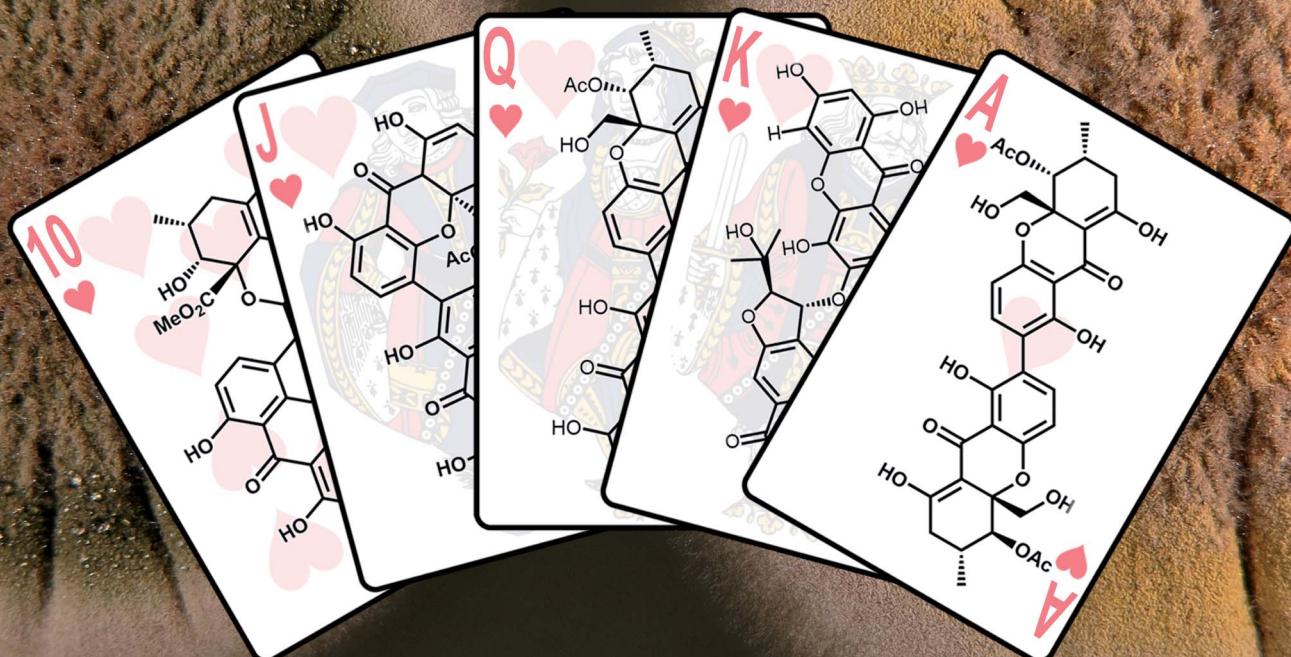


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Stefan Bräse, Kye-Simeon Masters *et al.*

Xanthone dimers: a compound family which is both common and privileged



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Xanthone dimers: a compound family which is both common and privileged†

Tim Wezeman,^a Stefan Bräse^{*ab} and Kye-Simeon Masters^{*c}

Covering: up to 2014

Xanthone dimers are a widespread, structurally-diverse family of natural products frequently found in plants, fungi and lichens. They feature an intriguing variety of linkages between the component xanthones (benzannulated chromanones). These synthetically elusive secondary metabolites are of great interest due to their broad array of bioactivities, which has led to the xanthones being designated as 'privileged structures'. We seek herein to give an overview of all reliably-described xanthone dimers, their structures, occurrence, and the bioactivities established to date. The possible biosynthetic pathways leading to members of this family are also discussed in light of our current knowledge.

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^aInstitute of Organic Chemistry (IOC), Karlsruhe Institute of Technology (KIT), Fritz-Haber-Weg 6, 76131 Karlsruhe, Germany. E-mail: stefan.braese@kit.edu; Fax: (+49) 721 6084 8581

^bInstitute of Toxicology and Genetics (ITG), Karlsruhe Institute of Technology (KIT), Hermann-von-Helmholtz-Platz 1, D-76344 Eggenstein-Leopoldshafen, Germany

^cDiscipline of Nanotechnology and Molecular Sciences, School of Chemistry, Physics and Mechanical Engineering, Faculty of Science and Engineering, Queensland University of Technology (QUT), GPO Box 2434, Brisbane, Queensland, 4001, Australia. E-mail: kye.masters@qut.edu.au

† Electronic supplementary information (ESI) available: In order to provide a clear overview of the structural features, natural sources and biological activities of the discussed xanthones, three tables detailing these topics have been added as ESI. See DOI: 10.1039/c4np00050a.

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1 Introduction

Xanthones comprise a group of structurally diverse, biosynthetically intriguing, biologically active and synthetically challenging natural products. Many xanthones have been found to exhibit pronounced biological activities, for example anti-tumour effects.^{1–5} Xanthone chemistry is satisfactorily rich, with the conjugated donor–acceptor motif of the central B-ring

ensuring that these compounds display a degree of personality greater than their apparently simple core structure might suggest.

The xanthone family has been studied for over a century since the early 1900s^{6,7} and has since been investigated and reviewed, particularly for the monomeric species and plant-derived species.⁸⁻¹⁴ Several reviews on xanthones have been published in recent years, covering xanthones from fungi,¹⁵ lichens and bacteria,¹⁶ structure-activity relationships,² anti-malarial properties of xanthones,¹⁷ biological activity,¹⁸ chemical synthesis of xanthone cores,¹⁹⁻²¹ and their biosynthesis.²²⁻²⁷ Nonetheless, the field of xanthone chemistry is growing rapidly; every few years see the publication of an update on novel naturally-occurring xanthones.^{8,21,28-30} Analytical techniques have been optimised specifically for xanthones, including crystallography³¹ and chromatographic methods.³² The



Kye-Simeon obtained his Doctorate with Prof. Bernard Flynn at Monash University (2007). After a year in Industry, he had a postdoctoral role with Prof. Bert Maes at the University of Antwerp (2008–2009). An Alexander von Humboldt Fellowship with Prof. Stefan Bräse at the Karlsruhe Institute of Technology (2010–2011) led to his interest in xanthones. Following a Michael J. Fox Fellowship at

Lundbeck, Copenhagen (2012) he returned to Australia, where he is currently a Vice Chancellors Fellow at the Queensland University of Technology. Activities in his newly-created research group focus on the discovery of synthetic methodology and natural product synthesis.



Tim Wezeman is a PhD candidate in the Institute of Organic Chemistry at the Karlsruhe Institute of Technology, under the supervision of Prof. Stefan Bräse. His current research interests involve heterocyclic chemistry and method development, as well as solid phase organic synthesis and natural product synthesis. He received his BSc (2010) and MSc (2012) in Molecular Life Sciences at the

Wageningen University in the Netherlands after a research stay with Prof. Ciufolini at the University of British Columbia in Vancouver.

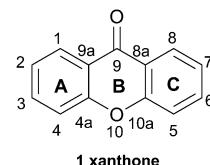


Fig. 1 Xanthone monomer core with numbering.

structure, activity and synthesis of polycyclic, yet monomeric xanthones has recently been reviewed,³³ as well as the specific class of the caged xanthones.^{34,35} Most of these articles focus mainly on monomeric xanthones, and to date none have focused on dimeric and trimeric xanthones.

For the sake of simplicity and easy comparison between structural features, structural numbering will follow the monomeric parent xanthone core (1, Fig. 1) throughout this review.

2 Biosynthesis

The biosynthesis of dimeric xanthones is a topic which stands in need of further research progress, especially the identification and description of the key dimerisation processes (see also ref. 15 and 16). As the biosynthesis of the dimeric structures is a point of considerable interest and remains largely unknown, we will give a necessarily brief outline of the achievements so far – the full elucidation of the biosynthesis is no doubt just over the horizon.

2.1 Monomer xanthones

It has been known for some time that the biosynthesis of the monomeric xanthone core proceeds differently in fungi^{25,36} and higher plants (Scheme 1).^{37,38} The synthesis of the xanthones in fungi has been studied in detail since the radiolabelled acetate



Stefan Bräse studied in Göttingen (Germany), Bangor (UK) and Marseille (France) and received his Ph.D. in 1995 with Armin de Meijere (Göttingen). Post-doctoral appointments at Uppsala University (Jan E. Bäckvall) and the Scripps Research Institute (K. C. Nicolaou) were followed by habilitation associated with Dieter Enders at RWTH Aachen in 1997. He became professor for organic chemistry at the University of Bonn in 2001. He has been full professor at the Karlsruhe Institute of Technology since 2003, and director of the Institute of Toxicology and Genetics since 2012. His research interests include drug-discovery and -delivery, combinatorial synthesis of bioactive compounds, total synthesis and nanotechnology.



feeding experiments of Birch in the 1950s, showing that polyketides are the biosynthetic precursors of the xanthone core for fungi. More recently, Simpson has reported findings resulting from the sequencing of the *Aspergillus nidulans* genome.³⁹ The sequencing revealed the presence of 32 clusters containing polyketide synthase (PKS) genes, which were studied by gene deletion to reveal information about the biosynthetic pathway which converts emodin *via* key steps of oxidative ring-scission and decarboxylative cyclisation to xanthones and eventually prenylated and cyclised products such as shamixanthone, tajixanthone and sterigmatocystin. As part of a comprehensive study on the formation of aflatoxin in *Aspergillus* species, Townsend and co-workers described the generation of xanthone monomers through a complex sequence of epoxidation, rearrangement, deoxygenation, Baeyer–Villiger oxidation, and further deoxygenation leading to the tricyclic xanthone core of sterigmatocystin.⁴⁰ Interpretation of the sequences may differ markedly⁴¹ in terms of the sequence of events, and the structural homology of biomolecules involved in the pathway in differing species may not, in fact, be a reliable method for determining correlations between pathways.

Müller and co-workers have recently discussed the importance of reductive steps in the synthesis of xanthones commencing from (polyketide-derived) anthraquinone, emodin, and passing through chrysophanol.⁴² Contrastingly, the biosynthesis of xanthones in plants results from a convergent synthesis of polyketide and shikimic acid pathways, whereby 3-hydroxybenzoyl-CoA (obtained from an early shikimate pathway intermediate) is condensed with a polyketide, which may be then cyclised to form the C-ring and then the B-ring.²⁶ A consequence of these two distinct pathways, both of which pass through a freely-rotatable benzophenone intermediate, is that the pattern of oxidation between xanthones from plants and fungi frequently differs. Fungal xanthones (or their biosynthetic precursors) almost ubiquitously display C-1 and -8-hydroxylation, whilst plants frequently display C-1, -3 and -5 or -7-hydroxylation.

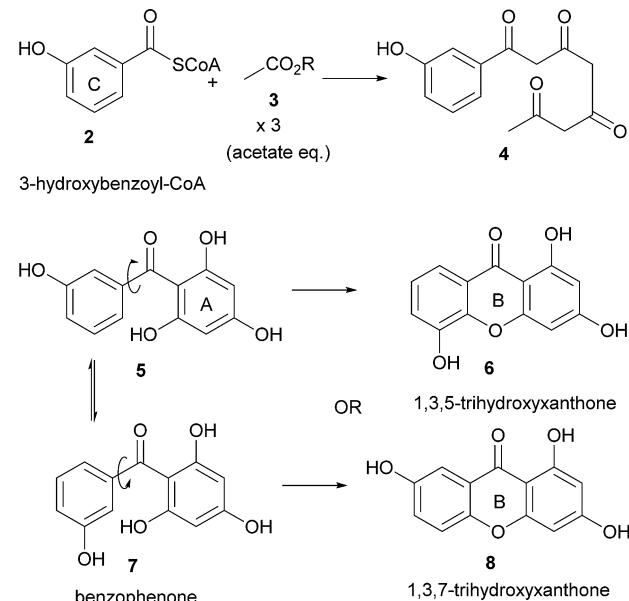
In terms of further diversification of the xanthone core, some processes are now understood better than others. The investigations of Elix and co-workers have led to a fuller understanding of some of the pathways of further structural elaboration of the xanthone nucleus in lichens. Particularly, cladistical/phylogenetic analysis of related species was studied to develop a fuller understanding of the relationship between enzymes responsible for methylation and chlorination.^{43,44} Prenylation of the dimers is frequently found in plant-derived xanthones.⁴⁵ Pendant sugars are also found as a structural feature of dimers, see for example puniceaside C (72, plant derived) and Hirtus-neanoside (120, lichenoid). It is also known that some substituents are removed from the xanthone core, such as the reduction of hydroxyl functions and reductive dearomatisation.

