceae, and whereas *O*-glycosylxanthonenes are common in the Gentianaceae [2], only two have been reported from the Guttiferae.

Xanthonenes occur sporadically throughout the remainder of the plant kingdom. The Moraceae contain several Guttiferae-type prenylated xanthonenes and the Polygalaceae, simple hydroxy- and alkoxyxanthonenes [1]. *C*-Glucosylxanthonenes have been found in certain ferns, and in over one hundred species of higher plants [3], and fungi produce xanthonenes with substitution patterns characteristic of their acetate derivation [4].

Several earlier reviews have summarized the literature on xanthonenes [5–10], with emphasis on biosynthesis [7, 8], synthesis [6, 9] or phylogeny [10]. In the last review in 1980, Sultanbawa listed 95 xanthonenes from the Guttiferae [1]. Since then there has been a steady stream of reports in which more than 80 new xanthonenes have been characterized and many known xanthonenes re-isolated from *ca* 60 species of Guttiferae.

The aim of this review is to summarize the recent work on xanthonenes from the Guttiferae as a supplement to the 1980 review. The combined data will then be studied from a chemotaxonomic point of view to determine if any patterns of xanthone distribution exist within the family, and the implications of new biosynthetic results will be discussed. Structure elucidation, synthesis, and pharmacology will also be covered.

DISTRIBUTION

The family Guttiferae numbers over 1000 species, mainly confined to the tropics—the major exception being the genus *Hypericum*, which occurs widely in

temperate regions. According to Engler's Syllabus [11] the family comprises six subfamilies (Table 1). The subfamily Hypericoideae has been treated by some taxonomists (notably Hutchinson [12]) as a separate family, but the recent surge of interest in the chemistry of this group has led to the isolation of several prenylated xanthonenes, supporting its inclusion in the Guttiferae. The only related family in which xanthonenes have been found is the Bonnetiaceae, which, in keeping with Thorn's recent classification [13], is included in the Kielmeyeroideae subfamily in Table 1.

Xanthonenes or the related benzophenones have been found in all the major and several minor genera of the Guttiferae. The approximate number of species [14], and the number that have been found to contain xanthonenes is given for each genus (Table 1). It appears from the number of chemically investigated species (for three of the larger genera), that a large proportion of the species contains xanthonenes. A species–xanthone index is included as an appendix.

Simple oxygenated xanthonenes

The symmetrical nature of the xanthone nucleus, coupled with its mixed biogenetic origin in higher plants, necessitates that the carbons be numbered according to a biosynthetic convention. Carbons 1–4 are assigned to the acetate-derived ring A, (often characterized by 1,3-dioxygenation) and carbons 5–8 to the shikimate-derived ring B (e.g. 1,3,7,8-tetrahydroxy rather than 1,2,6,8). All higher plant xanthonenes appear to have 5- and/or 7-oxygenation [8] and with this assumption, only xanthonenes with 2,5(or 4,7)-oxygenation have alternative names. In cases where only ring B is oxygenated the lowest numbers are used except in the biosynthetic discussion [e.g. 2- rather than 7-hydroxyxanthone (1)].

Structural assignments are based mainly on ¹H NMR and UV spectroscopy. Shifts of ring or side-chain protons due to acetylation or alkylation of adjacent hydroxyls and changes in the UV spectrum on addition of the usual shift reagents, are especially useful in determining substitution patterns. Confirmation of assignment may be

Table 1 The family Guttiferae

<u>SUB-FAMILY</u>	<u>TRIBE</u>	<u>GENUS</u>	<u>NO. OF SPECIES*</u>
Kielmeyeroideae	Kielmeyereae	<i>Kielmeyera</i>	20/9(10)
		<i>Mahurea</i>	8/1
	Caraipeae (Bonnetiaceae)	<i>Caraipta</i>	20/4
		<i>Haploclathra</i>	4/3
		<i>Bonnetia</i>	18/1
		<i>Archytaea</i>	2/1
Calophylloideae	Calophylleae	<i>Calophyllum</i>	110/21(32)
		<i>Mesua</i> (inc. <i>Kayea</i>)	40/5
		<i>Mammea</i> (inc. <i>Ochrocarpus</i>)	50/5
Clusioidae	Clusiaceae	<i>Clusia</i>	145/4
		<i>Toumita</i>	60/6
	Garcinieae	<i>Allanblackia</i>	8/1
		<i>Garcinia</i>	400/29(38)
		<i>Pentaphalangium</i>	7/1
		<i>Rhedia</i>	45/4
Moronoboideae		<i>Moronobea</i>	7/1
		<i>Pentadesma</i>	4/1
		<i>Platonia</i>	1-2/1
		<i>Symphonia</i>	20/1
Lorostemonoideae		<i>Lorostemon</i>	3/2
Hypericoideae	Cratoxyleae	<i>Cratoxylum</i>	6/2
	Hypericeae	<i>Hypericum</i>	400/18
	Vismieae	<i>Harungana</i>	1/1
		<i>Psorospermum</i>	40/1
		<i>Vismia</i>	35/3

*Approx. no. of species [14]/no. of species in which xanthenes and closely related compounds have been found. Numbers in parentheses refer to number of species chemically investigated.

obtained by derivatization (sometimes not possible due to the scarcity of material), although often synthesis is necessary.

The synthesis of xanthenes has been reviewed [15, 9]. The standard method is the Grover, Shah and Shah condensation between an *ortho*-oxygenated benzoic acid and a reactive phenol in the presence of phosphorous oxychloride and zinc chloride [16]. A recent variation of this method uses methylsulphonic acid and phosphorous pentoxide in place of the above reagents [17]. Alternatively, benzophenones, prepared by Friedel-Crafts acylation [18], may be converted to xanthenes by dehydrative or oxidative cyclization. Two other methods have appeared recently. The nucleophilic addition of salicylic acid derivatives to *p*-benzoquinones has been used to prepare a series of 1,4 (or 5,8)-dihydroxyxanthenes [19], and 3-hydroxyxanthone has been prepared in 50% yield

by a cycloaddition reaction between a diene and a benzopyranone [20].

Mono-oxygenated xanthenes. Two mono-oxygenated xanthenes (**1** and **2**, Table 2) have recently been isolated from seven species. 4-Hydroxyxanthone also occurs in Guttiferae. These compounds are now known to occur in eight genera from three subfamilies. Notable is their absence from Clusioidae (> 50 species investigated).

Dioxygenated xanthenes. The majority of the recently isolated dioxygenated xanthenes (Table 3) are from species of the Hypericoideae, reflecting the recent interest in this subfamily. Dioxygenated xanthenes are however, also common in species of *Calophyllum*, *Mammea* and *Mesua* and found in all subfamilies. 1,7-Dihydroxyxanthone (**4**), has now been isolated from 40 species of Guttiferae. The 2,5-dioxygenation pattern of xanthenes **5** and **10** is new from nature.

Table 2 Mono-oxygenated xanthenes

<u>2-Hydroxyxanthone(1)</u>	<u>2-Methoxyxanthone(2)</u>
<i>Calophyllum zeylanicum</i> Kosterm.[21]	<i>Caraipa psidifolia</i> Ducke [29]
<i>Hypericum balearicum</i> L. [22]	<i>Hypericum mysorensense</i> [27]
<i>Hypericum canariensis</i> L. [24]	<i>Vismia guaramirangae</i> Huber [28]
<i>Hypericum ericoides</i> L. [25]	
<i>Hypericum mysorensense</i> [26,27]	
<i>Vismia guaramirangae</i> Huber [28]	

*First reported occurrence in nature

†First reported occurrence in Guttiferae.

Table 3 Dioxygenated xanthenes

<u>1,5-Dihydroxyxanthone(3)</u>	<u>1-Hydroxy-7-methoxyxanthone(7)</u>
<i>Calophyllum zeylanicum</i> Kosterm.[21]	<i>Bonnetia stricta</i> (Nees) Ness Mart. [31]
<i>Garcinia xanthochymus</i> Hook. f. [30]	<i>Haploclathra paniculata</i> (Mart.) Benth. [33]
<u>1,7-Dihydroxyxanthone(4)(Euxanthone)</u>	<i>Haploclathra verticillata</i> Ducke[29]
<i>Bonnetia stricta</i> (Nees) Ness & Mart.[31]	<i>Hypericum mysorensense</i> [26]
<i>Calophyllum zeylanicum</i> Kosterm.[21]	<i>Mahurea tomentosa</i> Ducke [29]
<i>Garcinia indica</i> Choisy [50]	<i>Vismia guaramirangae</i> Huber [28]
<i>Garcinia xanthochymus</i> Hook. f.[30]	<u>2-Hydroxy-1-methoxyxanthone(8).</u>
<i>Haploclathra verticillata</i> Ducke[29]	<i>Vismia guaramirangae</i> Huber [28]
<i>Hypericum balearicum</i> L. [22]	<u>2-Hydroxy-3-methoxyxanthone(9)</u>
<i>Hypericum canariensis</i> L. [24]	* <i>Hypericum mysorensense</i> [26]
<i>Hypericum ericoides</i> L. [25]	<u>2-Hydroxy-5-methoxyxanthone(10)</u>
<i>Hypericum mysorensense</i> [26,27]]	* <i>Hypericum androsaemum</i> L. [34]
<i>Mahurea tomentosa</i> Ducke [29]	<i>Hypericum canariensis</i> L. [24]
<i>Vismia guaramirangae</i> Huber [28]	<u>3-Hydroxy-2-methoxyxanthone(11)</u>
<u>2,5-Dihydroxyxanthone(5)</u>	<i>Hypericum androsaemum</i> L. [34]
* <i>Hypericum canariensis</i> L. [24]	<i>Hypericum balearicum</i> L. [22]
<u>2,3-Dimethoxyxanthone(6)</u>	<i>Psorospermum febrifugum</i> Sprach [35]
* <i>Hypericum mysorensense</i> [27,32]	<i>Vismia guaramirangae</i> Huber [28]

Trioxxygenated xanthenes. Twenty trioxxygenated xanthenes, of which nine are new natural products, are shown in Table 4. Novel oxygenation patterns are 1,5,8-

(17), 1,4,7- (19) and 2,3,5- (27), and xanthenes 33 and 34 are the first trimethoxyxanthenes from the Guttiferae. Two 1,2,5-trioxxygenated xanthenes (12 and 14) have been

Table 4 Trioxxygenated xanthenes

<u>1,2-Dihydroxy-5-methoxyxanthone(12)</u>	<u>1-Hydroxy-3,5-dimethoxyxanthone(23)</u>
* <i>Garcinia xanthochymus</i> Hook f [30]	<i>Haploclathra paniculata</i> (Mart) Benth [33]
<u>1,3-Dihydroxy-2-methoxyxanthone(13)</u>	<u>1-Hydroxy-6,7-dimethoxyxanthone(24)</u>
* <i>Vismia guaramirangae</i> Huber [28]	<i>Hypericum mysorense</i> [26]
<u>1,5-Dihydroxy-2-methoxyxanthone(14)</u>	<u>1-Hydroxy-7,8-dimethoxyxanthone(25)</u>
* <i>Garcinia xanthochymus</i> Hook. f [30]	<i>Haploclathra paniculata</i> (Mart) Benth [33]
<u>1,5-Dihydroxy-3-methoxyxanthone(15)</u>	<u>2-Hydroxy-3,4-dimethoxyxanthone(26)</u>
<i>Garcinia xanthochymus</i> Hook f.[30]	<i>Hypericum canariensis</i> L [24]
<i>Haploclathra leantha</i> (Benth) Benth [36]	<i>Hypericum sampsonii</i> Hance [39]
<i>Haploclathra paniculata</i> (Mart) Benth [33]	<i>Kielmeyera rubiflora</i> Camb [40]
<i>Vismia guaramirangae</i> Huber [28]	* <i>Kielmeyera speciosa</i> St.Hill [40]
<u>1,5-Dihydroxy-6-methoxyxanthone(16)</u>	<u>3-Hydroxy-2,5-dimethoxyxanthone(27)</u>
* <i>Tovomita excelsa</i> Andrade-Lima et G Mariz [37]	* <i>Hypericum androsaemum</i> L [34]
<u>1,5-Dihydroxy-8-methoxyxanthone(17)</u>	<u>4-Hydroxy-2,3-dimethoxyxanthone(28)</u>
* <i>Vismia guaramirangae</i> Huber [28]	<i>Caraipa grandiflora</i> Mart. [29]
<u>1,6-Dihydroxy-5-methoxyxanthone(18)</u> (<u>Buchanoxanthone</u>)	<u>5-Hydroxy-1,3-dimethoxyxanthone(29)</u>
<i>Calophyllum zeylanicum</i> Kosterm [21]	<i>Haploclathra paniculata</i> (Mart.) Benth. [33]
<i>Tovomita excelsa</i> Andrade-Lima et G Mariz [37]	<u>1,3,5-Trihydroxyxanthone(30)</u>
<u>1,7-Dihydroxy-4-methoxyxanthone(19)</u>	<i>Garcinia xanthochymus</i> Hook. f [30]
* <i>Vismia guaramirangae</i> Huber [28]	<u>1,3,7-Trihydroxyxanthone(31)</u> (<u>Gentisein</u>)
<u>1,7-Dihydroxy-6-methoxyxanthone(20)</u>	<i>Haploclathra paniculata</i> (Mart.) Benth [33]
* <i>Tovomita excelsa</i> Andrade-Lima et G Mariz [37]	<i>Hypericum degentii</i> Bornm. [41]
<u>1,7-Dihydroxy-8-methoxyxanthone(21)</u>	<u>1,6,7-Trihydroxyxanthone(32)</u>
<i>Bonnetia stricta</i> (Nees) Ness & Mart [31]	<i>Platonia insignis</i> Mart [29]
<i>Haploclathra leantha</i> (Benth) Benth. [36]	<u>1,3,5-Trimethoxyxanthone(33)</u>
<i>Haploclathra paniculata</i> (Mart) Benth. [33]	† <i>Haploclathra paniculata</i> (Mart.) Benth [33]
<u>5,6-Dihydroxy-1-methoxyxanthone(22)</u>	<u>2,3,4-Trimethoxyxanthone(34)</u>
* <i>Tovomita excelsa</i> Andrade-Lima et G. Mariz [37]	* <i>Hypericum ericoides</i> L [25]

reported from *Garcinia xanthochymus*, but their physical data have not been published.

Three of the four xanthenes isolated from the trunk

wood of *Tovomita excelsa*, were assigned a 1,5,6-dihydroxymethoxy structure [37]. Two of these (16, 18) were known compounds, whereas the 1-methoxy structure (22)

Table 5. Tetra- and pentaoxygenated xanthenes

<u>1,3-Dihydroxy-5,6-dimethoxyxanthone</u> <u>(Leiaxanthone) (35)</u>	<u>2-Hydroxy-5,6,7-trimethoxyxanthone</u> <u>(43)</u>
† <i>Haploclathra lieantha</i> (Benth) Benth [36]	* <i>Hypericum ericoides</i> L [25]
<u>1,5-Dihydroxy-6,7-dimethoxyxanthone</u> <u>(36)</u>	<u>3-Hydroxy-1,2,4-trimethoxyxanthone</u> <u>(44)</u>
<i>Caraipa grandiflora</i> Mart [29]	* <i>Psorospermum febrifugum</i> Sprach [35]
<i>Caraipa psidifolia</i> Ducke [29]	<u>3-Hydroxy-1,5,6-trimethoxyxanthone</u> <u>(45)</u>
<i>Caraipa valloi</i> Paula [29]	<i>Haploclathra lieantha</i> (Benth) Benth [36]
<i>Tovomita brasiliensis</i> Walp [42]	<u>7-Hydroxy-1,3,8-trimethoxyxanthone</u> <u>(Anthaxanthone) (46)</u>
<u>1,6-Dihydroxy-5,7-dimethoxyxanthone</u> <u>(37)</u>	* <i>Haploclathra lieantha</i> (Benth) Benth [36]
<i>Hypericum canariensis</i> L [24]	<u>1,3,5,6-Tetrahydroxyxanthone (47)</u>
<u>1,7-Dihydroxy-3,8-dimethoxyxanthone</u> <u>(Gentiaculein) (38)</u>	<i>Hypericum androsaemum</i> L [34]
† <i>Haploclathra lieantha</i> (Benth) Benth [36]	<u>1,3,6,7-Tetrahydroxyxanthone (48)</u> <u>(Norathyriol)</u>
<i>Haploclathra paniculata</i> (Mart) Benth. [33]	<i>Cratoxylum pruniflorum</i> Kurz [130]
<u>2,5-Dihydroxy-1,6-dimethoxyxanthone</u> <u>(39)</u>	<i>Garcinia mangostana</i> L. [46]
* <i>Garcinia thwaitesii</i> Pierre [43]	<i>Hypericum androsaemum</i> L. [34]
<u>3,7-Dihydroxy-1,8-dimethoxyxanthone</u> <u>(40)</u>	<i>Hypericum aucheri</i> Jaub et Sprach. [47]
* <i>Haploclathra paniculata</i> (Mart.) Benth [33]	<u>1,3,5-Trihydroxy-6-methoxyxanthone</u> <u>(49)</u>
<u>3,8-Dihydroxy-1,7-dimethoxyxanthone</u> <u>(Isogentiaculein) (41)</u>	<i>Haploclathra lieantha</i> (Benth) Benth. [36]
† <i>Haploclathra paniculata</i> (Mart.) Benth [33]	<u>1,5,6-Trihydroxy-3-methoxyxanthone</u> <u>(50)</u>
<u>5,6-Dihydroxy-1,3-dimethoxyxanthone</u> <u>(Ferraxanthone) (42)</u>	† <i>Hypericum androsaemum</i> L. [34]
<i>Haploclathra lieantha</i> (Benth) Benth. [36]	<u>1,7,8-Trihydroxy-6-methoxyxanthone</u> <u>(51)</u>
* <i>Nesua ferrea</i> L. [23]	* <i>Archytaea multiflora</i> Benth. [31]
	<u>2,4,5-Trihydroxy-1-methoxyxanthone</u> <u>(BR-xanthone-B) (52)</u>
	* <i>Garcinia mangostana</i> L. [83]
<u>3,8-Dihydroxy-1,2,4-trimethoxyxanthone (53)</u>	
* <i>Psorospermum febrifugum</i> Sprach [35]	

was new. An unambiguous synthesis of **22** has been completed by the methylation of 1-hydroxy-5,6-dibenzyloxyxanthone followed by debenzylation, and the product shown to be different from the natural compound [38]. 4,5-Dihydroxy-3-methoxyxanthone is suggested as a likely alternative structure

The sodium acetate induced shifts in the UV spectrum of a 2,3,4-hydroxydimethoxyxanthone from *Kielmeyera* species, led to the assignment of the more acidic 3-

hydroxy structure. However, the ^{13}C NMR spectrum has indicated that both methoxy groups are di-*ortho*-substituted and that therefore the structure should be 2-Hydroxy-3,4-dimethoxyxanthone (**26**) [40]. Consequently, the structure of a xanthone from *Hypericum canariensis* should also be revised to **26**, as in Table 4.

Ten xanthenes have been isolated from the roots of *Vismia quaramirangae* [28]. Based mainly on ^1H NMR spectroscopy, one of the three new xanthenes was as-

signed the structure of 1,7-dihydroxy-4-methoxy-xanthone (**19**), rather than the 2-methoxy alternative that was indicated by a positive Gibbs test. Two para-dioxygenated analogues also gave positive Gibbs tests, suggesting that this test is unreliable in certain cases

Tetraoxygenated xanthones The newly isolated tetraoxygenated xanthones are shown in Table 5. In the first investigations of the genus *Haploclathra*, three species have yielded 17 different xanthones [29, 33, 36]. These include four 1,3,5,6- and four 1,3,7,8-tetraoxygenated xanthones, the latter being common in the Gentianaceae. Xanthone **35** has been synthesized [44] and shown to be identical to a compound from *Centaurium linarifolium*, but to differ somewhat from the *Haploclathra* xanthone.

Several unusual oxygenation patterns have been found. The stem bark and timber of *Garcinia thwaitesi* yielded 2,5-dihydroxy-1,6-dimethoxyxanthone (**39**), and the structure of 2-hydroxy-5,6,7-trimethoxyxanthone (**43**), isolated from *Hypericum ericoides* [24], has been confirmed by synthesis (Scheme 1) [45]. A 1,2,4,5-tetraoxygenated xanthone, BR-xanthone-B (**52**), has been isolated from the fruit of *Garcinia mangostana*.

The inclusion of *Archytaea* (Bonnetiaceae) in the subfamily Kielmeyeroideae is supported by the isolation of a 1,6,7,8-tetraoxygenated xanthone (**51**) from *Archytaea multiflora* [31]. This oxygenation pattern is only known in other three species, two of which belong to this subfamily.

Pentaoxygenated xanthones. These compounds are rare in the Guttiferae but common in the Gentianaceae. The new compound, 3,8-dihydroxy-1,2,4-trimethoxyxan-

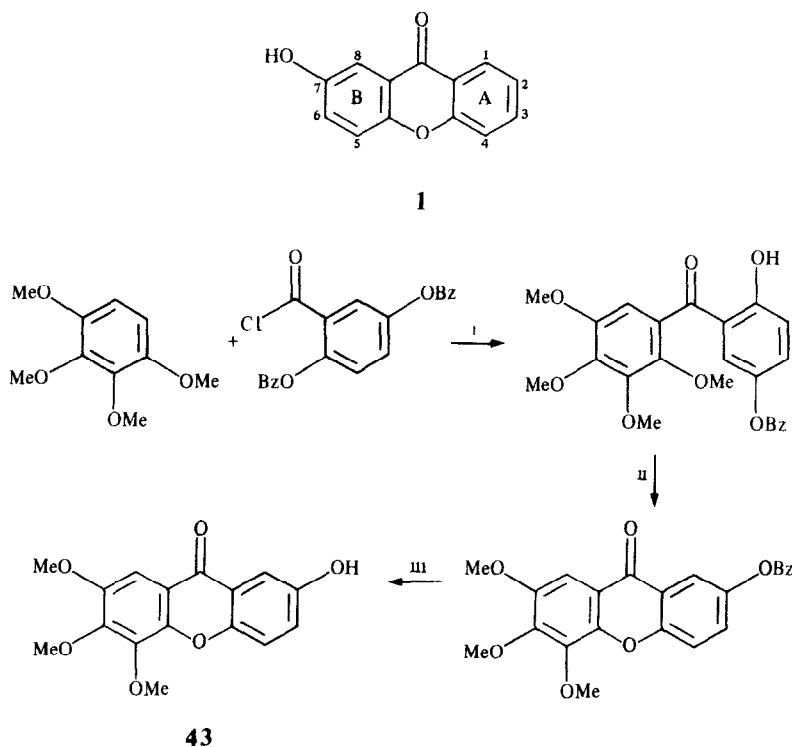
thone (**53**, Table 5) is related to **44** from the same plant, and brings the total number known to five.

Alkylated xanthones

Alkylated xanthones in Guttiferae are usually mono- or di- C_5 -substituted. The C_5 group may be 3-methylbut-2-enyl (as in **54**), or less often 1,1-dimethylprop-2-enyl (as in **89**), and these are frequently cyclized with *ortho* hydroxyls giving 2,2-dimethylpyrano (or dihydropyrano), 2,2,3-trimethylfurano (possible artefacts), or rarely 2-isopropenyldihydrofurano compounds. Occasionally, hydroxylation or hydration of the side chain occurs. C_{10} substituents, in which two prenyl groups are joined together, include geranyl (as in **112**) and lavandulyl (as in **115**).

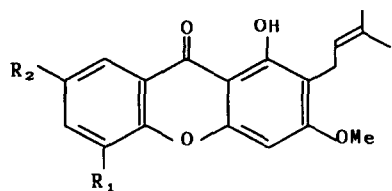
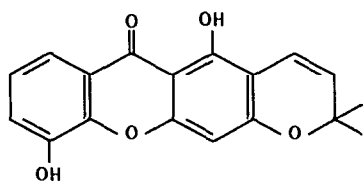
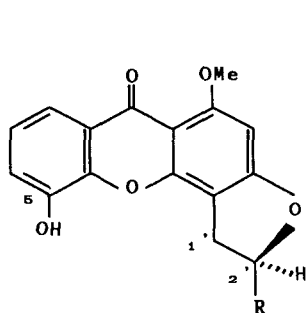
The oxygenation patterns of alkylated xanthones are less diverse than in the unalkylated compounds. Four patterns predominate: 1,3,5-, 1,3,7-, 1,3,5,6-, and 1,3,6,7-Di- and pentaoxygenated compounds are rare.