2.2 Xanthone dimerisation

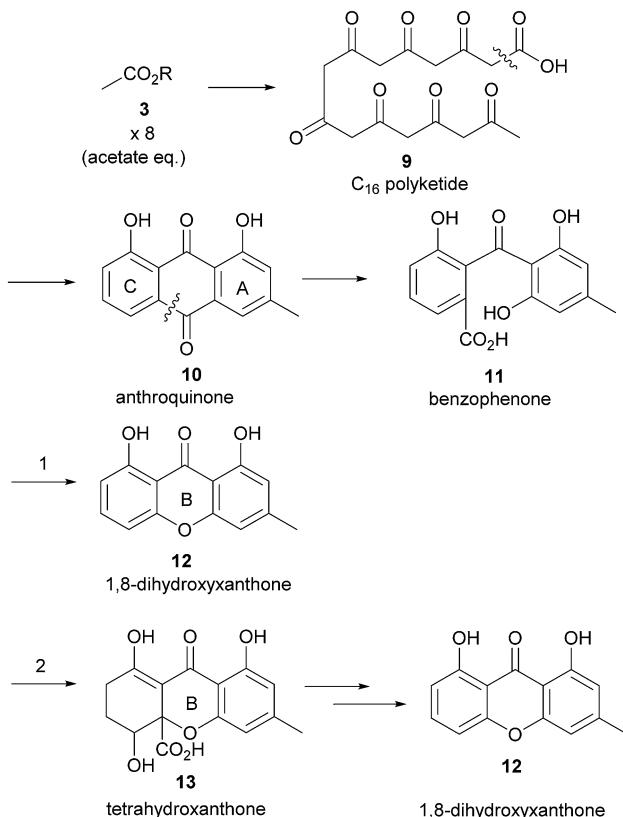
There are a variety of ways in which dimeric and trimeric xanthones can be linked: (i) a rotatable or atropisomeric biaryl C–C bond or (ii) a biaryl ether C–O–C linkage. Also, because

xanthones, particularly from plants, can be decorated with prenyl groups, the linkages are also often in the form of (iii) aryl–O–alkyl linkages, or (iv) prenyl derivative–prenyl derivative linkages. The incorporation of nitrogenous bases to form polyaromatic antibiotic xanthones is also frequently observed,

Xanthone Synthesis in Plants



Xanthone Synthesis in Fungi and Lichens



Scheme 1 Xanthone biosynthesis differs in plants and fungi.



and readers are directed to the recent review by Porco and co-workers on this subject.³³

One of the more common ways in which dimerisation is found in the xanthones is through the biaryl linkage. Regio- and stereo-selective biaryl linkage of phenolic compounds is common in plants, bacteria, lichen, and fungi – as Bringmann and co-workers noted in their seminal work on natural biaryl linkages, “wherever in Nature phenolic aromatics can be found... the corresponding homo- or hetero-dimeric biaryls have to be expected”.⁴⁶ Certainly, the discovery of the blennolides (hemisecalonic acids)⁴⁷ in 2008 supported the putative dimerisation of discrete monomers as the likely method of dimer formation, rather than a more complex and difficult to imagine tandem biosynthesis pathway – the side-by-side cyclisation of a double-length polyketide to form *e.g.* a secalonic acid seems unlikely. Despite efforts in the detailed study of the biosynthesis of dimeric xanthones, particularly of the secalonic acids by Frank and co-workers,^{23,40,48–50} the dimerisation of xanthones remains a tricky subject for both the synthetic and biosynthetic chemist – the extent of the involvement of enzymes, and their nature, is at present unclear. There has been, at the time of writing, no direct observation of enzymatic dimerisation of the monomer units.

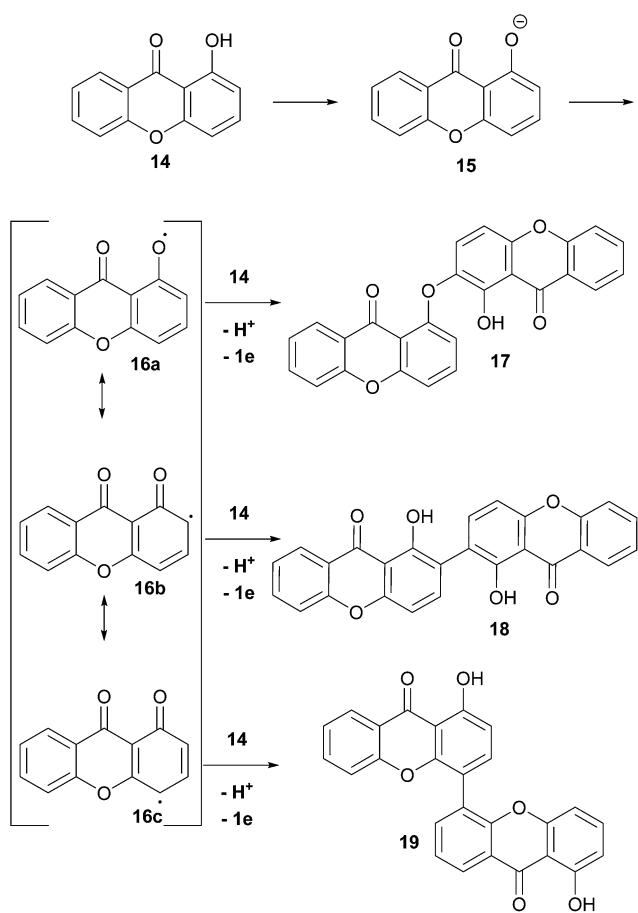
Enzymatically-mediated or not, the oxidative pathway which is most commonly invoked is shown here (Scheme 2), whereby single electron transfer deprotonation of a hydroxyxanthone, **14**, leads to an intermediate which can be readily oxidised by single-electron-transfer to xanthonyl radical **16**. Resonance contributors of the delocalised aryl radical, **16a–c**, can then couple to electron-donors, notably the *ortho* and *para*-C positions.

2.3 Regiochemical evidence for enzyme dimerisation

The resulting biaryl bond linkages are formed which, in the absence of an enzyme or other structurally-influencing substrate, is determined largely by the spin-density distribution of the radical generated, and presumably in some cases other inherent stereoelectronic characteristics of the substrate molecule. As Kozlowski has put it, in the absence of external factors, “the substrate typically dictates the available coupling products”.⁵¹ This leads us, as chemists, to an interesting corollary – if there are differing regio- and atropo-isomeric biaryl-linked dimers of the same substrate to be found in Nature, then it must be the case that there are external forces at play in their biosynthesis to lead to this divergence, rather than ‘spontaneous’ oxidative dimerisation with oxidants such as intracellular O₂. Prominent examples supporting this line of thinking can be seen in Scheme 3, where both secalonic acid B/chrysoxanthone and secalonic acid E/talaroxanthone are presumably generated with alternative (chemo- and regio-) selectivity from the monomer units, hemisecalonic acids B and F, respectively. The oxidative dimerisation of the xanthones is thus most likely mediated by enzymes, as shown for a number of model compounds.^{52–54} It can be mimicked by chemical oxidases.⁵⁵ In some cases, the dimerisation led to the formation of diaryl ethers (*e.g.* chrysoxanthone **22**, Scheme 3) instead of to a C–C bond formation.

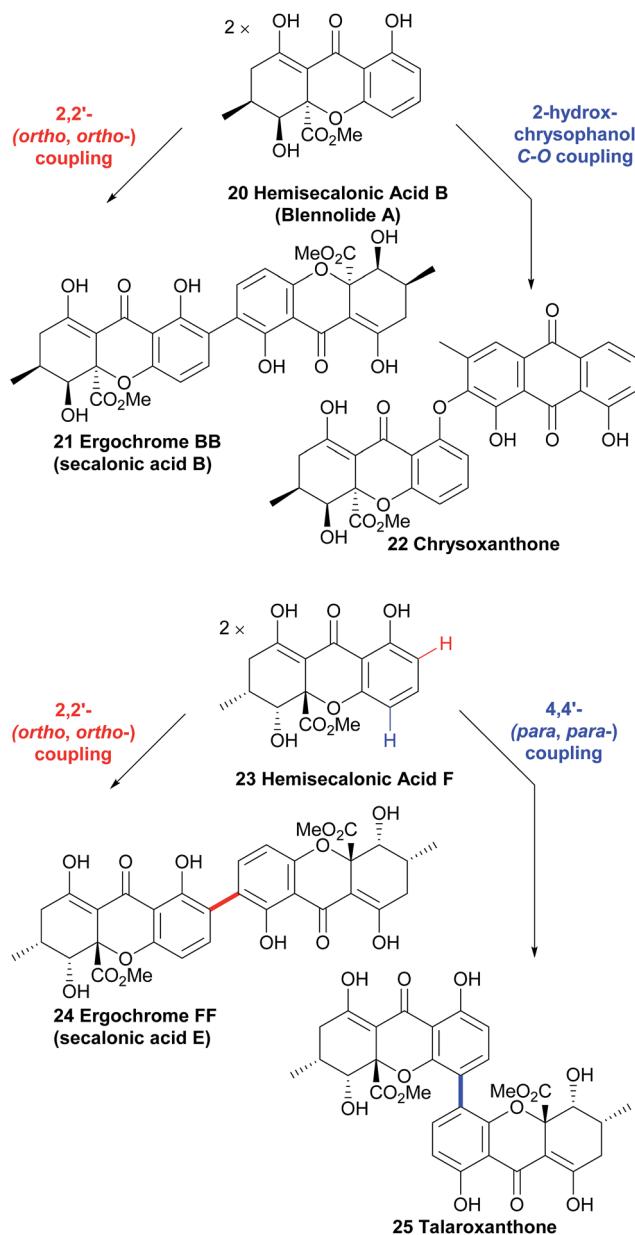
Hemisecalonic acid B⁴⁷ is dimerised at the *ortho*-positions to form secalonic acid B **21** (Scheme 3; see below for more information) in a variety of fungal species.^{49,148–150,153} From the same substrate in the case of the xanthone anthraquinone chrysoxanthone, the coupling of the hemisecalonic acid B with a 2-hydroxy variant of chrysophanol, a widespread polyketide natural product, dimerisation leads to the C–O bond formation of diaryl ethers *i.e.* chrysoxanthone.⁵⁶ Another example of biosynthetic divergence is likely involved in the regioisomeric selectivity which leads to the formation of talaroxanthone **25**, a 4,4'-linked xanthone dimer (from the endophytic fungi *Talaromyces* sp.)⁵⁷ and secalonic acid E **24**, a 2,2'-linked xanthone dimer (from *e.g.* *Phoma terrestris*).^{58,59} That a statistical mixture of the three regiosomers is produced by each of the organisms in question and the two differing regiosomers have been selectively isolated by the research groups in each case is highly unlikely (see Scheme 3). The selective discovery of these regiosomers can therefore be considered indirect evidence for the active role of yet-to-be discovered enzymes.

In the laboratory, regioisomeric control over the biaryl bond formation^{60–62} is often difficult to impose, and frequently requires innovative solutions.^{63–66} In natural biosynthesis, it has



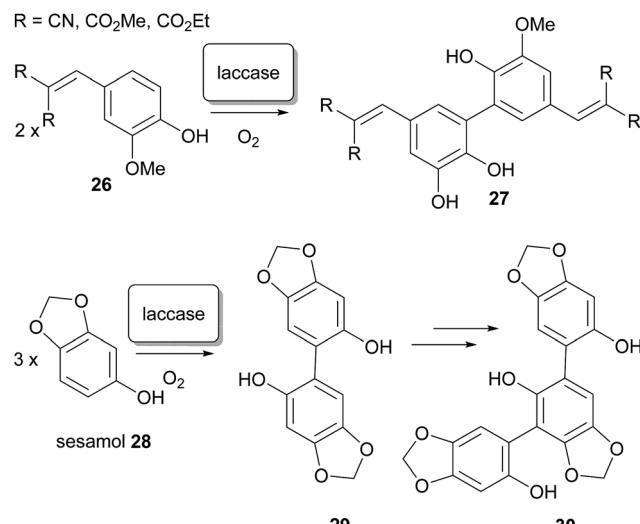
Scheme 2 Phenolic dimerisation reactions adapted for xanthones. For clarity, coupling of **16a–c** at the *ortho*-position of **14** is shown.