Mono- C_5 -trioxygenated xanthones Fractionation of the cytotoxic ethanol extract of the roots of *Psorospermum febrifugum* led to the isolation of the anti-leukaemic xanthones, psorospermin (**57**, Table 6) and its chlorohydrin (**58**) [51, 52]. Their ready interconversion suggested a possible artefact. Tandem mass spectrometry has confirmed that psorospermin is a natural product, while the possibility that **58** is an artefact has not been rigorously excluded [53]. The other derivatives (**59–61**), show no anti-leukaemic properties. The absolute stereochemistry of psorospermin has been determined as



Reagents (i) $AlCl_3$ / ether, (ii) Me_4NOH / reflux, (iii) H_2 , Pd / C

Scheme 1

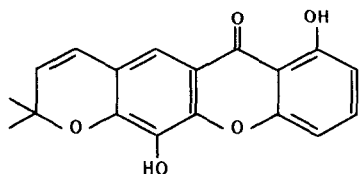
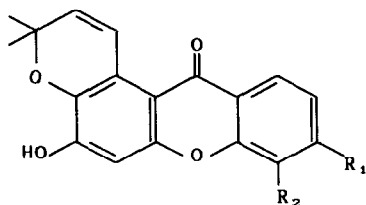
Table 6 Mono-C₅-trioxygenated xanthenes(54) R₁ = OH, R₂ = H**Garcinia mangostana* L. [48,49](55) R₁ = H, R₂ = OH**Garcinia mangostana* L. [48](56) **6-Deoxyjacareubin***Calophyllum zeylanicum* Kosterm. [21](57) R = : **Psorospermin** (2'R,3'R)

(58) R = (2'R,3'S)

(59) R = (2'R,3'R)

(60) R = : **3',4'-Deoxypsorospermin**

(61) R = (5-O-Methyl)

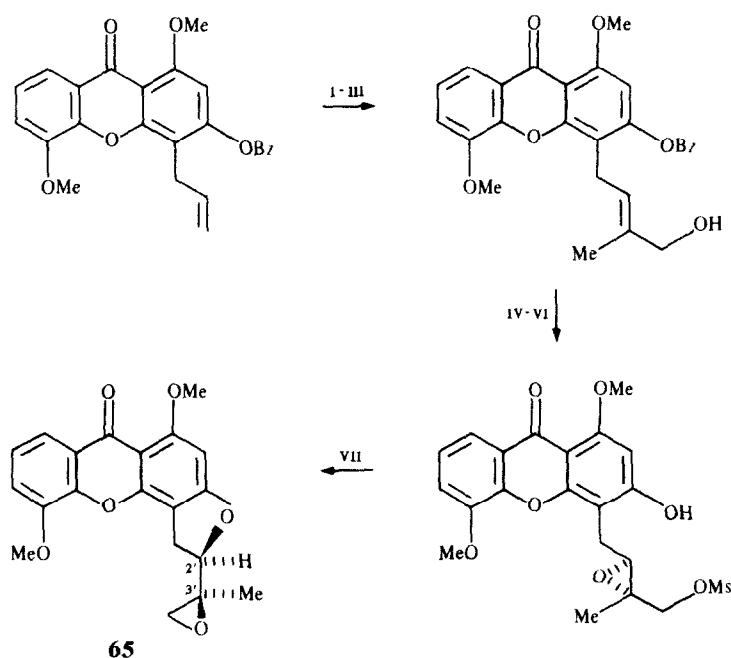
**Psorospermum febrifugum* Sprach.
[51,52,54](62) **Rheediachromenoxanthone***Rheedia brasiliensis* (Mart.)
Pl. and Tr. [58]**Rheedia gardneriana* Pl. and Tr. [59](63) R₁ = H, R₂ = OH; **Hypericanarin****Hypericum canariensis* L. [24](64) R₁ = OH, R₂ = H; **Hyperxanthone****Hypericum sampsonii* Hance [39]

2'R,3'R by an ORD, ¹H NMR and TLC comparison with analogous epoxides derived from rotenone [54].

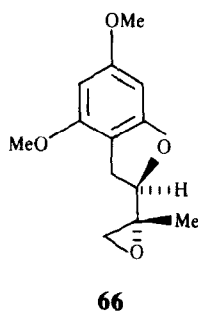
A stereoselective synthesis of 5-methyl-(±)-2'R,3'S-psorospermin (65) has been completed and is shown in part in Scheme 2 [55]. The final step in the synthesis involves two consecutive displacements, producing a racemic mixture of the 2R',3'S and 2'S,3'R epoxides (65).

Progress towards a total synthesis of psorospermin has been made by the enantioselective preparation of the 2'R,3'R-dihydrobenzofuran (66) [56]. A synthesis of the 5-methyl-1',2'-dehydro derivative of 3',4'-deoxypsorospermin (60) has also been reported [57].

Rheediachromenoxanthone (62) and hyperxanthone (64) appear to be derived from tetraoxygenated xan-



Reagents (i) $\text{OsO}_4/\text{NaIO}_4$, dioxane / H_2O (ii) $\text{Ph}_3\text{P}=\text{C}(\text{Me})\text{CO}_2\text{Et}$, benzene (iii) LiAlH_4 / THF (iv) MCPBA, CH_2Cl_2 (v) MsCl/py (vi) H_2 , Pd/C (vii) KO^tBu



Scheme 2

thones by nuclear reduction at the 3- and 1-positions respectively. In fact, **64** occurs with the expected precursor, toxylloxanthone **B** (**82**) in *Hypericum sampsonii*. The unusual 4,6,7- (or 2,3,5-?) oxygenation pattern of hypericanarin (**63**) is otherwise only known in **27** from *Hypericum androsaemum*. The structures of **63** and **64** are supported by significantly different physical data.

Di- C_5 -trioxygenated xanthenes Garcimone A from *Garcinia mangostana* was assigned a novel 1,3,6-trihydroxy structure (**68**, Table 7), an oxygenation pattern unknown in the Guttiferae. However, the synthetic substance, prepared by the prenylation of 1,3,6-trihydroxyxanthone, showed little similarity with the natural compound [66].

The stem bark of *Garcinia quadrifaria* produces xanthone **69**, which shows a rare prenylation, *para*, rather than *ortho* to a hydroxyl group. 6-Deoxy- γ -mangostin (**70**) has been isolated from the seed arils of *Garcinia mangostana* fruit and is a possible biosynthetic precursor

to mangostin (**70**; $\text{R}_1 = \text{Me}$, $\text{R}_2 = \text{OH}$), the major metabolite from the same plant (see Biosynthesis).

The stem bark of *Calophyllum walkeri* contains xanthenes **74** and **76–78**. The hydroxymethyl group in thwaitesixanthanol (**77**) was located by mild acetylation. Shielding of H-3'' was observed, while H-3' was unaffected [65]. The 2-isopropenyldihydrofuran group of **78** is otherwise only known (without the hydroxy) in 3',4'-deoxyporospermin (**60**). The *cis* configuration of **78** was assigned from the 6 Hz coupling between the 2'' and 3'' protons. The 3'' proton was also coupled to H-6 and strongly deshielded, suggesting that it is almost in the same plane as the xanthone nucleus.

Mono- C_5 -tetrahydroxyxanthenes Xanthone **79** (Table 8) is the second 1,3,5,8-tetra-oxygenated xanthone from *G. mangostana*, the other being gartanin (**79**; $\text{R}_1 = \text{H}$, $\text{R}_2 = \text{prenyl}$). This oxidation pattern is otherwise unknown in the Guttiferae but common in the Gentianaceae. Three 2-prenyl-3-methoxyxanthenes (**54**, **55**, and **79**) have been

Table 7 Di-C₅-trioxygenated xanthenes

	<p>(67) $R_1 = \text{OH}, R_2 = \text{H}$, 8-Desoxygartanin <i>Rheedia brasiliensis</i> (Mart.) Pl. and Tr [58] <i>Rheedia gardneriana</i> Pl. and Tr [60]</p>
	<p>(68) $R_1 = \text{H}, R_2 = \text{OH}$; Garcinone A * <i>Garcinia mangostana</i> L. [61]</p>
	<p>(69) 4,8-Di(3-methylbut-2-enyl)-1,3,5-trihydroxyxanthone * <i>Garcinia quadrifaria</i> Baill ex Pierre [62]</p>
	<p>(70) $R_1=R_2=\text{H}$; 6-Deoxy-γ-mangostin * <i>Calophyllum thwaitesi</i> Pl. and Tr [63] <i>Garcinia mangostana</i> L. [49]</p>
	<p>(71) $R_1=\text{H}, R_2=\text{Me}$; Calcocalabaxanthone (6-Deoxymangostin) <i>Calophyllum bracteatum</i> Thw. [63] * <i>Calophyllum calaba</i> var. <i>calaba</i> L. [63,64]</p>
	<p>(72) Trapezifolixanthone <i>Calophyllum calaba</i> var. <i>calaba</i> L. [63]</p>
	<p>(73) $R = \text{Me}$; Calabaxanthone <i>Calophyllum calaba</i> var. <i>calaba</i> L. [64] <i>Calophyllum zeylanicum</i> Kosterm. [21] <i>Garcinia mangostana</i> L. [49]</p>
	<p>(74) $R = \text{H}$; Demethylcalabaxanthone * <i>Calophyllum walkeri</i> Wight [65] <i>Garcinia mangostana</i> L. [49]</p>
	<p>(75) Calothwaitesixanthone * <i>Calophyllum thwaitesi</i> Pl. and Tr. [63]</p>

Table 7 Continued

	<p>(76) R = Me; <u>Thwaitesixanthone</u> <i>Calophyllum walkeri</i> Wight [65]</p>
	<p>(77) R = CH₂OH, <u>Thwaitesixanthanol</u> *<i>Calophyllum walkeri</i> Wight [65]</p>
	<p>(78) *<i>Calophyllum walkeri</i> Wight [65]</p>

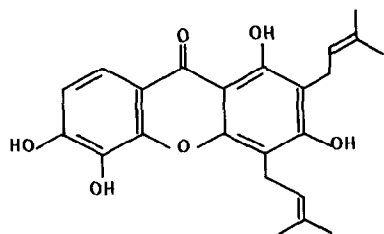
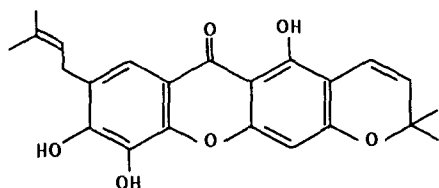
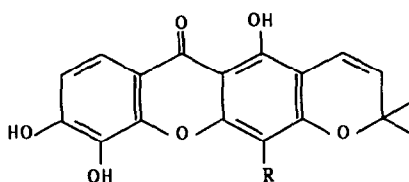
Table 8 Mono-C₅-tetraoxygenated xanthones

	<p>(79) R₁ = Me, R₂ = H <u>1,5,8-Trihydroxy-3-methoxy-2-(3-methylbut-2-enyl)xanthone</u> *<i>Garcinia mangostana</i> L [67]</p>
	<p>(80) <u>1,3,6,7-Tetrahydroxy-8-(3-methylbut-2-enyl)xanthone</u> *<i>Hypericum androsaemum</i> L [34]</p>
	<p>(81) <u>Jacareubin</u> <i>Calophyllum zeylanicum</i> Kosterm [21]</p>
	<p>(82) <u>Toxyloxanthone B</u> †<i>Hypericum androsaemum</i> L [34] <i>Hypericum sampsonii</i> Hance [39]</p>

found in *Garcinia mangostana*. The 3-methoxy group prevents further prenylation in the 4-position, however the presence of a methylating enzyme does not preclude 4-prenylation as is shown by the presence of **67** in the same plant.

Toxyloxanthone B (**82**), first found from the Moraceae, has been isolated from *Hypericum androsaemum* together with its probable biogenetic precursor (**80**).

Di-C₅-tetraoxygenated xanthones While the xanthones in this category (Table 9) are more numerous than

Table 9 Di-C₅-tetraoxygenated xanthenes(83) **Xanthone V_{1a}****Vismia guineensis* (L.) Choisy [68](84) **7-Prenyljacereubin****Rheedia gardneriana* Pl. and Tr. [60](85) R = 3-Methylbut-2-enyl; **Xanthone V₁****Vismia guineensis* (L.) Choisy [68]

(86) R = 1,1-Dimethylprop-2-enyl;

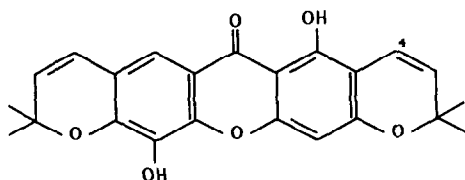
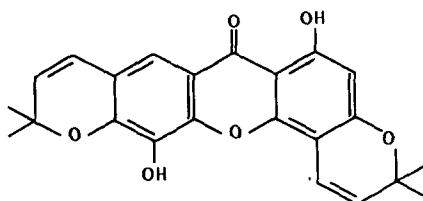
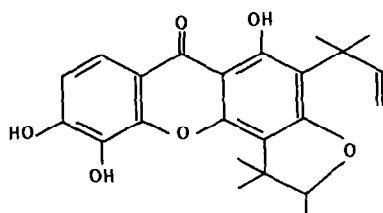
Macluraxanthone*Garcinia ovalifolia* Oliv. [69]†*Rheedia benthamiana* Pl. and Tr. [70]*Rheedia brasiliensis* (Mart.)
Pl. and Tr. [58]
Rheedia gardneriana Pl. and Tr. [59](87) **Pyranojacereubin***Rheedia brasiliensis* (Mart.)
Pl. and Tr. [58]**Rheedia gardneriana* Pl. and Tr. [60](88) **Rheediaxanthone A****Garcinia densivenia* Engl. [71]*Garcinia staudtii* Engl. [62]*Rheedia benthamiana* Pl. and Tr. [70]*Rheedia brasiliensis* (Mart.)
Pl. and Tr. [58]
Rheedia gardneriana Pl. and Tr. [59](89) **Rheediaxanthone B****Rheedia benthamiana* Pl. and Tr. [70]*Rheedia brasiliensis* (Mart.)
Pl. and Tr. [58]
Rheedia gardneriana Pl. and Tr. [59]