Scheme 3 Multiple pathways from the same monomer units strongly suggest the involvement of enzymatic processes in xanthone dimerisation.

been demonstrated that selectivity between 2,2'- 2,4'- and 4,4'- coupling can presumably be imposed by enzymes, although little direct evidence exists for this – no enzyme has yet been identified which fulfils the role of regioselective differentiation in the dimerisation of xanthones in Nature. Nonetheless, high selectivity for such processes has been demonstrated with enzymes in the laboratory; oxidative dimerisation at a specific *ortho*-position to form ferulic acid derivatives has been reported by Beifuss and co-workers, and has been achieved by application of the (single electron) oxidase enzyme^{67,68} laccase,⁶⁹ from *Trametes versicolor*, in the presence of O₂.⁷⁰ Additionally, selective coupling at the C-6 position of sesamol led to dimer intermediates and then trimers in good yield.⁷¹ The outcome was



Scheme 4 The synthetic usage of laccase enzymes in biphenolic coupling to biaryl-linked dimers.

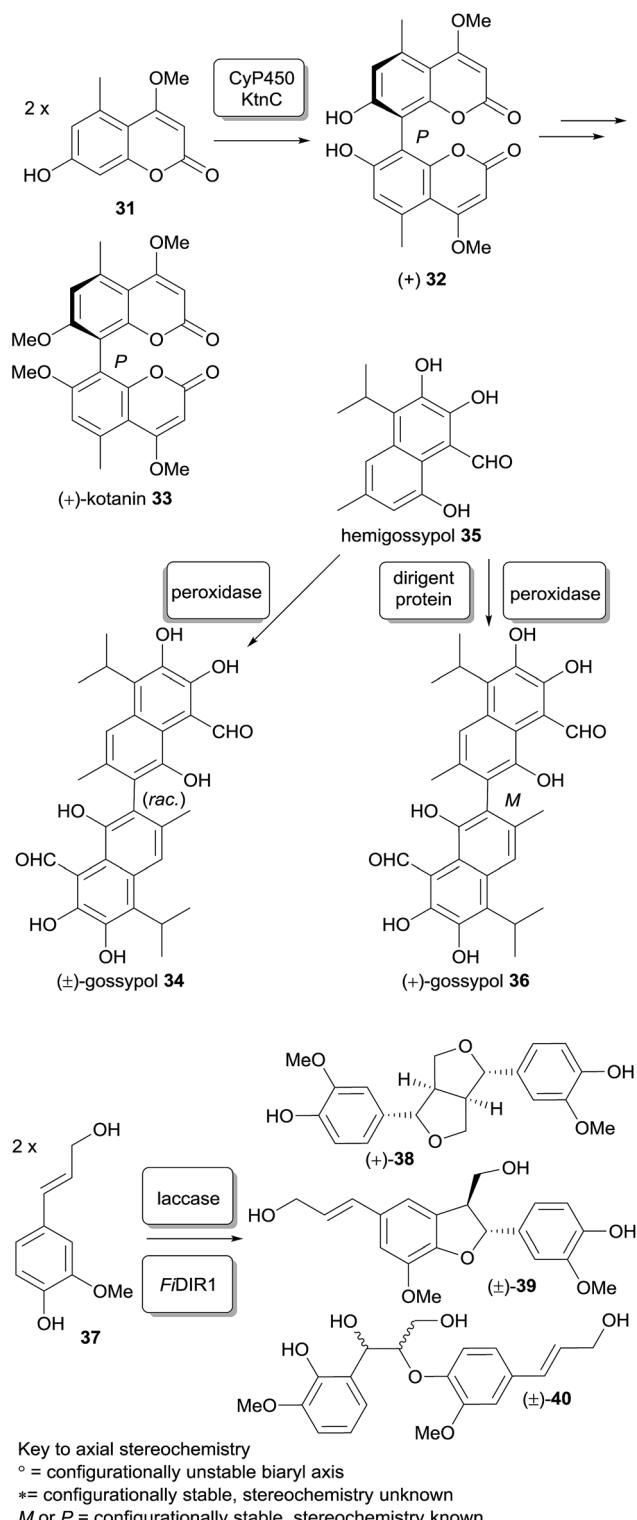
markedly different than in the application of laccase from *Agaricus bisporus*, and also from electrochemical non-enzymatic oxidative coupling of this substrate by the method of Waldvogel and co-workers.^{72,73} Aside from their potential relevance to xanthone dimerisation, the synthetic application of enzyme-catalyzed transformations has significant additional benefits: they can be performed in aqueous solvent systems and under mild reaction conditions, a remarkably broad substrate spectrum that can be expanded even more by using mediators (Scheme 4).⁷⁴

It is therefore logical to support the assertion of Bringmann and co-workers that the dimerisation of phenolic aromatics (both C–C and C–O coupling) is through “ – mostly – enzymatically-assisted biogenetic pathways”.⁴¹ Based on related processes in the chemical literature on both natural products^{75–79} and synthesis,^{80–82} one possibility is that a laccase is responsible. Another possibility is a monooxygenase, such as cytochrome P450; evidence for a likely mechanism for stereoselectivity is also found in this case.

2.4 Stereochemical features of the biaryl xanthone dimer

Substituted biaryls exhibit varying degrees of barrier to rotation, ranging from freely rotating (microsecond time-frame) through slow room-temperature interconversion to atropisomers which may be isolable and thereafter stable; dimeric xanthones show this feature also. Stereochemical biaryl bond formation is displayed in the secondary metabolism of bacteria, fungi, lichens and plants, and has herein been denoted alongside the structures of biaryl-linked dimers utilising the notation of Bringmann and co-workers (see Scheme 5).⁴⁴ As both chiral and *meso*-dimers have been found in the secalonic acid series, various mechanisms for the origin of the dimers can be invoked. The reasons for this could certainly be due to the different biosynthesis routes employed between the classes of species. In fungi, all xanthone dimers display a biaryl xanthan–xanthone bond between one or more di- or tetra-hydro xanthones. It could





Scheme 5 Examples of stereoselective phenolic coupling with enzymes *in vitro*.

possibly be argued that the dimerisation process could be influenced by the asymmetrically-substituted C-ring, a possible explanation for non-enzymatic axial chirality generation. However, the fully aromatic xanthone dimer ploiarixanthone 68, which is optically active ($[\alpha]_D^{25} = +23^\circ$),⁸³ must involve axial-

chirality influencing biomolecules in the stereospecific dimerisation step.

The involvement of stereoselectivity-mediated proteins may prove to be the origin of stereospecific pathways in the formation of dimeric xanthones. One case where the explanation for the regio- and stereo-chemical control observed in a biphenolic coupling has been uncovered is the mechanism of selective phenol coupling in *Aspergillus niger* by Müller and co-workers.⁵⁴ They showed that an oxidative phenol coupling is also the key stereo-divergent step in the formation of *P*-(+)-kotanin (33, Scheme 5) from two equivalents of coumarin 31 catalyzed by cytochrome P450 monooxygenase KtnC, which was identified by targeted gene deletion in the *A. niger* strain. The facially-selective interaction of the two monomer subunits with the heme-containing active site of the CyP450 was modelled *in silico* by homology studies and docking of both the substrates and product. Stipanovic and co-workers showed enzymatic coupling of hemigossypol (35, Scheme 5); this compound by *in vitro* application of a pure peroxidase only led to a racemic product (±)-gossypol, whereas the addition of a cotton-plant dirigent protein gave predominantly (+)-(S)-gossypol.^{84a} Naturally-occurring gossypol is found as a mixture of (+) and (−) isomers, which have been found to exist in a continuum of ratios between 68 and 2% (−)-(R)-gossypol, depending on the strain of the cotton plant *G. barbadense*,^{84b} possibly indicating a variation in the type or extent of dirigent protein control in the coupling. Stereoselective phenol coupling of 37 resulting in C and O-alkylation products (+)-38 or racemic 39 and 40, reminiscent of the kind of transformations of some linked and/or cyclized dimeric xanthones, has also been achieved with a combination of laccase and dirigent protein FiDIR1.⁸⁵ It may be the case that for xanthones there are also mediator proteins associated with oxidase enzymes involved in the stereoselective generation of the biaryl bonds.

The absolute stereochemical assignation of regioisomerically identical xanthone dimers isolated from different biological sources is, unfortunately, in many cases incomplete. X-ray structures have been used to determine the relative and absolute stereochemistry of the stereochemical centres and axes (e.g. for phomoxanthone A 106 and phomalevone A 103).⁸⁶ In some cases, however, modern DFT calculations have allowed CD spectra to be computed which were used for comparison with experimental data (blennolides).⁴⁷ In particular, the quantitative consideration of vibronic effects, conformer equilibria and solvent effects were included in most of the models.⁸⁷ As biological activity and their biological targets are different for different diastereomers (e.g. compare secalonic acid A and E), it is very important to address stereochemical issues of dimeric xanthones. Unfortunately, many data are not available (due to the lack of instrumentation and/or authentic material).

2.5 The true biological sources of xanthone dimers?

In some cases researchers have reported dimeric xanthones as being isolated from endophytes.^{57,86,88–90} Certainly culture broth production of dimeric xanthones has demonstrated that this class of molecules can be produced by fungi alone. This raises the question for 'plant-derived' compounds: are the plants really



the true producers, or fungi? Is there a mixed origin, or is it really only the (undetected) fungi which produce the xanthone dimers?

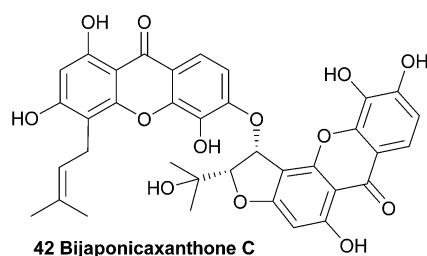
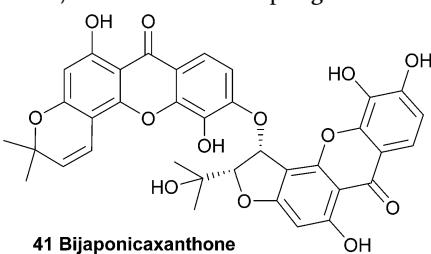
For monomeric xanthones, the plant-fungus contributions have been at least partially unveiled on at least two occasions. In its biosynthesis, the monomeric xanthone 2,7-dichlorolichexanthone (not shown) was isolated from samples of the lichen *Lecanora dispersa*. In cultures of the fungal species in the absence of the alga, the xanthone biosynthesis was halted.⁹¹ In an example for xanthone dimers, the eumitrins (see below) are produced by the lichen (algae-fungal symbiotic organism), and not by only the fungus in isolation.⁹² It is conceivable that in some situations that involve plant-microbe combinations, the biosynthesis may even involve a to-and-fro sequence of enzymatic transformations carried out by each organism. Such a scenario, if it does exist, is likely to be complex and case-specific.

3 Xanthone dimers from plants

Xanthone dimers from plants, like the monomers themselves, commonly feature prenylated and multi-prenylated cores. The presence of these many prenylated monomeric xanthones shows that prenylation occurs readily prior to dimerisation, although there is no evidence to suggest that it does not also occur post-dimerisation.