Table 9 Continued

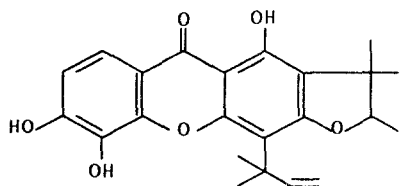
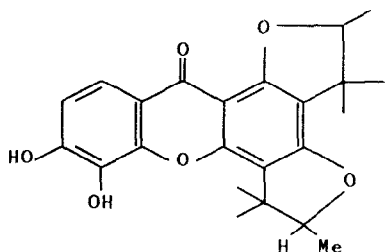
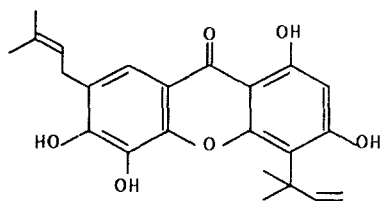
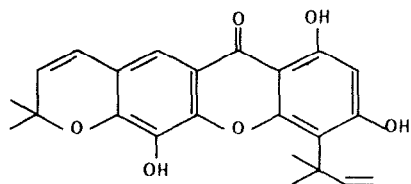
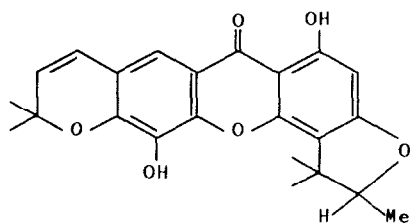
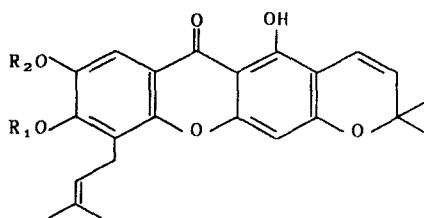
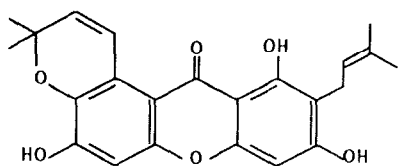
(90) **Isorheediaxanthone B***Garcinia polyantha* Oliv. [72]*Rheedia brasiliensis* (Mart.)Pl and Tr. [58]
Rheedia gardneriana* Pl and Tr. [59](91) **Rheediaxanthone C**Rheedia benthamiana* Pl. and Tr [70]*Rheedia brasiliensis* (Mart.)Pl and Tr. [58]
Rheedia gardneriana Pl. and Tr [59](92) **Rheedia brasiliensis* (Mart.)
Pl. and Tr [58](93) **Rheedia brasiliensis* (Mart.)
Pl. and Tr [58](94) **Rheedia brasiliensis* (Mart.)
Pl and Tr. [58](95) $R_1 = \text{Me}, R_2 = \text{H}$; **Manglexanthone****Touomita mangle* G Maritz [73](96) **Garcinone B****Garcinia mangostana* L [61,147]

Table 9 Continued

	(97) * <i>Garcinia mangostana</i> L. [74]
	(98) 3',4'-Dihydro 93; <u>3-Isomangostin</u> * <i>Garcinia mangostana</i> L. [49]
	(99) R = H; <u>Garcinone C</u> * <i>Garcinia mangostana</i> L. [61]
	(100) R = Me; <u>Garcinone D</u> * <i>Garcinia mangostana</i> L. [75]
	(101) 3',4'-Dihydro 96; <u>3-Isomangostin Hydrate</u> * <i>Garcinia mangostana</i> L. [49]
	(102) R = -C=CHMe ₂ ; <u>1-Isomangostin</u>
	(103) R = -CH ₂ C(OH)Me ₂ ; <u>1-Isomangostin Hydrate</u> * <i>Garcinia mangostana</i> L. [49]
	(104) <u>BR-xanthone-A</u> * <i>Garcinia mangostana</i> L. [83]

the monoprenylated compounds, there is little structural variation. This is due in part to the fact that most of these xanthenes have been isolated from only a few species of the closely allied genera *Garcinia* and *Rheedia*. Dialkylated xanthenes, especially dipyrano compounds, give rise to doubly-charged ions in their mass spectra due to the simultaneous loss of two alkyl fragments [58]. Apart from macluraxanthone (86), which is known from a Moraceae species, all the xanthenes in Table 9 are new natural products.

A dipyranoxanthone from *Garcinia densivenia* was originally assigned the linear structure of pyranojacereubin (87) on the basis of the size of the diamagnetic shift of H-4' upon O-acetylation [71]. Subsequently, when a similar compound was isolated from *Rheedia gardneriana* and the two compounds compared, the structure of the *G. densivenia* xanthone was revised to that of rheediaxanthone A (88), and the new compound assigned the structure of pyranojacereubin (87) [60]. This example serves to highlight the problem of differentiating between substituents at the 2- and 4-positions. Upon acetylation, H-4' of 87 was deshielded by $\delta 0.26$ (expected $\sim \delta 0.30$), compared to $\delta 0.19$ for H-4' of 88 [60].

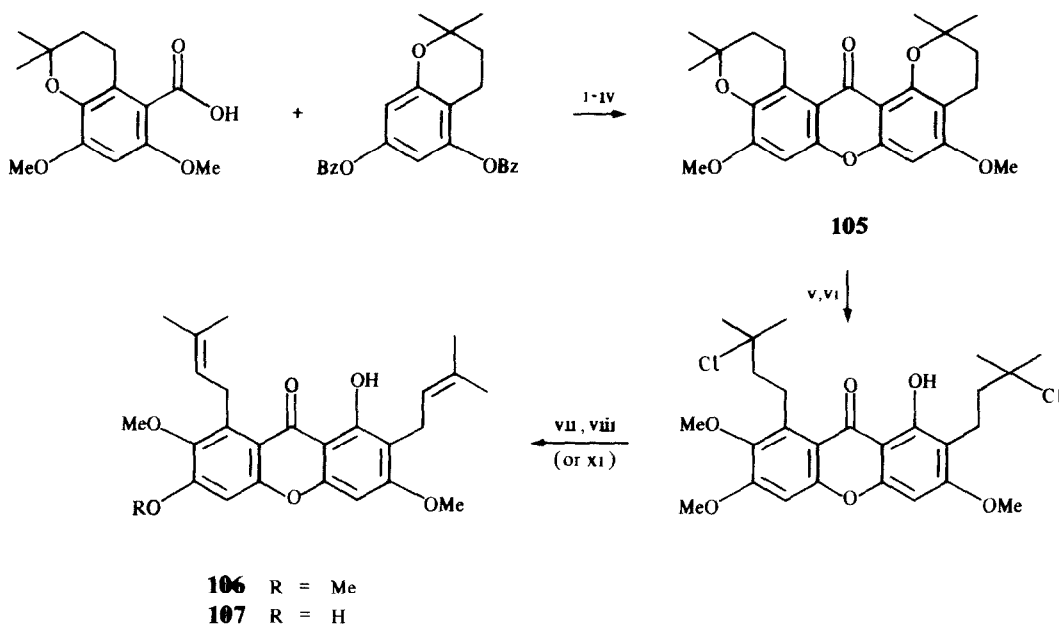
Compounds which contain a 1,1,2-trimethyldihydrofuran ring are possible artefacts due to the ease with which the parent 1,1-dimethylallyl group cyclizes with an

ortho-hydroxyl. Rheediaxanthone B (89) and xanthenes 90 and 94 are optically active and should therefore be true natural products; however rheediaxanthone C (91), although also optically active, is thought to be an artefact of 89 [70].

Nine of the xanthenes in Table 9 were isolated from *Rheedia brasiliensis* and are biogenetically closely related (eg. 92–94). The location of the methoxyl in manglexanthone (95) was confirmed by its ^{13}C NMR chemical shift ($> \delta 60$), indicating di-*ortho*-substitution [70]. Therefore 95 is apparently different from tovopyrifolin A (95; $\text{R}_1 = \text{H}$, $\text{R}_2 = \text{Me}$) which is found in the same genus [76].

During the last eight years an equal number of papers have described new xanthenes from *Garcinia mangostana*. Those shown in Table 9 all contain the 1,3,6,7-tetraoxygenation pattern and are cyclized or hydrated derivatives of the major metabolite, mangostin (70; $\text{R}_1 = \text{Me}$, $\text{R}_2 = \text{OH}$) or γ -mangostin (70) $\text{R}_1 = \text{H}$, $\text{R}_2 = \text{OH}$). The chromanoxanthenes (98 and 101–104) had been synthesised previously by the *p*-toluenesulphonic acid catalysed cyclization of mangostin [77]. Whereas this reagent gave both linear and angular chroman rings, methanolic HCl gives only the linear product [78].

Dimethylmangostin (106), which may be selectively demethylated to give the natural mangostins, has been synthesized by Lee (Scheme 3) [79]. The problem of



Reagents (i) TFAA/CH₂Cl₂ (ii) H₂, Pd/C (iii) CH₂N₂ (iv) Me₄NOH/py (v) BCl₃/CH₂Cl₂
 (vi) MeI/K₂CO₃ (vii) MeCOCl/K₂CO₃ (viii) LiCl/DMF (xi) KO^tBu/DMSO

Scheme 3

selective prenylation in the 2- and 8-positions was overcome by the preparation of the dichromanoxanthone (105). Ring opening was accomplished with boron trichloride to give a dichloride which underwent dehydrochlorination with lithium chloride in DMF, producing 106. Alternatively, treatment of the dichloride with potassium-*t*-butoxide in DMSO gave β -mangostin (107).

Tri-C₅-tetraoxygenated xanthones. Five xanthones of this type are now known, and the two new ones are shown in Table 10. The structure of garcinone F (108) is tentative, and based mainly on the formation of a trichromanoxanthone compound with methanolic HCl [78].

The novel ring B substitution pattern of nervosaxanthone (109) was suggested by the deshielded methylene of one of the prenyl groups, placing it *peri* to the carbonyl, and the ¹H NMR resonance of the single aromatic proton [72]. The arrangement of the 2- and 4-substituents was determined by an ¹H NMR study of the tri- and tetra-acetates.

Di-C₅-penta-oxygenated xanthones. The first natural examples of this type are xanthones V₂ (110) and V_{2a} (111) (Table 10) from *Vismia quineensis*. They are the 7-methoxy derivatives of xanthones V₁ (83) and V_{1a} (85) (Table 9). Together these constitute the only occurrences of prenylated xanthones in the genus *Vismia*.

More complex xanthones. *Garcinia pyrifera* shows a link with *G. cowa* and *G. rubra* by producing 8-geranyl-1,3,6,7-tetraoxygenated xanthones (112–114; Table 11) [72]. The optically active maculatoxanthone (115) is the first xanthone with a lavandulyl side chain, although this arrangement is found in three benzophenones—tovo-phenones A and B (141, 142) and xanthochymol (144). Calozeyloxanthone (116) contains a novel C₁₀ arrangement, presumably a cyclized geranyl group.

Calophyllum wightianum produces a palmitic acid clathrate which yielded a xanthone named wightianone [81]. A 5,5-diprenyl structure was proposed. However, on comparison [21], it was found to be identical with a sample of zeyloxanthone (117) isolated from *Calophyllum zeylanicum*. Gambogic acid (118) has been re-isolated from *Garcinia hanburyi* together with a new derivative, neo-gambogic acid (119) [82].

Xanthone glycosides

The C-glucosylxanthones, mangiferin (120; Table 12) and isomangiferin (121) have been found in several species of *Hypericum* and *Cratoxylum pruniflorum* but are not known in the rest of the family. The taxonomic significance of this will be discussed later. There are only two examples of O-glycosides in Guttiferae (122, 123), although neither has been properly characterized. Both O- and C-glucosylxanthones are common in the Gentianaceae [2].

Xanthonolignoids

Kielcorin was first isolated from *Kielmeyera* species, and its structure has been defined as a racemate of the two 5,6-*trans* isomers (124) [88]. It has recently been isolated from several *Hypericum* species and *Vismia guaramirangae*. The other xanthonolignoids in Table 13 also have the *trans* configuration and are optically inactive.

Biogenetically, xanthonolignoids are thought to be formed by the coupling of a cinnamyl alcohol with an *ortho*-dihydroxyxanthone. Kielcorin and 2,3,4-(or 5,6,7)-trioxygenated xanthones co-occur in *Hypericum calycinum* and *H. ericoides* as well as in three *Kielmeyera*

Table 10 Tri-C₅-tetraoxygenated and Di-C₅-pentaxygenated xanthenes

	<p>(108) <u>Garcinone E</u> *<i>Garcinia mangostana</i> L. [78]</p>
	<p>(109) <u>Nervosaxanthone</u> *<i>Garcinia nervosa</i> Miq. [72]</p>
	<p>(110) <u>Xanthone V₂</u> *<i>Vismia guineensis</i> (L.) Choisy [68]</p>
	<p>(111) <u>Xanthone V_{2a}</u> *<i>Vismia guineensis</i> (L.) Choisy [68]</p>

species. The 1,5,6,7-tetraoxygenated candensins A and C (125, 126) are found alongside similarly substituted xanthenes in the two *Caraipa* species and *H. canariensis*. Furthermore, syringaresinol (129), a likely precursor to the C₆C₃ moiety of candensins A and C, has been isolated as the diacetate from *Vismia quaramirangae*.

Kielcorin (124) has been synthesized by two different routes [91, 92]. One involves the biomimetic coupling of 2-methoxy-3,4-dihydroxyxanthone (130, Scheme 4) and conferyl alcohol (131) [91]. *cis*-Kielcorin (124, 5,6-*cis*) was a minor product.

Candensin D and hypericorin from *Hypericum canariensis* and *H. mysorens* respectively, have been assigned the same structure (127) and their physical data compare reasonably well. An isomer of kielcorin, named kielcorin B has been identified in *Kielmeyera coriacea* and tentatively assigned structure 128.

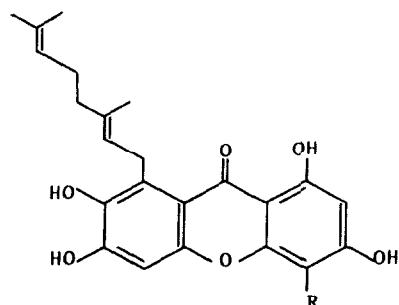
Benzophenones

Benzophenones are of interest as they have been implicated in xanthone biosynthesis. Simple benzophenones are rare in the Guttiferae. They have been found in the heartwood of five species (Table 14)—usually with analogous xanthenes. For example, maclurin (134) occurs with 1,3,6,7-tetrahydroxyxanthone (48) in *Garcinia mangostana*, with 48 and 1,3,5,6-tetrahydroxyxanthone in *Symphonia globulifera* and with xanthochymol (144) in *Garcinia xanthochymus*.

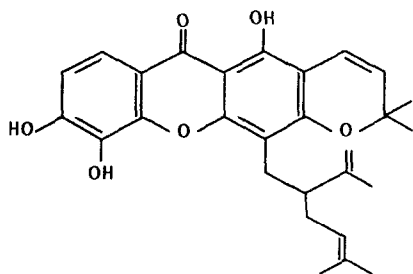
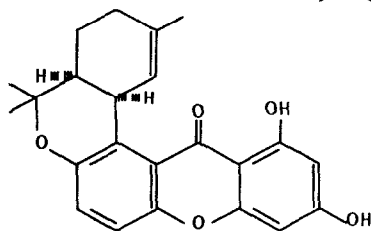
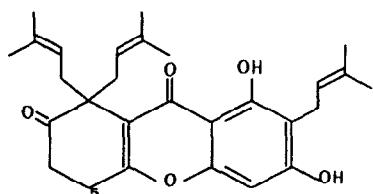
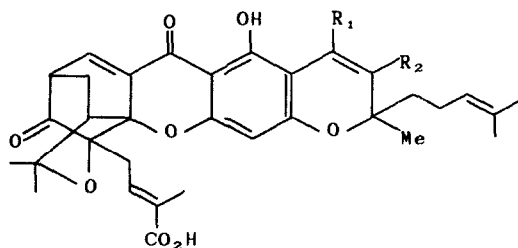
Prenylated benzophenones

These compounds shown in Table 15, may be divided into two groups, true benzophenones (136–142) and those with reduced A rings, the so-called polyisoprenylated

Table 11 More complex xanthenes

(112) R = H, Rubraxanthone(113) R = 3-Methyl-2-butenyl; *Isowanin

(114) R = 3-Methyl-4-hydroxy-2-butenyl;

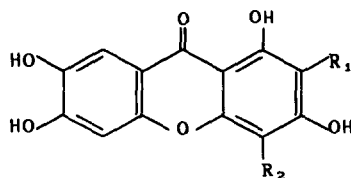
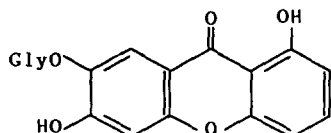
*Isocowanol*Garcinia pyrifera* Ridl. [72](115) Maculatoxanthone**Hypericum maculatum* Crtz. [80](116) Calozeyloxanthone**Calophyllum zeylanicum* Kosterm. [21](117) Zeyloxanthone (Wightianone)*Calophyllum wightianum* T Anders [81](118) $R_1=R_2=H$, Gambogic acid(119) $R_1=H, OH, R_2=H_2$ (no d b);Neogambogic acid**Garcinia hanburyi* Hook f [82]

benzophenones (143–148). Of the former group, five contain unsubstituted B rings, and therefore have no known xanthone analogues. The vismiaphenones (136–139) have been synthesized by the prenylation of suitably substituted benzophenones [98–100]. Tovo-phenones A and B (141, 142) contain the rare lavandulyl side chain, and kolanone (143) from the edible fruit of

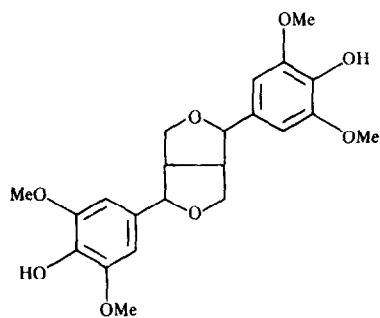
Garcinia kola, shows broad antimicrobial activity [102]

There has been some controversy regarding the structures of the four closely related compounds containing a bicyclononene moiety (144–147). The X-ray determined structures of xanthochymol (144) and isoxanthochymol (146) are shown in Table 15. Cambogin (147) was shown to be the antipode of the latter and camboginol (145),

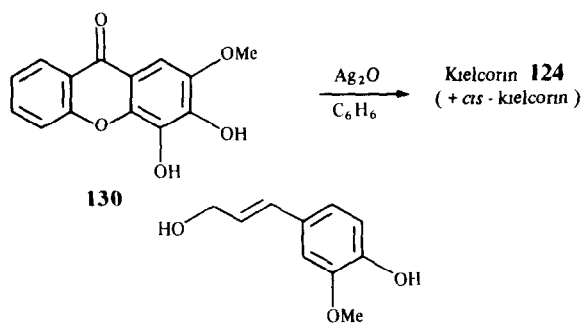
Table 12. Xanthone glycosides

(120) $R_1 = \beta\text{-D-Glucopyranosyl}$, $R_2 = \text{H}$;**Mangiferin***Cratoxylum pruniflorum* Kurz. [130]*Hypericum aucheri* Jaub et Sprach [47]*Hypericum barbatum* Jack [84]*Hypericum boissieri* Petrovic [85][†]*Hypericum humifusum* L. [86]*Hypericum maculatum* Crantz [84]*Hypericum rocheli* Griseb. and Schenk. [85]*Hypericum rumeliacum* Boiss [84]*Hypericum sampsonii* Hance [39](121) $R_1 = \text{H}$, $R_2 = \beta\text{-D-Glucopyranosyl}$;**Isomangiferin***Cratoxylum pruniflorum* Kurz. [130]*Hypericum boissieri* Petrovic [85]*Hypericum hirsutum* [87]*Hypericum rocheli* Griseb[†]*Hypericum sampsonii* Hance [39] and Schenk [85](122) $R_1 = R_2 = \text{H}$; **1,3,6,7-Tetrahydroxy-
O-glucosylxanthone***Garcinia mangostana* L. [46](123) **1,6,7-Trihydroxyxanthone-
7-O-glycoside***Platonia insignis* Mart. [29]

(Gly: unidentified sugar)



129



130

131

Scheme 4

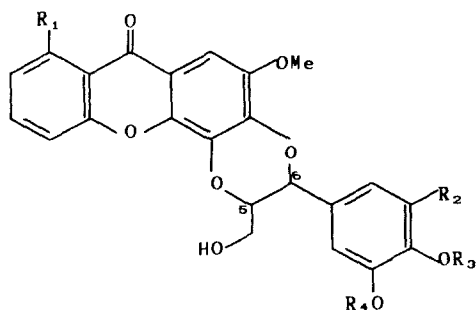
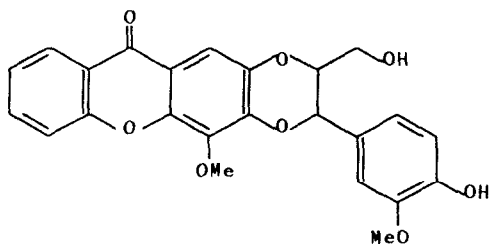
which can be prepared from cambogin, was assigned the structure of the enantiomer of xanthochymol with a shifted double bond [106].

Two related pigments, namely garcinol and isogarcinol were isolated from *Garcinia indica* [107]. Garcinol was assigned the structure of camboginol (145), however, their UV spectra in ethanol differed significantly, and it was suggested that camboginol lacked the extended conjugation in the cyclohexane ring. The UV spectrum of camboginol was repeated in cyclohexane and shown to be compatible with the originally proposed structure. A solvent-dependent UV spectrum had already been noted for xanthochymol [108]. Ultimately, the X-ray crystal structure of isogarcinol indicated its identity with cambogin (147), and the formation of isogarcinol from garcinol, as well as physical data, suggest that the latter is identical to camboginol (145) [111].

A related compound, nemorosanol, isolated from the fruits of *Clusia nemorosa* has been assigned the tricyclodecane structure (148). It is suggested that the various polyisoprenylated benzophenones may originate from a common precursor (150, R = prenyl or 2-isopropenylhex-4-enyl) by cyclization between C-1 and C-6 to form the bicyclic compounds and C-10-C-1 and C-6-C-7 to form 148.

Sarothralin (149) from *Hypericum japonicum* has antimicrobial properties, and is closely related to other filicinic acid derivatives (which lack a benzoyl group) found in this and other *Hypericum* species. Three other

Table 13 Xanthonolignoids

(124) $R_1=H$, $R_2=H$, $R_3=H$, $R_4=Me$.**Kielcorin***Hypericum androsaemum* L [34,88]*Hypericum calycinum* L [88]*Hypericum canariensis* L [24]*Hypericum ericoides* L [89]*Hypericum maculatum* Crantz [88]*Hypericum perforatum* L. [88]*Vismia guaramirangae* Huber [28](125) $R_1=OH$, $R_2=H$, $R_3=OMe$, $R_4=H$ **Candensin A***Cariapa grandiflora* Mart [29]*Cariapa valor* Paula [29]*Vismia guaramirangae* Huber [28](126) $R_1=OH$, $R_2=OMe$, $R_3=H$, $R_4=Me$.**Candensin C***Hypericum canariensis* L [24]**Vismia guaramirangae* Huber [28](127) $R_1=H$, $R_2=OMe$, $R_3=H$, $R_4=Me$.**Candensin D (Hypericorin)****Hypericum canariensis* L [24]*Hypericum mysorensense* [90]**(128) Kielcorin B****Kielmeyera coriacea* Mart. [91]

prenylated benzophenones (bronianone, clusianone, and marupone) were mentioned in the earlier review [1].

Finally, one other compound of interest is hermionic acid. Originally isolated from *Garcinia hermonii*, and assigned a diphenyl structure [1], it has now been re-

isolated from *G. quaesita*, and on the basis of its conversion to a xanthone and ^{13}C NMR evidence, its structure has been revised to the diphenylether (149) [113]. Decarboxylated hermionic acid (150) and its demethyl derivative (151), named quaesitol were also isolated [114].

Table 14 Benzophenones

	<p>(132) R = H; (Hydrocotoin) 2-Hydroxy-4,6-dimethoxybenzophenone</p>
	<p>(133) R = OH 2,3'-Dihydroxy-4,6-dimethoxybenzophenone <i>Allanblackia floribunda</i> Oliv. [93]</p>
	<p>(134) 2,3',4,4',6-Pentahydroxybenzophenone (Maclurin) <i>Garcinia mangostana</i> L. [46] <i>Garcinia xanthochymus</i> Hook. f [30] <i>Symphonia globulifera</i> L. [94]</p>
	<p>(135) 2,3',4,5',6-Pentahydroxybenzophenone <i>Garcinia pedunculata</i> Roxb. [95]</p>

Biogenetically, these compounds appear to be very closely related to xanthenes.

CHEMOTAXONOMY

Recent reviews have outlined the principles of chemotaxonomy and the taxonomic importance of various classes of secondary metabolites [115-117]. As xanthenes occur widely in only a few, unrelated families of higher plants, their potential taxonomic value is restricted to within these few families. This section will examine the distribution of xanthenes within the Guttiferae.

The oxygenation pattern is the most variable structural feature of Guttiferae xanthenes, and almost forty different patterns are known. The major oxygenation patterns are shown at the top of Table 16, grouped mainly according to B-ring oxygenation. The number of species in which xanthenes with each oxygenation pattern occur is given for various sub-divisions of the family (cf. Table 1).