3.1 Bijaponicaxanthones

Several interesting xanthones were isolated from the ethanol extracts of the dried aerial parts of *Hypericum japonicum*, a Chinese medicinal plant, including one dimeric xanthone named bijaponicaxanthone (41). By comparing the NMR spectra with known compounds such as the monomeric xanthone isojacareubin (not shown), and by analysing additional 2D NMR spectra, its structure could be assigned. The relative configuration between the two chiral centres was determined to be *cis* for the two protons, based on their coupling constant.^{93–95}

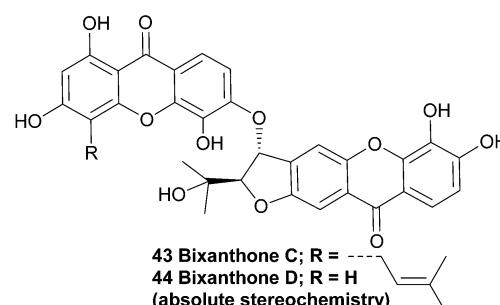


In 2005, a similar dimeric xanthone was isolated from *Hypericum japonicum*. After detailed structural analysis the

xanthone was found to be prenylated instead of having a fourth ring and the natural product was named bijaponicaxanthone C (42).⁹⁶ Bijaponicaxanthone C was also reported to have been isolated from the roots of the *Hypericum riparium* plant.⁹⁷

3.2 Bixanthones

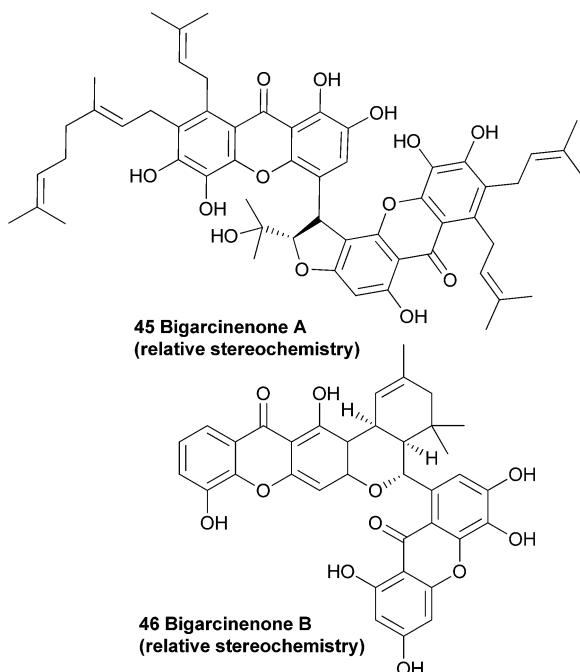
The structurally very similar bixanthones C and D (43, 44), as reported in a Chinese patent, were also isolated from *Hypericum japonicum*. The patent claims these compounds are active ingredients in a traditional Chinese medicine which is used to treat a variety of liver diseases, and describes the use of the flavone component of that plant species as a medicament for treating hepatic fibrosis.⁹⁸ These compounds are missing OH groups on the furannulated xanthone component, presumably due to enzymatic reduction.



3.3 Bigarcinenones

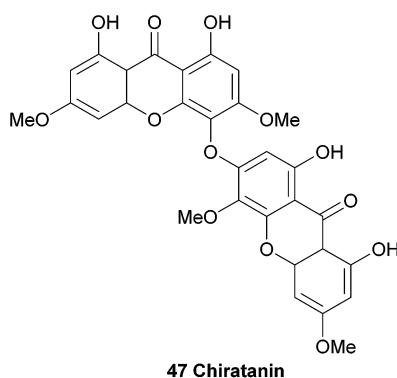
Since the plant *Garcinia xanthochymus* is widely used as traditional Chinese medicine, Yang *et al.* investigated the extracts from the bark of the plant. They observed a strong antioxidant activity (IC_{50} : 4.6 μ g mL⁻¹, as determined by a 1,1-diphenyl-2-picrylhydrazyl (DPPH) radical scavenging bioassay) in the ethyl acetate soluble fraction from the ethanol extracts, and proceeded with a more detailed study on the contents of these extracts. Along with seven known xanthones, they found a novel bisxanthone called bigarcinenone A (45).

Using mainly 2D-NMR techniques, the authors identified the dimeric xanthone to be coupled through a side-chain of one of the two subunits. Bigarcinenone A performed relatively well in a DPPH radical scavenging activity assay (IC_{50} : 9.2 μ g mL⁻¹), outperforming all other isolated xanthones from this extract (IC_{50} : 16.3–250 μ g mL⁻¹) as well as BHT (IC_{50} : 20.0 μ g mL⁻¹), a well-known synthetic antioxidant.⁹⁹ In 2011, bigarcinenone B was isolated from the bark of the same plant and its structure and relative stereochemistry was also elucidated using mainly 2D NMR techniques such as HMBC, HSQC and ROESY. A possible biosynthetic pathway was proposed by the authors that would lead to the unique connection through the two six membered rings. A Diels–Alder reaction between the two prenyl groups of the monomers would lead to the cyclohexene derivative, which would undergo a second cyclisation to form the final ring. The antioxidant activity of bigarcinenone B (7) was tested with a DPPH assay (IC_{50} : 20.14 μ M *versus* 13.16 μ M for ascorbic acid) and a H₂O₂ assay (IC_{50} : 2.85 μ M *versus* 0.76 μ M for ascorbic acid).¹⁰⁰



3.4 Chiratanin

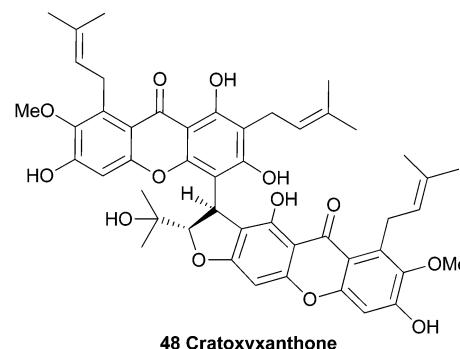
An interesting biaryl-ether linked dimeric xanthone named chiratanin (47) was isolated together with several monomeric xanthones and other secondary metabolites from the benzene extracts of the plant *Swertia chirata*.¹⁰¹ At the time, this was the first report of the occurrence of a dimeric xanthone in a higher plant. After detailed analysis of the NMR spectra, the authors identified the positions of the oxygen atoms in the molecule, and with D₂O exchange experiments all the phenolic hydroxyl groups were identified. However, the actual connection of the two xanthones remained difficult to determine, and three possible structures were identified. The molecular asymmetry was instructive in determining the final structure, which they named chiratanin.



3.5 Cratoxyxanthone

Several natural products were isolated from the bark of *Cra-toxylum cochinchinense* in 1995 by Sim *et al.*¹⁰² After extensive HPLC purification of the isolated compounds, detailed high

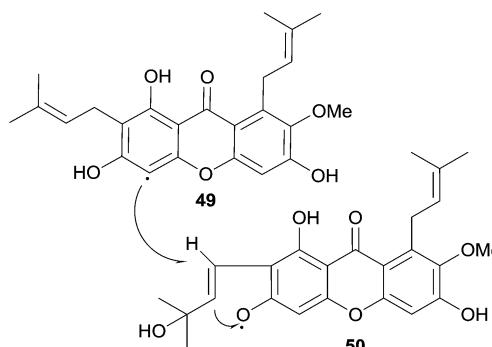
field NMR spectroscopy was used to identify a novel dimeric xanthone, named cratoxyxanthone (48).



The authors speculated on a biosynthetic pathway involving the coupling of two mangostin-derived radicals, as depicted in Scheme 6. Cratoxyxanthone was also isolated from the chloroform soluble extracts of the stem bark of *Garcinia mangostana* in 2009.¹⁰³ In this study, it was found that cratoxyxanthone had a very poor *in vitro* activity in cytotoxicity (HT-29 cell line) and ELISA NF- κ B (p65 and p50) assays.

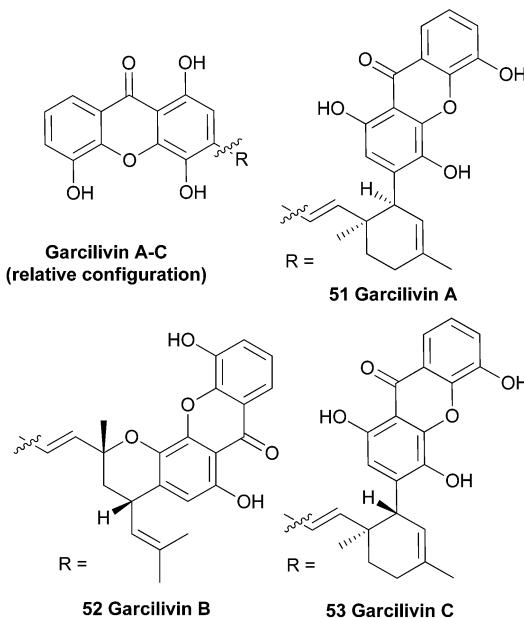
3.6 Garcilivins A-C

Bark from the roots of the South African plant *Garcinia livingstonei* was phytochemically investigated in a larger study aiming to find reversible monoamine-oxidase inhibitors for antidepressant drugs. This study resulted in the discovery of several prenylated xanthones, including three dimeric xanthones named the garcilivins (51–53).¹⁰⁴ The structures were elucidated using extensive mass and NMR spectroscopy and by comparing with monomeric xanthone spectra.¹⁰⁵ In 1992 it was reported that these xanthones were currently being assayed for their biological activities, but as far as the authors know, no follow-up paper was published until Pieters *et al.* in 2006.¹⁰⁶ They report the isolation of several xanthones and flavonoids from a Tanzanian *G. livingstonei* and tested the isolated compounds for their antiparasitic activity and cytotoxicity. Garcilivin A and C showed a very interesting difference in their toxicity tests, considering they are diastereoisomers. Both compounds were tested against four parasites and garcilivin A



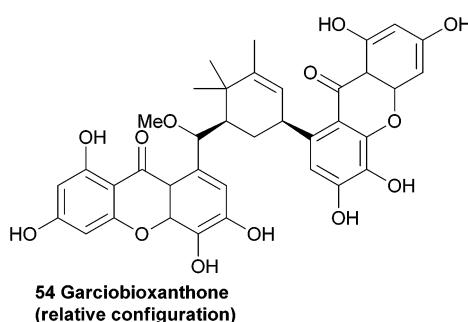
Scheme 6 Proposed biosynthetic coupling of two mangostin-derived radicals to form cratoxyxanthone.

showed a very strong non-selective activity in all the assays, whereas garciliavin C only showed significant activity against *Trypanosoma brucei*. Especially noteworthy is the score for cytotoxicity against MRC-5 cells of 2.0 μM for garciliavin A and 52.3 μM for garciliavin C.



3.7 Garciobioxanthone

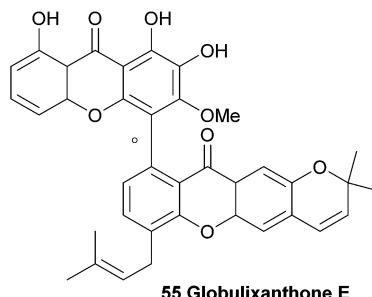
Recently, the novel garciobioxanthone (54) was isolated from the ethanol extracts from the bark of *Garcinia oblongifolia*, along with several known compounds. Using 2D NMR techniques the relative configuration and structure was determined.¹⁰⁷



3.8 Globulixanthone E

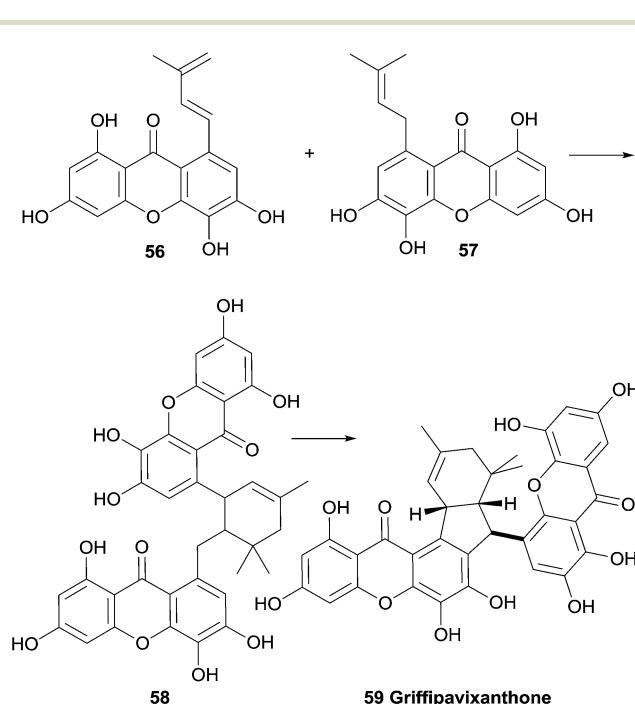
Three prenylated xanthones were isolated from the root bark of the large forest tree *Sympomia globulifera* and named globulixanthone C-E. The tree is widely used in Cameroon as a medicinal plant and laxative for pregnant women. Globulixanthone C and D were found to be monomeric xanthones, but globulixanthone E was found to be a bisxanthone and its structure (55) was elucidated using ESI-TOF MS, IR and NMR spectroscopic methods. Using NOESY spectra and by comparing the obtained spectra with the other isolated xanthones, the structure was determined. The three xanthones

were tested for their *in vitro* antimicrobial activities against *Staphylococcus aureus*, *Bacillus subtilis*, *Vibrio anguillarum* and *Escherichia coli* in an agar well diffusion assay. The monomeric xanthones showed a moderate activity to *S. aureus* (MIC: 8.05–14.05 $\mu\text{g mL}^{-1}$) and *B. subtilis* (MIC: 8.24–12.5 $\mu\text{g mL}^{-1}$), but globulixanthone E showed a remarkably high activity against all tested organisms (MIC: 5.56–3.12 $\mu\text{g mL}^{-1}$) except *E. coli*, and outperformed the positive control in the case of *S. aureus* (MIC: 4.51 $\mu\text{g mL}^{-1}$ versus 6.25 $\mu\text{g mL}^{-1}$ for streptomycin sulfate).¹⁰⁸