It can be seen from Table 16 that the variation of xanthone oxygenation pattern is of some systematic significance. Some oxygenation patterns appear throughout the family, while others are restricted to certain plant groups. Clearly, the first three of the six subfamilies each produce xanthenes with a different range of oxygenation patterns, suggesting that there are different biogenetic constraints on the species of each group. A similar or somewhat narrower range of oxygenation patterns is shown by the tribes or genera of each of these subfamilies. This overall picture may become clearer as more plants are studied.

A species from a particular genus generally does not contain the whole range of oxygenated xanthenes found

in the genus. Some species appear not to synthesize xanthenes, e.g. *Calophyllum macrocarpum*. This plant accumulates a C_6C_1 compound, 3,4-dihydroxybenzaldehyde, perhaps, it is suggested, due to the absence of a key enzyme [65]. Similarly, some species appear to produce only benzophenones, or only simple rather than prenylated xanthenes.

At first sight, such wide variations within a genus may appear to suggest that xanthenes are of little taxonomic value at this level; however, when considered with other taxonomic characters, xanthenes may be useful in aiding infra-generic classifications. For example, closely related species of *Calophyllum* have many xanthenes in common.

The xanthone distribution within each subfamily will now be examined in more detail. Besides xanthenes, the Guttiferae contain a wide variety of other metabolites many of which are also of taxonomic value and these will be mentioned briefly.

Kielmeyeroideae

The species of this South American subfamily are characterized by an abundance of simple oxygenated xanthenes. The two tribes, Kielmeyereae and Caraipeae, show nine oxygenation patterns in common (Table 16). Several rather unique patterns containing 7,8-oxygenation are evident, and interestingly, xanthenes with 6,7-dioxygenated B-rings have not been found. Simple 1,3,5-trioxygenated xanthenes are common only in this subfamily, and perhaps related to this is the rare occurrence of prenylated xanthenes. Prenylation is absent in the Caraipeae tribe, but three prenylated xanthenes have been found in species of the Kielmeyereae tribe. Two of

these xanthenes are very common in *Calophyllum* species.

A further difference between the two tribes is that xanthenes without 1-oxygenation are common only in the Kielmeyereae tribe. Similarly, xanthonolignoids, which otherwise only occur in the Hypericoideae subfamily, are 5,6,7-trioxygenated in *Kielmeyera* species but 1,5,6,7-tetraoxygenated in *Caraipa* species.

Remarkably, the two genera of the Caraipeae tribe, so far have no oxygenation patterns in common (*Caraipa*: 7-, 1,3,7-, 1,5,6,7-, 5,6,7- and 1,6,7,8-oxygenation, and *Haploclathra*: 1,7-, 1,3,5-, 1,3,5,6-, 1,3,7,8- and 1,7,8-oxygenation).

For a long time a morphological link has been recognised between the Kielmeyeroideae and the family Bonnetiaceae (*Bonnetia* and *Archytaea* genera), and Hutchinson has even grouped the two together as a single family [118]. However, the Bonnetiaceae have also been classified as a tribe in Theaceae [119], a related family in which xanthenes have not been found. The xanthenes that have recently been isolated from two Bonnetiaceae species show the characteristic oxidation patterns of the Kiel-

meyeroideae [31], and have led one taxonomist to submerge the Bonnetiaceae into this subfamily [13].

Calophylloideae

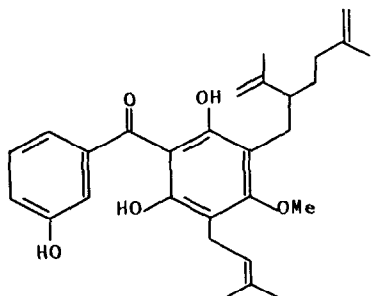
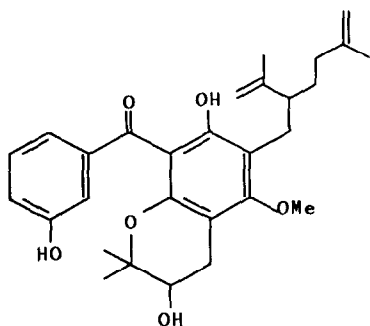
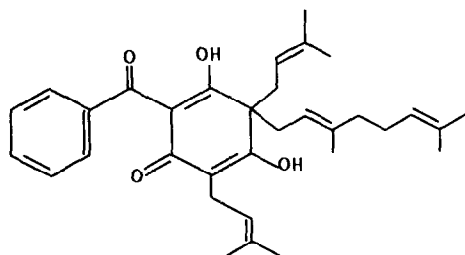
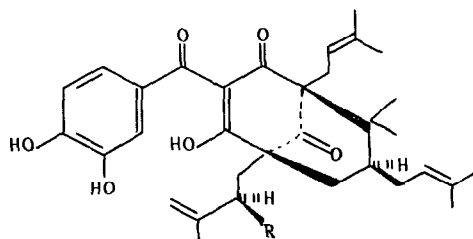
In Table 16 this subfamily has been divided into two groups; *Calophyllum* and *Mesua/Mammea*. Both groups contain a similar range of simple xanthenes, and show the same common oxygenation patterns (1,5-, 1,7- and 1,5,6-), but there is a clear distinction, in that the latter group shows an almost total absence of prenylated xanthenes. Only two are known from a species of *Kayea* (= *Mesua*) [1].

The 180 species of *Calophyllum* are found mainly from India to New Guinea [120]. The genus appears to be the most homogeneous in the family as far as xanthone distribution is concerned. Almost all the 21 xanthone-containing species show both simple and prenylated xanthenes. Jacareubin (81) occurs in 17 of these species (and with 6-deoxyjacareubin (56) in 11), and is therefore classed as a taxonomic marker for the genus [121]. This

Table 15 Prenylated benzophenones

	<p>(136) $R_1 = H, R_2 = Me$. Vismiapphenone A *<i>Vismia decipiens</i> Schlecht-Cham [96] <i>Vismia guaramirangae</i> Huber [28]</p>
	<p>(137) $R_1 = Me, R_2 = H$; Vismiapphenone C *<i>Vismia guaramirangae</i> Huber [28]</p>
	<p>(138) Vismiapphenone B <i>Clusia ellipticifolia</i> Cuatr. [97] *<i>Vismia decipiens</i> Schlecht-Cham [96]</p>
	<p>(139) Isovismiapphenone B <i>Clusia ellipticifolia</i> Cuatr. [97] *<i>Vismia decipiens</i> Schlecht-Cham [96]</p>
	<p>(140) *<i>Clusia ellipticifolia</i> Cuatr. [97]</p>

Table 15 Continued

(141) **Tovophenone A****Tovomita mangle* G.Maritz [101](142) **Tovophenone B****Tovomita mangle* G.Maritz [101](143) **Kolanone****Garcinia kola* Heckel [102]

(144) R = 3-Methylbut-3-enyl

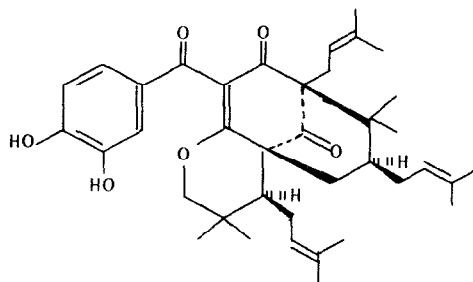
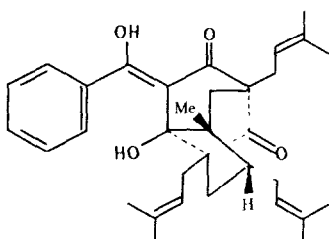
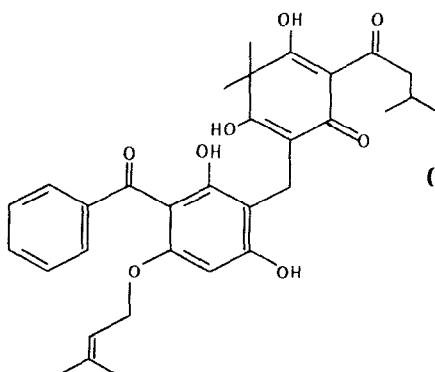
Xanthochymol*Garcinia mannii* Oliv. [103]*Garcinia ovalifolia* Oliv [69]*Garcinia polyantha* Oliv. [72]*Garcinia staudii* Engl. [62]*Garcinia xishuanbannanensis*
Y H.Li [104]*Rheedia madrunno* (HBK)
Pl and Tr [105]

(145) R = 3-Methylbut-2-enyl

"enantiomer" of 144

Camboginol (Garcinol)**Garcinia cambogia* Desr. [106,108]*Garcinia huillensis* [109]*Garcinia indica* Choisy [107,111]

Table 15 Continued

**(146) Isoxanthochymol***Garcinia ovalifolia* Oliv [69]*Garcinia polyantha* Oliv [72]*Garcinia xishuanbannanensis*
Y H Li [104]**(147) Enantiomer of 146****Cambogin (Isogarcinol)****Garcinia cambogia* Desr [106]*Garcinia indica* Choisy [107, 111]**(148) Nemorosonol***Clusia nemorosa* G F.W Meyer [110]**(149) Sarothralin***Hypericum japonicum* Thunb [112]

uniformity may be a reflection of the rather narrow geographic range, and close relationship of many of the species investigated

Interestingly, the xanthone content of *Calophyllum* species often varies greatly with the part of the plant. For example, the three xanthones found in the bark of *C. zeylanicum*, were not found in the timber, which contained seven other xanthones [21]. This phenomenon, as well as the structural similarities of many of the xanthones, makes the chemical comparison of *Calophyllum* species difficult

In a recent review of the genus [120], Stevens equated *C. zeylanicum* (now *C. lankaensis*? [21]) with *C. trapezifolium*. They do in fact have seven xanthones in common, and only differ in that they each produce two xanthones not found in the other [21, 122]. A possible geographical variation is suggested in *C. walkeri* [65]. A plant from India yielded four xanthones that were not found in the Sri Lankan plant of the same name, however, the bark of the latter was not thoroughly investigated [123]. Significant differences have also been found in two varieties of *C. calaba* [63, 64, 124].

Oxygenation patterns

*Numbers in Table refer to number of species in which xanthones with the particular oxygenation have been found.

+Genera in which several plants have been examined from the rest of the subfamily or tribe (cf Table 1)

†These minor oxygenation patterns, apart from one, have been found in one species; 2,5-dioxygenation found in two species

\$1,5-Dioxygenation and 1,6,7-trioxygenation found in eight and two species respectively, remainder found in one species

|XG, xanthone glycosides, XL, xantholignoids, BZ, simple and prenylated benzophenones

Includes three species which contain xanthenes with reduced B-ring

Neoflavonoids (and other 4-substituted coumarins) and chromones occur in many *Calophyllum* species [125, 126].

Clusiodeae

Generally xanthenes occur less frequently in the Clusiodeae than in the preceding subfamilies, and the range of major oxygenation patterns is rather restricted. Xanthenes in which 1-oxygenation is absent are not known, but several unique oxygenation patterns containing 2- and 4-oxygenation occur. Species of both tribes produce prenylated benzophenones, often as the major metabolites, but otherwise the tribes show little in common.

In the Clusiaceae tribe, xanthenes and benzophenones occur in *Tovomita* species, but the large, poorly-studied *Clusia* genus has so far not yielded any xanthenes. Recently, Delle Monache and co-workers have begun to study a number of *Clusia* species [110].

The Garcineae tribe is divided into two groups in Table 16: *Garcinia* and *Rheedia*/*Pentapthalangium*/*Allanblackia*. The tribe appears fairly homogeneous, with each group showing a similar range of simple and prenylated xanthenes.

Xanthenes have been found in about half the *Garcinia* species studied. Biflavonoids occur in almost all species and benzophenones in 11. Waterman and co-workers have systematically examined several *Garcinia* species from tropical Africa. The subsequent chemotaxonomic review of the genus shows that in certain cases xanthone structure correlates well with the infra-generic classification [127].

Three species from the section *Rheediopsis* of the genus have been investigated, and each contains prenylated xanthenes with 1,3,5,6-tetraoxygenation as well as xanthochymol (144). The benzophenones cambogin (147) and camboginol (145) occur in two species from the *Gamogin* section, and the two species which produce gambogic acid (118) and other xanthenes with reduced B-rings are from the section *Hebradendron*. 8-Geranylated xanthenes are only known in three *Garcinia* species, two of which belong to section *Oxycarpus*. It has been suggested that the third species, which has so far not been assigned to a section, may also go here [72]. Such correlations were not found in many other sections, but as only 10% of *Garcinia* species have been studied it is perhaps too early to tell.

Also of interest is the apparent geographical variation of tetraoxygenation patterns [127]. African *Garcinia* species contain 1,3,5,6- (and no 1,3,6,7-) tetraoxygenated xanthenes, whereas Asian species, apart from one, show only 1,3,6,7-tetraoxygenation. Some xanthenes from African *Garcinia* also occur in species of the related South American genus, *Rheedia*, which so far has also yielded only 1,3,5,6-tetraoxygenated xanthenes.