3.9 Griffipavixanthone

An investigation into the secondary metabolites of the Malaysian plants *Garcinia pavifolia* and *G. griffithi* lead to the discovery of a new bisxanthone named griffipavixanthone.¹⁰⁹ Using advanced spectroscopic methods, including NOESY, HMBC and INEPT NMR studies, the relative structure was elucidated. In a later study, the bisxanthone was also found in *G. maingayii*.¹¹⁰ Griffipavixanthone was the first example of a dimeric xanthone that is connected *via* a 5 and a 6 membered ring.¹⁰⁹ The authors suggest a possible biosynthetic pathway involving an initial Diels–Alder reaction of two prenyl groups on

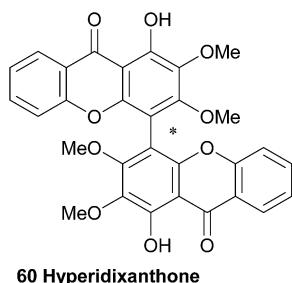


Scheme 7 Proposed biosynthetic pathway.

the two xanthones (Scheme 7). The resulting cyclohexene is then followed by another cyclisation, either ionic or radical, to yield the 5-membered ring. The bisxanthone was also isolated from *Garcinia oblongifolia*¹⁰⁷ (along with several other xanthones, such as garciobioxanthone) and *G. virgate*¹¹¹ (along with two new monomeric xanthones named virgataxanthone A and B, not shown). The isolated compounds in the latter study were tested for antioxidant capacity, and it was noted that griffipavixanthone showed a notably high radical scavenging ability, with an EC₅₀ lower than the references used in the study (EC₅₀: griffipavixanthone: 11.5 μg 100 mL^{-1} , 2,6-di-*tert*-butyl-4-hydroxy-anisol: 13.6 μg 100 mL^{-1} , α -tocopherol: 13.8 μg 100 mL^{-1}). In *in vitro* cell line cytotoxicity tests griffipavixanthone showed high activity against P388, LL/2 and Wehl64 cell lines with very promising ED₅₀ between 3.40 and 6.80 μg mL^{-1} .¹⁰⁹ A Chinese patent reports promising anti-cancer properties for griffipavixanthone. They claim a strong inhibition effect of the bisxanthone or its salts on human lung, breast, prostatic and intestinal cancer cells, while showing no cytotoxicity to normal kidney epidermal cells. For lung cancer cells H520 in particular, they found that the cell cycle was blocked in the S stage, thereby preventing propagation of the cancers.¹¹²

3.10 Hyperidixanthone

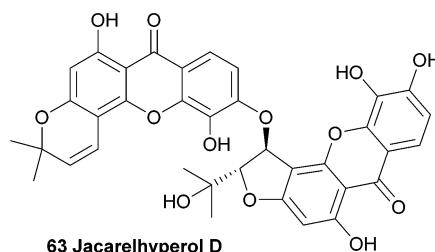
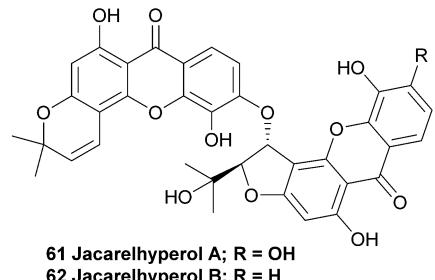
Hyperidixanthone (**60**) is a symmetrical dimer isolated from the plant *Hypericum chinense*.¹¹³ The compound was identified by a barrage of NMR techniques, including HMBC, and mass spectral data. Although hyperidixanthone is almost certainly axially chiral, the authors reported no details of optical activity.



3.11 Jacarelhypers A, B and D

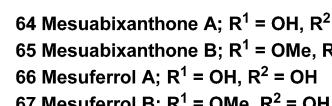
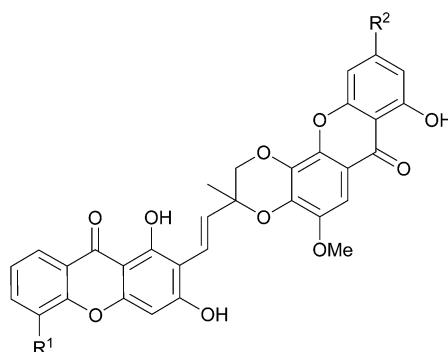
Jacarelhypers A and B (**61**, **62**) were isolated from *Hypericum japonicum* only a few years later than the bijaponicaxanthones, and the measured NMR spectra were found to be very similar to the spectra obtained from the bijaponicaxanthones. The relative structure of jacarelhypers A was elucidated to be an epimer of bijapoincaxanthone with a *trans*-configuration of the two protons at the chiral centres. Jacarelhypers B was subsequently identified as a 6'-dehydroxy variant of jacarelhypers A. Both jacarelhypers A and B were found to significantly inhibit platelet-activating factor (PAF)-induced hypertension in mice at 10 mg kg^{-1} without causing their blood pressure to rise. Therefore PAF inhibitors are considered to be potential drugs against allergic diseases.¹¹⁴ The isolation and structure elucidation of jacarelhypers D from the

same plant species was reported in 2007 in a short communication paper by Chen *et al.*¹¹⁵



3.12 Mesuabixanthones A and B

The bark of the south-east Asian tree *Mesua ferrea* has been widely used in traditional local medicine and is reported to exhibit antimicrobial and antiasthmatic activity. Therefore, the stem bark was investigated in an attempt to isolate the active compounds. Interestingly, the two novel bisxanthones mesuabixanthone A and B (**64**, **65**) were isolated from an extracted fraction that showed little biological activity in their initial bioassays. Using conventional mass spectrometric and NMR spectroscopic methods they identified the two novel bisxanthones. Mesuabixanthone B differed from mesuabixanthone A only by the methylation of the hydroxyl group at C-8, as determined by a NOESY experiment. No biological testing on the isolated compounds was reported.¹¹⁶

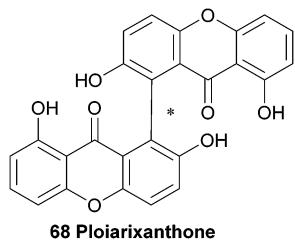


3.13 Mesuferrols A and B

The very similar mesuferrols A and B (**66**, **67**) have been isolated from an acetone extract of the bark of the same tree and were purified using chromatography.¹¹⁷ Mesuferrol A and B were identified by means of NMR spectroscopy and HR FAB-MS.

3.14 Ploiarixanthone

In 1990 a novel dimeric xanthone metabolite was isolated from branches of the shrub *Ploiarium alternifolium*. By synthesising a tetracetylated derivative and detailed comparison of the obtained NMR spectra with the data from the natural product, they identified ploiarixanthone (**68**) to be 8,8'-linked.⁸³ Ploiarixanthone is optically active ($[\alpha]_{D}^{25} = +23^{\circ}$), providing strong evidence for an enzymatic dimerisation process in its formation.



3.15 Puniceasides

In 1991 the first isolation and structure elucidation of a bis-xanthone C-glucoside named swertipunicoside (**77a**, Fig. 2) was reported. It was isolated from extracts from the whole plant *Swertia punicea* and the structure was confirmed by NMR experiments, particularly the selective INEPT technique (Insensitive Nuclei Enhanced by Polarisation Transfer).¹¹⁸ Only one year later the same group was also able to isolate and characterise 3-O-demethylswertipunicoside (**77b**) using several NMR techniques including APT, HETCOR and selective INEPT.¹¹⁹ Puniceasides B and C and isopuniceaside B have all been partially reduced to feature one tetrahydroxanthone core.

Recently, investigations showed that 3-O-demethylswertipunicoside has potent neuroprotective capabilities. Several oxidative toxicity tests were performed on xanthone treated and untreated PC12 cells. Increased cell viability was found in a 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) cell death assay where the cells were exposed to 1-methyl-4-phenylpyridinium ion (MPP⁺), rotenone or hydrogen peroxide. These neuroprotective effects were found to be caused by the elevation of TH and DJ-1 protein levels.¹²⁰

Hostettmann *et al.* isolated several xanthones from *Gentianella amarella* ssp. *acuta*, including one new dimeric xanthone. After structure elucidation with the use of 2D-NMR experiments and mass spectrometry they named the compound swertiabixanthone I 8'-O- β -D-glucopyranoside (**73b**).¹²¹

Remarkably, its aglycone, swertiabixanthone I (**73a**), was first isolated and identified more than twenty years earlier from *Swertia macrosperrm* plants.¹²² Recently both the glycone and the aglycone of swertiabixanthone I have also been isolated from

Swertia punicea.¹²³ In 2010, Guo *et al.* isolated two new dimeric O-glycoside xanthones, one new trimeric O-glycoside xanthone, two new trimeric C-glycoside xanthones and 12 known xanthones from the plant *Swertia punicea*.¹²³ The five new xanthones were identified using extensive HRESIMS and NMR spectroscopic experiments and were named puniceasides A–E (**69a**, **70a**, **72a**, **75a**, **76a**). For puniceaside B the *cis* configuration between H-5 and H-8 was confirmed using NOESY experiments. Also, it was found that the H-6' resonance pattern indicated that two different rotameric forms of puniceaside B exist. By heating to 120 °C in a ¹H NMR experiment, the two singlets of H-6' merged into one singlet. Puniceaside C was confirmed to be a trimeric xanthone glycoside by close examination of the results from mass spectrometric methods and HMBC NMR experiments. As far as is known, this was the first trimeric xanthone reported in the literature. By subjecting the compound to acid hydrolysis, the glycosidic moiety was found to be D-glucose. Two more trimeric xanthones were identified as puniceasides D and E. HMBC correlations showed the presence of the OMe group in puniceaside E. The absolute configurations of **75a** and **76a** could not be determined due to insufficient quantities of compounds. The presence of different rotameric configurations is likely, since heating the ¹H NMR samples to 120 °C resulted in the merging of the two singlets of H-6' into one singlet, similar to observations for puniceaside B. Puniceasides A–E, swertiabixanthone I 8'-O- β -D-glucopyranoside, 3-O-demethylswertipunicoside and swertipunicoside were tested for their neuroprotective activity against hydrogen peroxide-induced PC12 cell damage. In particular puniceaside B was found to be very potent with a cell viability of 98.1 ± 6.8% at a concentration of 25 μ g mL⁻¹ compared to hydrogen peroxide treated cells. Interestingly, swertiabixanthone I 8'-O- β -D-glucopyranoside and 3-O-demethylswertipunicoside were found to potently stimulate the damaged PC12 cells to grow, resulting in cell viability scores of 123% and 158%. In 2012, Guo *et al.* reported the characterisation of 16 new xanthone compounds from extracts of *Swertia punicea*. After studying the ESI-MS fragmentation behaviours of 17 known xanthones very closely, they could apply this knowledge to high-performance liquid chromatography diode-array detection/tandem mass spectrometric results and thus identify 11 new dimeric and 4 new trimeric xanthones (**69b–d**, **70b** and **c**, **71a** and **b**, **72b**, **75b**, **76b**, **74**, **73c–e**, **77c**).¹²⁴

4 Xanthone dimers from fungi

Fungi have yielded some beautiful xanthone dimers, the majority of which possess a common biaryl bond as the method of their linkage. The biaryl bond is most often arranged in a 2,2'-manner with respect the common xanthone hydroxyl moiety; in a couple of cases, natural products with 2,4'-biaryl bonds have been described, as has the somewhat rarer 4,4'-biaryl linkage. Additionally, the position of methylation of the xanthone core is found at the C3, C6 or sometimes C10a positions, which can provide insight to the biosynthetic origin of the compounds (see Table S-1† for structural details).



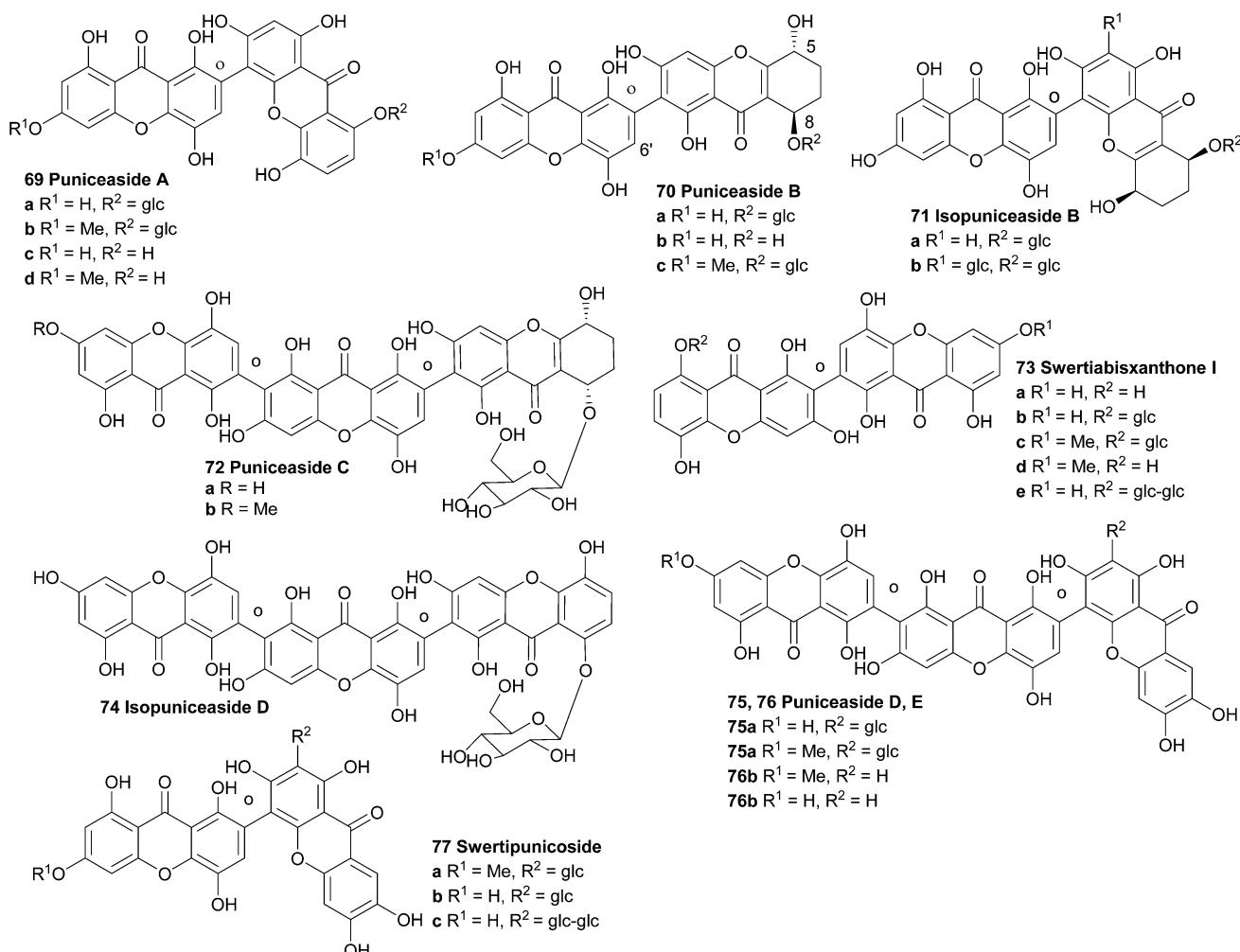
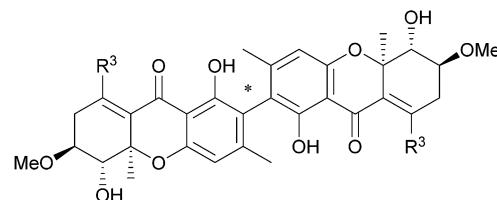


Fig. 2 Puniceaside and swertia bisxanthone derivatives (absolute stereochemistries shown).

4.1 Ascherxanthones

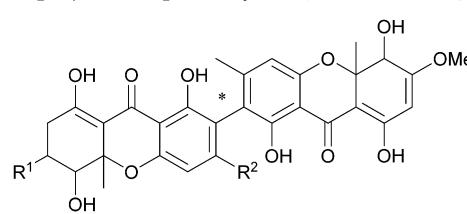
Ascherxanthone A (78) was isolated from the methanol extract of the mycelia of *Aschersonia* sp. BCC 8401, after significant antimalarial activity in *Aschersonia* cultures was observed.¹²⁵ Evidence for the symmetric homodimeric structure of ascherxanthone A was provided by mass spectrometry and the presence of only 16 carbon signals in the ^{13}C NMR spectrum. The tetrahydroxanthone structure was established on the basis of the HMBC correlations, and using COSY and HMQC NMR spectroscopic methods the entire structure was elucidated. Intense correlation from the phenolic proton to the quaternary carbon placed this carbon at C2, confirming a 2,2'-biaryl-linked dimer. Interestingly, the structure of ascherxanthone A is very close to TMC 315A₁, TMC 315A₂, TMC 315B₁ and TMC 315B₂ (80–83); four dimeric xanthones that were isolated from *Ceu-thospora* sp. TMC1678 and reported in a Japanese patent to be RANKL (receptor activator of NF- κ B ligand) antagonists that can prevent or control osteoporosis.¹²⁶ It should be noted the Japanese patent does not provide any stereochemical information for TMC 315A₂ (81), which may therefore be in fact identical to ascherxanthone B (79). Note that ascherxanthone A bears one less hydroxyl than the other compounds in this family, and one

component of each TMC 315B₁ and B₂ has been reduced to the relatively rare dihydroxanthone tricycle.



80 TMC 315A; $R_1 = \text{Me}, R_2 = \text{OMe}, R_3 = \text{OH}$ (no stereochemistry known)

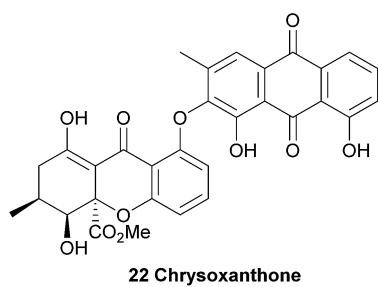
81 TMC 315A₂; $R_1 = \text{OMe}, R_2 = \text{Me}, R_3 = \text{OH}$ (no stereochemistry known)



In a search for new agents against the rice blast fungus *Magnaporthe grisea* that has developed a fungicide resistance, culture broth extracts from approximately 800 fungal strains were investigated in an *in vitro* screening assay. A culture of *Aschersonia luteola* BCC 8774 was found to produce ascherxanthone B, a compound active against *M. grisea* with an IC_{90} value of $0.58 \mu\text{g mL}^{-1}$. The structure of ascherxanthone B (79) was elucidated using COSY, HMBC and HMQC NMR experiments and found to be almost identical to ascherxanthone A, with the only difference being the replacement of the olefinic proton at $\delta_{\text{H}} 6.87$ (H-8) in ascherxanthone A with a chelated hydroxy resonating at $\delta_{\text{H}} 13.43$ (C-8 hydroxyl, broad singlet). Interestingly, it was found that ascherxanthone A was nearly inactive in the same biological assay against *M. grisea* with an IC_{90} value of over $50 \mu\text{g mL}^{-1}$.¹²⁷ However, ascherxanthone A did show significant activity against *Plasmodium falciparum* K1 ($IC_{50} = 0.20 \mu\text{g mL}^{-1}$). Furthermore, it also showed cytotoxicity to Vero cells ($IC_{50} 0.80 \mu\text{g mL}^{-1}$) and three cancer cell lines (IC_{50} values between 1.7 and $0.16 \mu\text{g mL}^{-1}$).¹²⁵

4.2 Chrysoxanthone

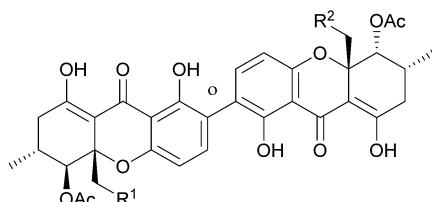
Chrysoxanthone, although not a xanthone dimer, is included here as an unusual example of the mixed-biological derivative 'xanthraquinones', which also include the beticolins^{128–131} and the xanthoquinodins.^{132–135} Chrysoxanthone, as reported by Anke and co-workers,⁵⁶ features a form in which 2-hydroxy-chrysophanol is coupled to blennolide A (hemisecalonic acid B) through a biaryl ether linkage. The authors determined the novel structure with 2D NMR and mass spectral techniques, and it was found to possess antimicrobial and antifungal properties. These most likely result directly from the discrete bioactivities of each the xanthone and anthraquinone components. This compound, alongside the xanthoquinodins and beticolins, represents a biosynthetic junction of a xanthone and one of its likely biosynthetic precursors (anthraquinone). The synthesis of an ether linkage between two xanthenes has been recently reported by Sahin, Nieger and Bräse.¹³⁶



4.3 Dicerandrols A–C

Dicerandrols A–C (84–86) were isolated from the culture broth of the *Phomopsis longicolla*, an endogenous fungi from the mint species *Dicerandra frutescens*. Extensive NMR experiments were complemented by a positive FeCl_3 experiment, indicating the 2,2'-linkage mode of the dimer (the test is positive for a *para*-unsubstituted phenol). The ^1H NMR spectrum of dicerandrol C

was found to be simple, suggesting a plane of symmetry. 2D NMR experiments were used to determine the relative stereochemistry for this compound, and by inference, those of dicerandrol B and C.¹³⁷ In 2013 the absolute configurations of dicerandrols B and C were determined by TDDFT electronic circular dichroism (ECD) calculations.¹³⁸ It is likely that dicerandrol A has the same absolute stereochemical configuration, but to date only the relative configuration has been reported.



84 Dicerandrol A (relative configuration); $R^1 = \text{OH}$, $R^2 = \text{OH}$

85 Dicerandrol B (absolute configuration); $R^1 = \text{OAc}$, $R^2 = \text{OH}$

86 Dicerandrol C (absolute configuration); $R^1 = \text{OAc}$, $R^2 = \text{OAc}$

The dicerandrols have antimicrobial activities (*Bacillus subtilis* and *Staphylococcus aureus*) with increased activities related to the decreased degree of acetylation of the molecules (*i.e.*, A > B > C). The dicerandrols were also active against two cancer cell lines, HCT-116 and A549 (colon and lung tumour, respectively).¹³⁷ In 2008, dicerandrol A was also isolated from *Phomopsis* sp. PSU-D15 and used in comparison studies to help identify a new member of the phomoxanthones (deacetylphomoxanthone B 101).⁹⁰ Dicerandrol C was isolated from a *P. longicolla* obtained from red seaweed *Botrychia radicans* and identified with the use of 1D and 2D NMR spectroscopic methods as well as by mass spectrometry.⁸⁹ Another study performed several biological tests on dicerandrols A, B, and C that were also isolated from *P. longicolla*. Using disk diffusion assays they found that dicerandrols A and B possess antibacterial activity against *Xanthomonas oryzae* KACC 10331, a cause for bacterial blight in rice. Subsequently, dicerandrol A was also tested for antimicrobial activity against seven other *X. oryzae* strains, several Gram-positive and Gram-negative bacteria, a fungus and a yeast. It was found that dicerandrol A showed a relatively high activity in this broad spectrum of species, although it was often bested by commercial antibiotics. By performing growth and time-dependent production of secondary metabolites studies, it was found that the dicerandrols were produced mainly after 4 days of fermentation and that the maximal production was at day 10. This led to a maximal antibacterial activity after day 14.¹³⁹ These results are useful in the production of a natural preventive medicine against bacterial blight of rice, which is very relevant since it was found that *X. oryzae* has developed a resistance to the available antibiotics. A more detailed continuation study on the time-dependent production of dicerandrols and other antibacterial compounds in *P. longicolla* confirmed that the optimum fermentation time for maximal antibacterial activity of the fungus is indeed 14 days.¹⁴⁰ Dicerandrols A–C were also isolated in 2013 from the mycelium of a culture broth of *Phomopsis longicolla* S1B4 and were tested against a strain of *X. oryzae*. In this antimicrobial



activity test the dicerandrols A–C showed a MIC of 8, 16 and 16 $\mu\text{g mL}^{-1}$, respectively.¹⁴⁰