The Moronboideae subfamily is rather small, and the four investigations so far have produced xanthenes similar to those in the Clusiodeae. The Lorostemoideae subfamily contains only two species from which one prenylated xanthone has been isolated.

Hypericoideae

This subfamily of three tribes is closely related to the rest of the Guttiferae [128, 129], although it is sometimes classed as a separate family, the Hypericaceae [12]. The

following evidence supports such a relationship. Prenylated xanthenes, have been found in all three tribes of the Hypericoideae, and although they have unique structures, they are similar to those found in other subfamilies (e.g. 63, 80). The presence of xanthenes with 7,8- and 5,6,7-oxygenation and xanthonolignoids shows a link with the Kielmeyeroideae subfamily, and two benzophenones (138, 139) from *Vismia decipiens* also occur in a species of *Clusia*.

Unlike the rest of the family, the Hypericoideae contain C-glucosylxanthenes. They have been found in several species of *Hypericum* [84, 85] and *Cratoxylum prunifolium* [130], and occur with both simple and prenylated xanthenes, as well as flavonoids. As it appears that these glycosides may be biogenetically different from the xanthenes in the rest of the family (see Biosynthesis), their presence should not be regarded as evidence of 'xanthenes' to support the inclusion of the subfamily in the Guttiferae. Interestingly, simple xanthenes with 6,7-oxygenated B-rings are unusually common in *Hypericum* species. They occur with the similarly substituted glycosides and may be biogenetically related.

A chemotaxonomic survey of the *Hypericum* genus has recently appeared [131]. In addition to xanthenes, many flavonoids and other phloroglucinol derivatives have been found. *Vismia* species produce prenylated anthranoids and other anthracene derivatives (vismiones) [132]. Similar metabolites also occur in *Psorospermum* species [133].

BIOSYNTHESIS

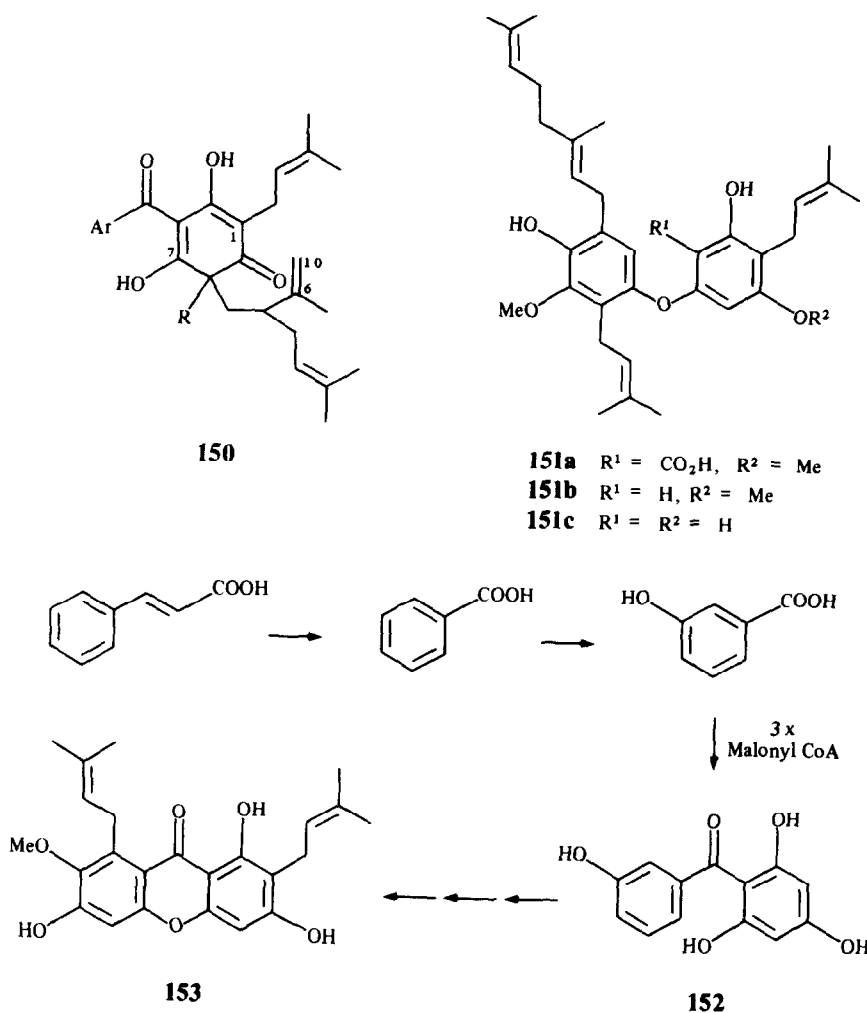
The characteristic oxygenation patterns of xanthenes from higher plants were recognised early on as being due to a mixed shikimate-acetate biogenesis [5]. Along these lines various biosynthetic pathways were proposed [7, 8, 10] and these have been reviewed [1].

Early biosynthetic studies were limited to the 1,3,7-trioxygenated xanthenes of *Gentiana lutea* (Gentianeaceae) [134, 135]. The results indicated that the xanthone nucleus was formed from acetate (ring A) and a C₆C₁ unit derived from phenylalanine. The participation of an intermediate benzophenone (152) was demonstrated by the incorporation of tritiated 152 [135].

The first study on the biosynthesis of xanthenes in the Guttiferae has recently been reported [136]. Cinnamic acid, benzoic acid, *m*-hydroxybenzoic acid and the benzophenone (152) as well as malonic acid were efficient precursors to mangostin (153) in *Garcinia mangostana*, implying the pathway depicted in Scheme 5.

Furthermore, the labelled benzophenone (152) was significantly incorporated into 8-desoxygartanin (67) and gartanin (79, R₁=H, R₂=prenyl) in the same plant. These findings, coupled with the earlier studies on *Gentiana lutea*, indicate the involvement of benzophenone 152 in the biosynthesis of xanthenes with four different oxygenation patterns (1,3,5-, 1,3,7-, 1,3,6,7- and 1,3,5,8-), and suggest that it may be an intermediate in the biosynthesis of most higher plant xanthenes.

The proposed direct *m*-hydroxylation of benzoate (Scheme 5) was suggested, in part, by the poor incorporation of *p*-substituted precursors. Earlier biosynthetic proposals, such as the choice of maclurin (134) as a universal xanthone precursor [10] and the idea of spirodieneone intermediates [7], were formulated with the assumption that shikimate derivatives were necessarily



Scheme 5.

oxygenated in the *p*-position. As it now appears that xanthone biosynthesis may not involve such shikimate derivatives, the earlier proposals require re-evaluation.

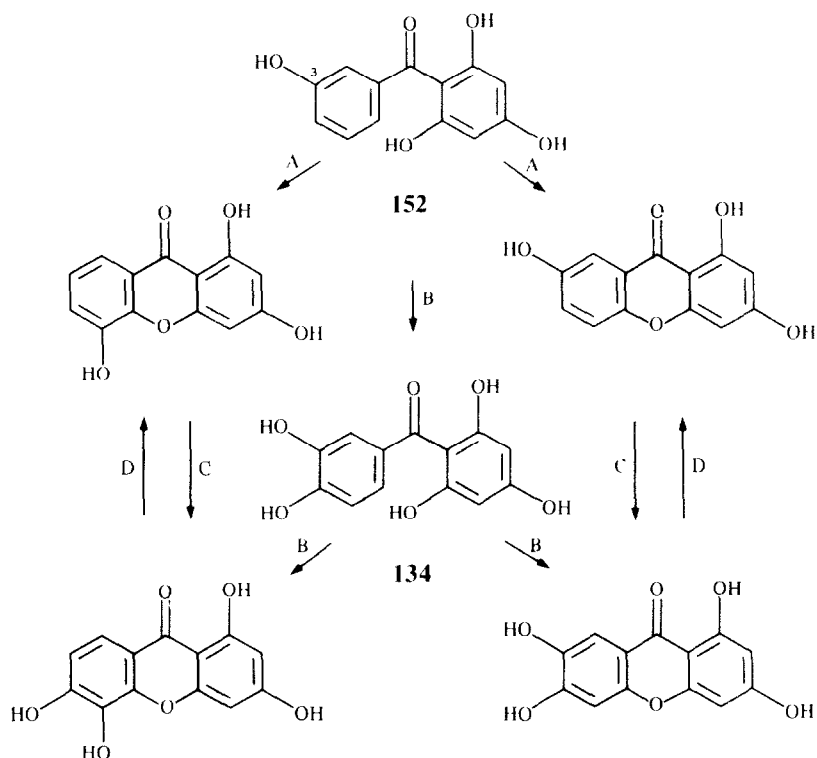
Of the various modes of xanthone formation proposed to date, phenol oxidative coupling [137, 138] can most neatly account for the large variety and co-occurrence of oxygenated xanthenes [8]. Carpenter *et al.* postulated the oxidative coupling of a series of suitably hydroxylated benzophenones to account for the major xanthone oxygenation patterns [8]. This suggestion is compatible with the new biosynthetic results, as the benzophenones all require the *meta*- or 3'-oxygenation for oxidative coupling, and may be derived from benzophenone **152**. For example, the oxidative coupling of **152** can give 1,3,5- and 1,3,7-trihydroxyxanthenes (Scheme 6, Route A) and maclurin (**134**), which may be formed by the 4'-hydroxylation of **152**, can produce 1,3,5,6- and 1,3,6,7-tetrahydroxyxanthenes (Route B).

An alternative to the view that xanthenes are formed from a series of benzophenones, is the proposal of Rezende and Gottlieb that the xanthone oxygenation pattern is modified after initial xanthone formation from a single benzophenone—maclurin (**134**) [10]. The 'pri-

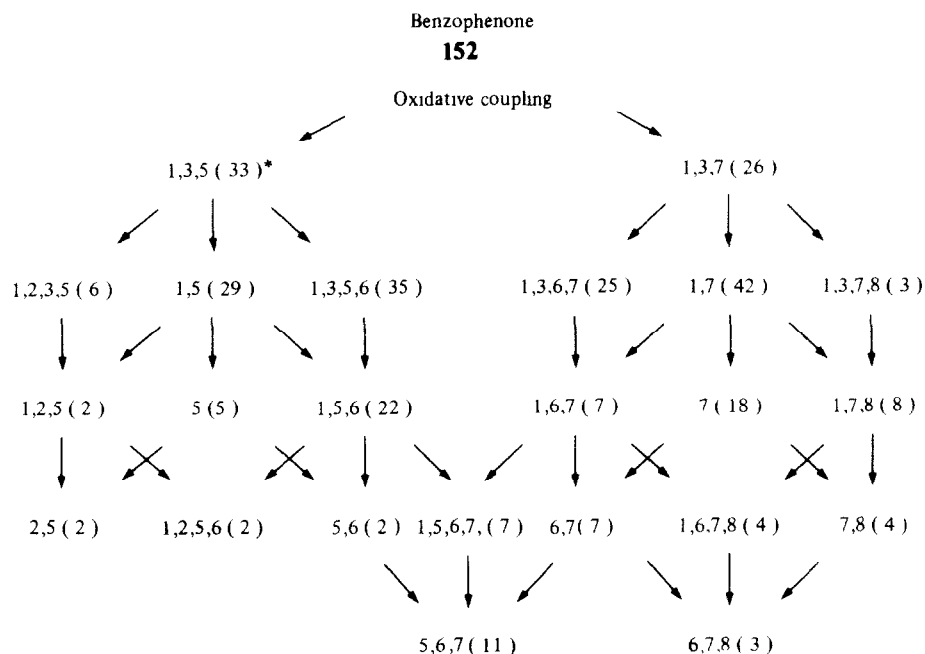
mitive', maclurin-derived 1,3,5,6- and 1,3,6,7-tetraoxygenated xanthenes could lead to all other oxygenation patterns by successive nuclear oxidations and/or reductions. For example, the reduction of the 6-position of these xanthenes would give 1,3,5- and 1,3,7-trioxygenated xanthenes (Scheme 6, Route C).

In view of the recent biosynthetic studies, the 1,3,5- and 1,3,7-trioxygenated xanthenes appear more suited to occupy the positions of 'primitive' xanthenes, as they are clearly more likely to be formed by the direct oxidative coupling of benzophenone **152** (Scheme 6, Route A), than by a three-step route via maclurin (**134**) (Route B + Route C). The 1,3,5,6- and 1,3,6,7-tetraoxygenated xanthenes may alternatively be formed by the 6-oxygenation of the 1,3,5- and 1,3,7-trioxygenated xanthenes (Scheme 6, Route D).

Similar nuclear oxidations/reductions of the trioxygenated xanthenes, can lead to the major oxygenation patterns of Guttiferae xanthenes, as illustrated in Scheme 7. It can be seen that the most common oxygenation patterns (particularly 1,3,5-, 1,3,7-, 1,7-, 1,5-, and 7-) are more directly accessible from **152** than from maclurin (**134**) in the earlier proposal.