The dicerandrols and the structurally similar penexanthone A were submitted to several biological activity tests *versus* a broad range of tumour cell lines.¹⁴¹ The majority of the *in vitro* screening tests were performed in the presence of non-malignant accessory cells, such as bone marrow stromal cells, since it was found that the activities of potential drugs can be affected by microenvironment-dependent drug resistance or sensitisation of the tumour cells.¹⁴² Among the tested compounds, dicerandrol B was found to be the most promising candidate for further investigations. It showed moderate activity against Dox40, Farage, H929, HT, OPM2 and RPMI8226 cell lines in the presence of stromal cells with IC₅₀ values of 2.3, 1.3, 3.4, 1.3, 1.5, and 1.2 μM , respectively. More importantly, it was found that dicerandrol B showed a relatively low toxicity against human immortalised non-malignant cells, such as HS-5 bone marrow stromal cells, HOBIT osteoblast-like cells, THLE-3 hepatocytes, and SVGp12 astrocytes, compared to the values found for cancer cell lines RPMI8226 and H929. This relative selectivity is cause for further investigations into the biological activity of this compound.¹⁴¹ Dicerandrols A–C were also isolated from *Phomopsis* sp. HNY29-2B and tested for their cytotoxicity against human breast cancer, colon cancer, lung cancer, liver cancer and breast epithelial cell lines. Dicerandrol A showed a broad anti-tumour activity, but was also cytotoxic to the breast epithelial cells. Dicerandrol B and penexanthone A (101) showed also some cytotoxicity to the cancer cells, but were found to be more selective and less strongly damage the breast epithelial cells. In this assay dicerandrol C was found to be not very cytotoxic, suggesting that the free hydroxy groups are important as a key pharmacophore.¹⁴³ In a study that mainly focusses on the biological activity of phomoxanthone A (106), dicerandrols B and C are also investigated. It was found that the dicerandrols were both cytotoxic to murine lymphoma cancer cell lines (IC₅₀ values of 10 and 1.1 μM , respectively) and that dicerandrol C is slightly pro-apoptotic.¹³⁸

4.4 Ergochromes

Ergochrome dimers comprise a colourful group of xanthones first reported by Kraft in 1906 as a single compound of the formula C₁₄H₁₄O₆ after separation from *Claviceps purpurea*.¹⁴⁴ The contamination of rye with ergochrome-containing *C. purpurea* led to epidemics known as ergotism or 'St. Anthony's Fire' in middle-ages Europe. The mixture of toxins produced by this fungus was responsible for the toxicity and debilitating effects on the affected population. The ergochrome dimers (21, 23, 87 to 99, Fig. 3) include the wide-spread secalonic acids, ergochrysin, ergoflavin and chrysergotic acid, many of which possess a huge variety of potent bioactivities. The ergochrome dimers are composed of several monomeric units (hemi-secalonic acids A–F, aka 'blennolides'),⁴⁷ which are arranged in a variety of dimers and heterodimers. In all known secalonic acids, the methyl and methoxycarbonyl substituents are found to be *trans*-configured. Whilst secalonic acids are ester homodimers, other heterodimer ergochromes are known. For

example, ergochrysin are lactone/ester heterodimers, ergoflavin is a homodimer,¹⁴⁵ whilst ergoxanthin has undergone structural rearrangement to form a pendant butyrolactone ring on one half of the molecule.¹⁴⁶

Due to the early time at which they were first studied, modern spectral methods were not applicable to facilitate the structural elucidation of the ergochromes, and early structural assignations on these compounds were frequently revised in later years. For example, the initial misidentification of ergochrysin and ergoflavin as secalonic acids was later revised by Bergmann in 1932 as dimers with the formula C₂₈H₂₈O₁₂.¹⁴⁷ A number of papers followed in the 1950s to 70s, over which time more secalonic acids were isolated and identified. One major point of contention in the early days of exploration of this compound class was as to whether the natural products were 2,2'-, 4,4'- or even 2,4'-linked.^{148,149} The biaryl linkage was finally definitively assigned as 2,2'-linked when Mayo and Aberhart reported on the isolation, in crystalline form, of six compounds from a Portuguese ergot drug in 1965.¹⁵⁰ In contrast to earlier reports,¹⁵¹ these authors showed that ergochrome dimers ubiquitously gave positive results in the Gibbs test, indicating the presence of an unsubstituted aryl methine in a position *para*- to the phenolic moiety, and thus a 2,2'-linkage.

A comprehensive analysis of the ergochrome constituents of *C. purpurea* by Franck and co-workers led to the reporting of a total of ten component compounds, including a new homodimer named ergochrome DD (94, Fig. 3), as well as the remaining six heterodimers possible with a combination of one subunit of each of those four species A, B, C and D. The majority of which they identified as the ergochromes are diastereomeric at positions 6,6', 5,5', 10 and 10'. The authors also identified the absolute stereochemistry at these centres.¹⁵²

Büchi and co-workers reported novel secalonic acid F (88, ergochrome BE) and secalonic acid D (91, ergochrome EE) in 1977, following isolation from the fungus *Aspergillus aculeatus*.¹⁵³ The following year, then-novel secalonic acid G (90) was isolated alongside secalonic acids A and E from *Pyrenopeziza terrestris* and the structure elucidated by CD and NMR spectroscopy.²³

Secalonic acid D (SAD) is a major environmental toxin, being isolated from *Penicillium oxalicum*, a major microbial contaminant of freshly-harvested corn (in one study present on 44% of pre-harvest corn crops).¹⁵⁴ Teratogenic effects were observed in the development of rats that were exposed to SAD injected during fetal development. SAD was lethal to mice when injected intraperitoneally in the 25–50 mg kg^{-1} range.¹⁵⁵ The teratogenic and toxic nature of these compounds is alarming in light of their propensity to contaminate foodstuffs.¹⁵⁶ The rate of birth-defects and teratogenicity-initiated spontaneous abortions in humans is also alarmingly frequent.

Despite the widespread occurrence of the ergochromes, none had been synthesised until the end of 2013, and neither had any other xanthone dimers or trimers. In early 2014, Porco and co-workers at long last described the synthesis of secalonic acids A and D, utilising a Cu(I) catalysed di-destannylative coupling.^{157,158} This reaction represents a significant breakthrough for the study of the xanthone



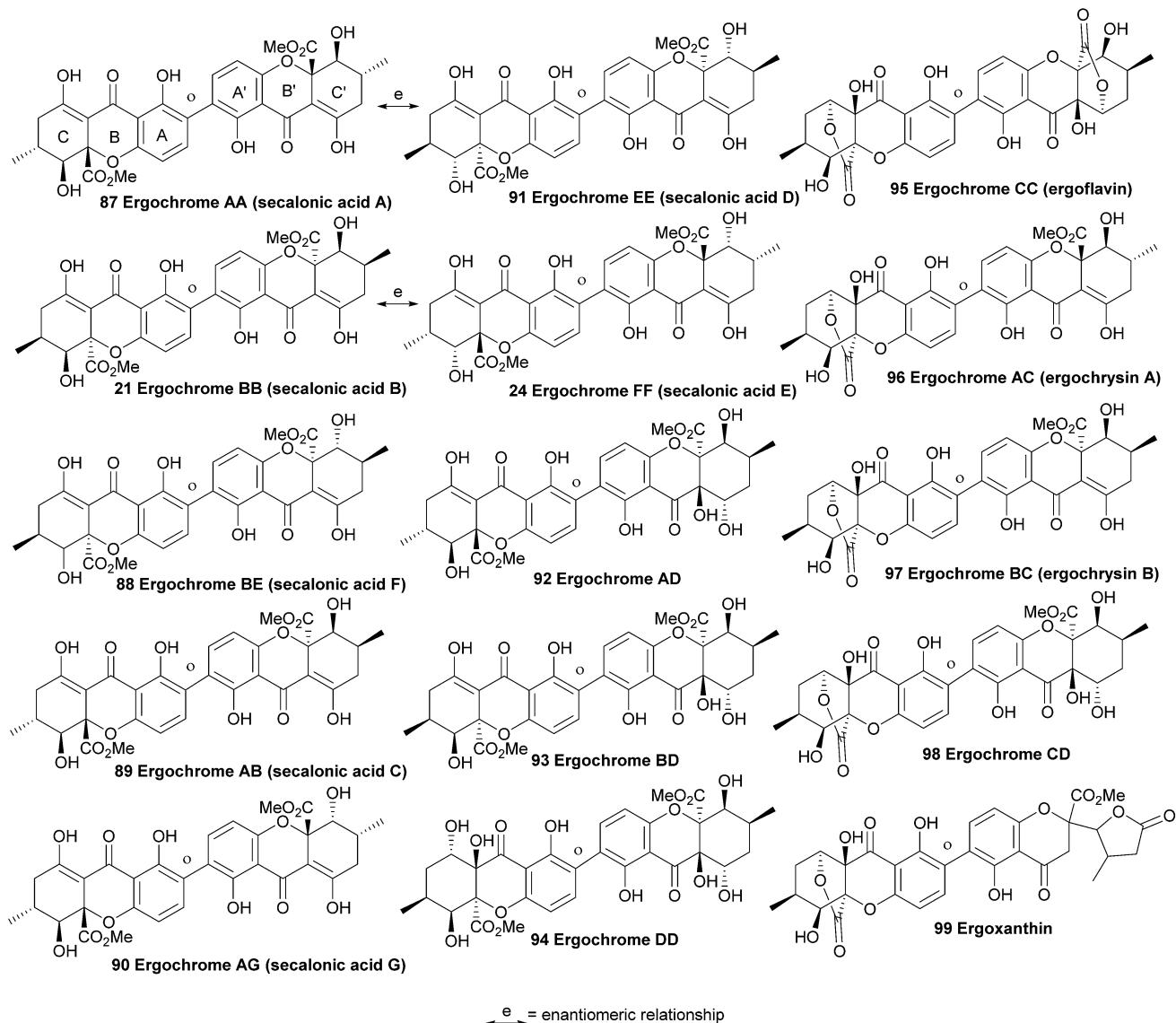


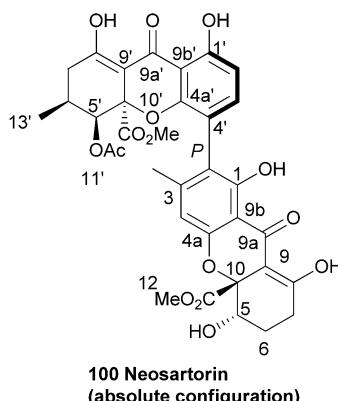
Fig. 3 Ergochromes showing heterodimeric structures, stereochemistry and enantiomeric relationships.

dimers,⁶² and may open up the field to the synthesis of interesting analogues for medicinal chemistry studies. Recent advances in Ullmann coupling may make the strategy of Whalley and co-workers towards ergoflavin a more feasible route to the natural products than was originally the case in 1971.¹⁵⁹ Bräse and co-workers have described a novel one-pot methodology for the synthesis of symmetrical biaryls¹⁶⁰ as a part of their ongoing investigations in the synthesis of the secalonic acids. Sahin, Nieger and Bräse have also published on the oxidative coupling of various hexahydroxanthenols. The application of an iron complex as an oxidant converted 2-hydroxy substituted xanthenes to the 3,3'-bis-coupled bixanthenes.¹³⁶ It appears that these two papers represent, until the end of 2013, the only successful chemistry published on the topic of xanthone or xanthene biaryl-coupling.