Scheme 6



*No of species that produce xanthones with each oxygenation pattern

Scheme 7

The discussion so far has centred on the four predominant oxygenation patterns of prenylated xanthones, but avoided the issue of prenylation. Prenylation is almost

always found *ortho* to an oxygen function. 2-Prenylation is the most common, but is only found in the presence of 1,3-dioxygenation, suggesting perhaps that the prenyl

group inhibits reduction of the 3-hydroxyl, which is very often absent in simple xanthenes

Monoprenylation occurs less often in the 4-, 8-, or 7-positions. A second prenyl group frequently occurs in the 4- or 8-positions. 2,4-Diprenylation is associated with 1,3,5-, 1,3,5,6- or 1,3,5,8-oxygenation, but interestingly, not known with 1,3,7- or 1,3,6,7-oxygenation, and 2,8-diprenylation is limited to 1,3,7- and 1,3,6,7-oxygenation patterns

As with oxygenation, there is little evidence to suggest whether prenylation occurs at the benzophenone or xanthone stage, although the latter is preferred, e.g. ref [139]. 2-Prenyl-1,3,5-trihydroxyxanthone has been postulated as a precursor to 6-deoxyjacareubin (**56**) and jacareubin (**81**) [140], a route involving 6-oxygenation at the xanthone stage, and in keeping with Scheme 7. These three xanthenes are found together in four *Calophyllum* species. Similarly, 2-prenyl-1,3,7-trihydroxyxanthone could lead to mangostin (**153**) via 6-deoxy- γ -mangostin (**70**). The alternative, that **153** and **70** derive from two benzophenones, is less attractive, as it does not indicate such a close relationship between these co-occurring xanthenes. It can be seen that a series of modifications (oxygenation, further prenylation, cyclization etc.) to the two basic 2-prenylxanthenes can produce over 40, or about half of the known prenylated xanthenes.

It had been suggested that mangiferin (**120**) was biogenetically related to flavonoids [5, 141], due to its occurrence in some plants in the presence, or apparently in place of C-glucosylflavones [142], rather than with other xanthenes. Fujita and Inoue have conducted a thorough study on the biosynthesis of mangiferin (**120**) and isomangiferin (**121**) in *Anemarrhena asphodeloides* (Liliaceae) [143, 144]. The results indicate that the xanthone nucleus is indeed formed from a flavonoid-type C_6C_3 precursor (*p*-hydroxycinnamate) coupled with two malonates (Scheme 8). The labelled benzophenones, iriflophenone (**154**) and maclurin (**134**) were significantly

incorporated into **120**, whereas labelled 1,3,6,7-tetrahydroxyxanthone was not, suggesting that the glucosylation occurs at the maclurin stage, and that both **120** and **121** are formed by the oxidative coupling of 3-glucosylmaclurin (**156**).

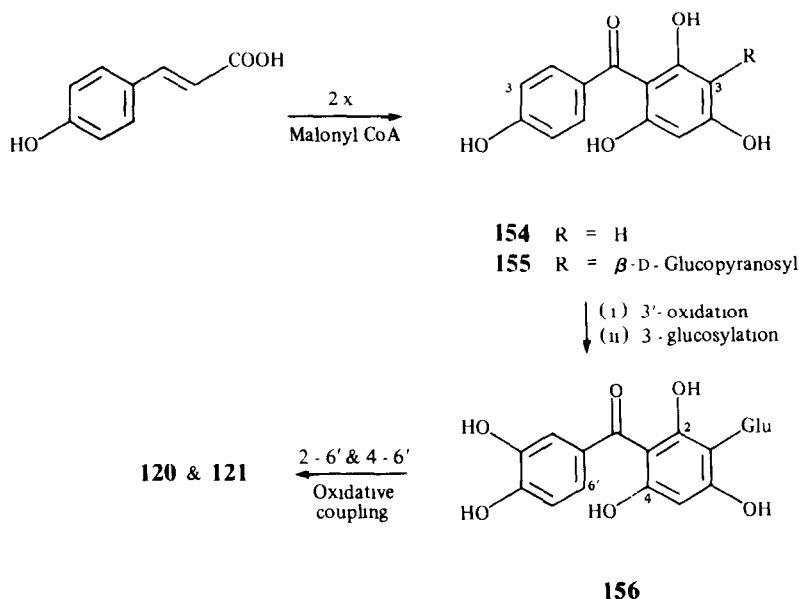
Interestingly, mangiferin (**120**) has recently been found together with 3-C-glucosyliriflophenone (**155**) in two ferns (*Hypodematum*) [145], and with the 1,3,7-xanthone analogue of **155** in *Gentiana lactea* [146].

PHARMACOLOGY

Probably the most important development in this area has been the identification of the cytotoxic xanthone psorospermin (**57**) [51, 52, 54]. It shows significant *in vivo* activity in P338 leukaemia, colon (C6) and mammary (CD) tumour systems [148]. Synthetic studies on psorospermin are in progress [55–57] (see earlier), and analogues are being prepared and tested for similar activity [149].

Mangiferin (**120**) has been the subject of several studies. Anti-inflammatory [150], antihepatotoxic [151] and antiviral (herpes) [152] properties have been reported. Mangiferin has also been shown to cause the *in vitro* activation of the lymphocytes of tumour-bearing rats [153].

In contrast to mangiferin (**120**), which is reported to be a CNS stimulant [154], mangostin (**153**) and several of its derivatives [155] as well as xanthenes from *Calophyllum inophyllum* and *Mesua ferrea* [156] produce CNS depression in rats and mice. Many of these xanthenes, especially mangostin, exhibit significant anti-inflammatory properties at doses of 50 mg/kg [155, 156]. They have no analgesic or antipyretic effect, but mangostin shows anti-ulcer activity. The interference of mangostin with inflammatory and immunopathological responses has been further studied [157]. The synthetic 3,6-di-O-glucosyl-mangostin produces myocardial stimulation and a rise in blood pressure in dogs [155, 158].



Scheme 8

More xanthenes have been tested for *in vitro* inhibition of monoamine oxidases (MAO's) [159]. Two compounds, 1-hydroxy-3,8-dimethoxyxanthone and 1,3-dihydroxy-7,8-dimethoxyxanthone (swertinin) were identified as the most effective inhibitors of type A MAO, but were weak type B MAO inhibitors. In another study 1,5,8-trihydroxy-3-methoxyxanthone also showed selective type A inhibition [146]. A number of xanthenes also inhibit xanthine oxidase [160]. 1,3,6,7-Tetrahydroxyxanthone was the most potent of the compounds tested.

Several prenylated xanthenes possess significant antimicrobial properties. Mangostin shows broad spectrum antibacterial activity, including the inhibition of penicillin-resistant strains of *Staphylococcus aureus*, as well as antifungal properties [161, 162]. Xanthenes from *Calophyllum inophyllum*, particularly 6-deoxyjacareubin (56) and jacareubin (81) [163], and the benzophenones kolaniol (143) [102] and garcinol (145) [109] also exhibit antimicrobial properties.

A tuberculostatic effect has been noted in many natural and synthetic xanthenes [164]. A quantitative structure-activity relationship (QSAR) study has found a correlation

between ^{13}C NMR chemical shifts of C-4b and C-7 of various 1,3-oxygenated xanthenes and tuberculosis inhibition [165]. Three potent xanthenes were identified, of which gentsin (31) was the most active.

CONCLUSION

In conclusion, the Guttiferae produce a wide variety of both simple and prenylated xanthenes. The growing interest in these compounds is shown by the large number isolated during the last eight years. An attempt has been made in this review to correlate xanthone structure with the classical taxonomic divisions of the Guttiferae. The new results on xanthone biosynthesis support the idea of a common benzophenone precursor, although further studies are still required. Lastly, the pharmacology of xanthenes has been summarized, indicating their potential as medicinal agents.

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APPENDIX

Species-compound index

Xanthenes except (ital)-xantholignoid, { }-glycoside ()-benzophenone

<i>Allanblackia floribunda</i> Oliv	(132), (133)
<i>Archytea multiflora</i> Benth	51
<i>Bonnetia stricta</i> (Nees) Nees & Mart	4, 7, 21
<i>Calophyllum bracteatum</i> Thw	71
<i>C. calaba</i> var. <i>calaba</i> L	71-73
<i>C. thwaitesii</i> Pl. and Tr	70, 75
<i>C. walkeri</i> Wight	74, 76-78
<i>C. wightianum</i> T. Anders	117
<i>C. zeylanicum</i> Kosterm (= <i>C. lankaensis</i> Kosterm.)	1, 3, 4, 18, 56, 73, 81, 116
<i>Caraipa grandiflora</i> Mart	28, 36, (125)
<i>C. psidifolia</i> Ducke	2, 36
<i>C. vahl</i> Paula	36, (125)
<i>Clusia ellipticifolia</i> Cuatr	(138-140)
<i>Clusia nemorosa</i> G. F. W. Meyer	(148)
<i>Cratoxylum pruniflorum</i> Kurz	48, {120}, {121}
<i>Garcinia cambogia</i> Desr	(145), (147)
<i>G. densueta</i> Engl	88
<i>G. hanburyi</i> Hook. f	118, 119
<i>G. huillensis</i>	(145)
<i>G. indica</i> Choisy	4, (145), (147)
<i>G. kola</i> Heckel	(143)
<i>G. mangostana</i> L	48, 52, 54, 55, 68, 70, 73, 74, 79, 96-104, 108, {122}, (134)
<i>G. manni</i> Oliv	(144)
<i>G. nertosa</i> Miq	109
<i>G. ovalifolia</i> Oliv	86, (144), (146)
<i>G. pedunculata</i> Roxb	(135)
<i>G. polyantha</i> Oliv	90, (144), (146)
<i>G. pvrifera</i> Ridl	112-114
<i>G. quadrifaria</i> Baill. ex Pierre	69
<i>G. quaeqita</i> Kosterm	[151 a-c]
<i>G. staudtii</i> Engl	88, (144)
<i>G. thwaitesii</i> Pierre	39
<i>G. xanthochymus</i> Hook. f	3, 4, 12, 14, 15, 30, (134)
<i>G. xishuanbannanensis</i> Y. H. Li	(144), (146)

APPENDIX *Continued*

<i>Haploclathra leantha</i> (Benth.) Benth	15, 21, 35, 38, 42, 45, 46, 49
<i>H. paniculata</i> (Mart.) Benth	7, 15, 21, 23, 25, 29, 31, 33, 38, 40, 41
<i>H. verticillata</i> Ducke	4, 7
<i>Hypericum androsaemum</i> L.	10, 11, 27, 47, 48, 50, 80, 82, (124)
<i>H. aucheri</i> Jaub et Sprach	48, {120}
<i>H. balearicum</i> L.	1, 4, 11
<i>H. barbatum</i> Jack	{120}
<i>H. boissieri</i> Petrovic	{120}, {121}
<i>H. calycinum</i> L.	(124)
<i>H. canariensis</i> L.	1, 4, 5, 10, 26, 37, 63, (124), (126), (127)
<i>H. degeni</i> Bornm	31
<i>H. ericoides</i> L.	1, 4, 34, 43, (124)
<i>H. hirsutum</i>	{121}
<i>H. humifusum</i> L.	{120}
<i>H. japonicum</i> Thunb	{149}
<i>H. maculatum</i> Crantz	115, {120}, (124)
<i>H. mysorens</i>	1, 2, 4, 6, 7, 9, 24, (127)
<i>H. perforatum</i> L.	(124)
<i>H. rochei</i> Griseb. and Schenk	{120}, {121}
<i>H. rumeliacum</i> Boiss	{120}
<i>H. sampsonii</i> Hance	26, 64, 82, {120}, {121}
<i>Kielmeyera coriacea</i> Mart.	(128)
<i>K. rubiflora</i> Camb	26
<i>K. speciosa</i> St. Hill	26
<i>Mahurea tomentosa</i> Ducke	4, 7
<i>Mesua ferrea</i> L.	42
<i>Platonia insignis</i> Mart.	32, {123}
<i>Psorospermum febrifugum</i> Sprach	11, 44, 53, 57–61
<i>Rheedia benthamiana</i> Pl. and Tr	86, 88, 89, 91
<i>R. brasiliensis</i> (Mart.) Pl. and Tr	62, 67, 86–94
<i>R. gardneriana</i> Pl. and Tr	62, 67, 84, 86–91
<i>R. madruno</i> (HBK) Pl.	(144)
<i>Symphonia globulifera</i> L.	(134)
<i>Tovomita brasiliensis</i> Walp	36
<i>T. excelsa</i> Andrade-Lima & G. Maritz	16, 18, 20, 22
<i>T. mangle</i> G. Maritz	95, (141), (142)
<i>Vismia decipiens</i> Schlecht-Cham	(136), (138), (139)
<i>V. guaramirangae</i> Huber	1, 2, 4, 7, 8, 11, 13, 15, 17, 19, (124–126), (136), (137)
<i>V. guineensis</i> (L.) Choisy	83, 85, 110, 111

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