4.5 Neosartorin

Neosartorin (**100**) was isolated in 1998 from the cultured mycelium of the soil mould *Neosartorya fischeri*. A combination of 2D NMR experiments determined that neosartorin was an isomer of eumitrin A₁. However, close examination of the long range coupling constants of nuclei around the two arene units revealed a 2,4'-biaryl linkage instead of the 4,2-linkage found in eumitrin A₁.¹⁶¹ A closer investigation of the low-intensity long-range NOEs by a 1D double-pulsed field gradient spin-echo NOESY NMR experiment resulted in confirmation of the relative configuration of the two xanthone subunits. The absolute stereochemistry of neosartorin was axially *P* and found to be (a*R*,5*S*,10*R*,5'*S*,6'*S*,10'*R*), this was determined on the basis of its electronic circular dichroism (ECD) spectra in conjunction¹⁶² with TDDFT-ECD calculations. They found that the nonplanar configuration of the aromatic rings connecting

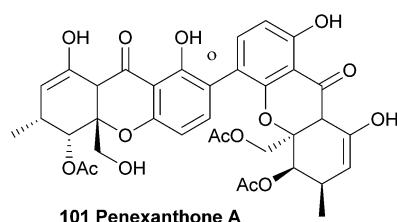
the two subunits was stabilised by long-range nonbonding interactions between the substituents of different xanthone units.¹⁶³



A taxonomical study of *Aspergillus* species conducted in 2005 reports the production of neosartorin in the new species *A. lentinus*, *A. fumigatiaffinis* and *A. novofumigatus*, but not in *A. fumigatus*. The presence of neosartorin was based on evaluation of UV spectral evidence. Based on these findings and other similarities the authors conclude that the taxa of *Neosartorya fischeri* and the *Aspergillus* species are chemically related.¹⁶⁴ Neosartorin was found to show a strong activity against Gram-positive bacteria such as *Staphylococcus aureus* (MIC: 8 µg mL⁻¹) and *Bacillus subtilis* (MIC: 4 µg mL⁻¹), while it did not affect the tested Gram-negative bacteria. In cytotoxicity tests against several cancer cell lines it scored very badly, showing almost no notable cytotoxicity.¹⁶²

4.6 Penexanthone A

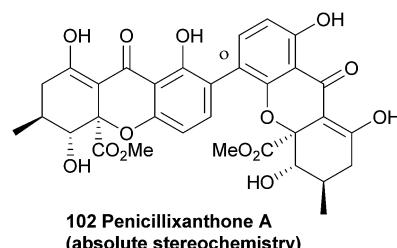
Recently, Clardy *et al.*¹⁴¹ isolated dicerandrols A–C and a novel dimeric xanthone named penexanthone A (101) from CR1642D, a *Penicillium* sp. obtained from the Costa Rican rainforest. The structure, including relative configuration, of penexanthone A was elucidated using mainly 2D NMR techniques (HSQC, HMBC, NOESY and ROESY) and molecular modelling with Chem3D Ultra (9.0). The two monomers were found to be connected through a 2,4'-linkage. The dicerandrols were identified by comparison of several physical and spectroscopic data sets (UV, IR, ¹H NMR, [α]_D and MS) with the literature values. The authors note that the structure of penexanthone A was already reported in a Korean patent, although the ¹H NMR spectra was unclean, suggesting an impure sample.¹⁶⁵ Penexanthone A is identical to monodeacetylated phomoxanthone B (*vide infra*). Penexanthone A was submitted to several biological activity tests *versus* a broad



range of tumour cell lines, however the observed IC₅₀ values indicate that penexanthone A is only weakly active in the presence of stromal cells.¹⁴¹ Penexanthone A was also isolated from *Phomopsis* sp. HNY29-2B and tested for its cytotoxicity against human breast cancer, colon cancer, lung cancer, liver cancer and breast epithelial cell lines. The dimeric xanthone showed relatively high cytotoxicity to the cancer cells and a reduced cytotoxicity towards the immortalised breast epithelial cells, which is a very useful feature when developing new anti-cancer drugs.¹⁴³

4.7 Penicillixanthone

The 2,4'-biaryl-linked dimer penicillixanthone was reported around ten years ago^{88,166} by Chinese groups, who described the absolute stereochemistry. This compound has been re-isolated in 2013.¹⁶⁷

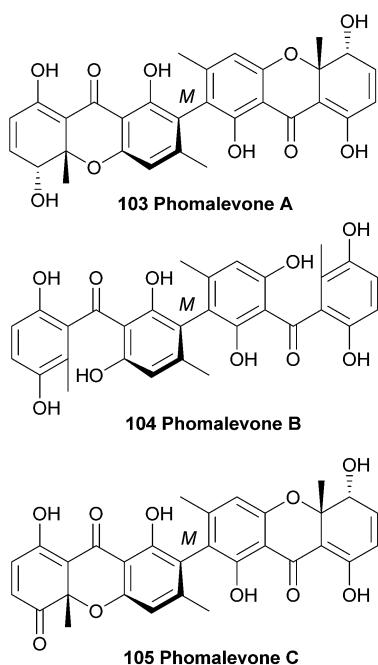


4.8 Phomalevones

In 2011, phomalevones A–C (103–105) were isolated from a Hawaiian isolate of *Phoma* sp. (MYC-1734 = NRRL 39060; Cucurbitaceae). The structures of phomalevones A–C were determined by detailed analysis of NMR and MS data.¹⁶⁸ The absolute configuration of the sp³ stereogenic centres of phomalevone A were revealed by analysis of NMR spectra of Mosher type derivatives.

Phomalevone A was found to resemble ascherxanthone B. The ¹³C NMR spectrum of phomalevone B showed only 15 carbon signals, thus indicating a symmetrical, homodimeric structure. Via HR-ESI-MS data it was determined that phomalevone B is an isomer of phomalevone A. However, the NMR spectra showed a significant downfield shift of the only carbonyl signal relative to its location in the spectrum of phomalevone A and the presence of two sp² carbon signals instead of sp³ signals as in the spectrum of phomalevone A. By comparing the NMR spectroscopic data with that of acremonidin D,¹⁶⁹ a symmetrical dimer with two benzophenone monomer units connected *via* a methylene linkage, a structural assignment could be made.¹⁶⁸ The occurrence of phomalevone A together with the benzophenone analogue, phomalevone B, is consistent with several reports that hydroxylated benzophenones are very likely to be key intermediates in xanthone biosynthesis.^{25,170,171} HMBC spectra of phomalevone C revealed its heterodimeric nature, with one subunit the same as the monomer unit in phomalevone A, while the other has one secondary alcohol group oxidised to a carbonyl group. Phomalevones A–C showed antibacterial activity in agar disk diffusion assays at 100 µg per disk against *Bacillus subtilis* and *Staphylococcus aureus*. Additionally, phomalevone B was also active against *Candida*

albicans and *Escherichia coli* and phomalevone C showed activity against *Fusarium verticillioides* and *Aspergillus flavus* (IC_{50} value of $4 \mu\text{g mL}^{-1}$, compared to an IC_{50} value of $5 \mu\text{g mL}^{-1}$ for the antifungal standard Nystatin).¹⁶⁸

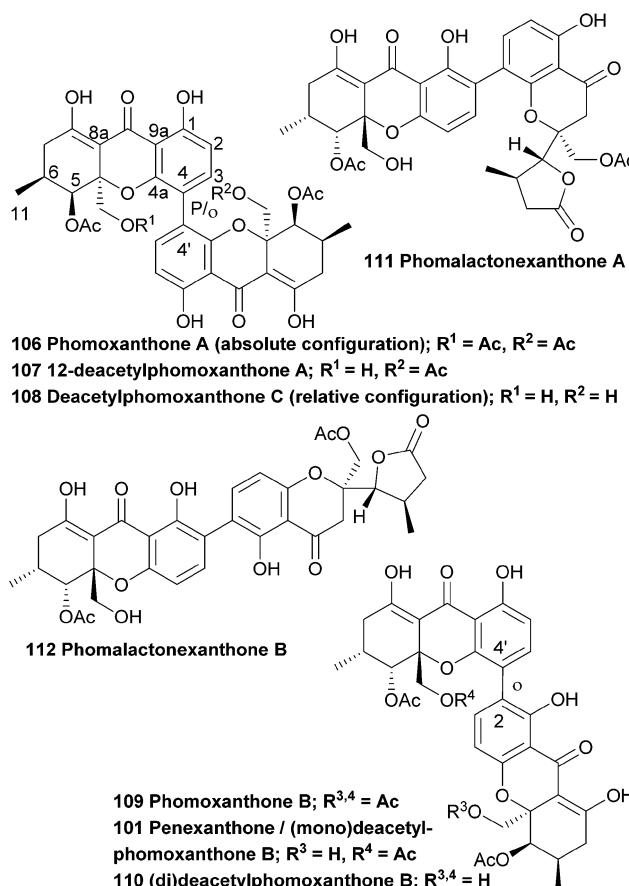


4.9 Phomoxanthones

The antimalarial activity of methanolic extracts of mycelia of the endophytic fungus *Phomopsis* sp. BCC 1323 led to researchers isolating and identifying phomoxanthones A and B (106, 109). The relative stereochemistry could be determined by synthesising a deacetyl derivative of phomoxanthone A and comparing the change in coupling signals. Phomoxanthone A and B were the first reported examples of naturally occurring xanthone dimers with 4,4' or 2,4' linkages.¹⁷² In 2008, the deacetyl derivative of phomoxanthone B (110) was isolated from the same fungus species by Kirtikara *et al.*⁹⁰ Phomolactonexanthone A and B are both derived either from coupling of a lactone-analogue of a xanthone (as the blennolides)⁴⁷ and a xanthone itself, or from further derivatisation from a dimeric xanthone.

Biological tests with deacetylphomoxanthone B showed a higher antibacterial effect against the pathogen *Xanthomonas oryzae* KACC 10331, a cause for bacterial blight in rice, than a positive control with 2,4-diacylphloroglucinol.¹³⁹ Deacetylphomoxanthone B and C (108) and phomolactonexanthone A and B (111, 112), that were isolated from *Phomopsis* sp. HNY29-2B, were tested for their cytotoxicity against a series of human cancer lines and it was found that while deacetylphomoxanthone B has a reasonable cytotoxicity against all tested cancer cell lines, deacetylphomoxanthone C and the phomolactonexanthones were virtually inactive against all cell lines.¹⁴³ Phomoxanthone A and B showed antimalarial (*Plasmodium falciparum* K1, multi drug resistant strain, IC_{50} of 0.11 and 0.33 $\mu\text{g mL}^{-1}$, compared to IC_{50} of 0.16 and 0.0011 $\mu\text{g mL}^{-1}$ for chloroquine diphosphate and artemisinin, respectively) and

antitubercular activity (*Mycobacterium tuberculosis*, H37Ra strain, IC_{50} of 0.5 and 6.25 $\mu\text{g mL}^{-1}$, compared to IC_{50} of 0.05 and 2.5 $\mu\text{g mL}^{-1}$ for isoniazid and kanamycin sulfate, respectively). Biological tests on cancer cell lines (KB cells, BC-1 cells and Vero cells) showed that the natural products are cytotoxic (IC_{50} $\mu\text{g mL}^{-1}$; 0.51–1.4 for phomoxanthone A and 0.7–4.1 for phomoxanthone B), although it was noted that standard drugs are more potent in both the bacterial and cancer cells tests. The deacetyl derivative was also investigated in these assays and was found to be mostly inactive in all the tests – it is speculated that this might be due to its lower lipophilicity.¹⁷²



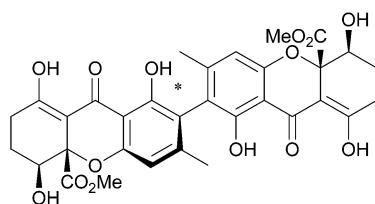
The absolute configuration (aS,5R,6R,10 aR,5'R,6'R,10a'R) of phomoxanthone A was determined by a combination of single crystal X-ray analysis with measured and calculated CD spectra, in 2005.⁸⁶ However, in 2013 the absolute configuration was revised to be (aR,5S,6-S,10 aS,5'S,6'S,10a'S), when a team of researchers performed a detailed X-ray analysis study and TDDFT-ECD calculations after they suspected that the configuration might have been wrongfully assigned.¹³⁸ The suspicion arose when they isolated phomoxanthone A together with dicerandrols B and C from *Phomopsis longicolla* and found that all their stereocenters had the S absolute configuration. In this same study they also found a new natural product from the same endophyte, which was identified by NMR studies as 12-deacetylphomoxanthone A (107).

Biological tests showed that phomoxanthone A has promising activity against Gram-positive bacteria and fungi.⁸⁶ In 2013

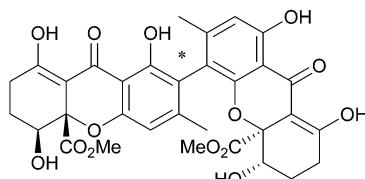


Lee *et al.* isolated several secondary metabolites from the mycelium of a culture broth of *Phomopsis longicolla* S1B4, including dicerandrols A–C (84–86) and a compound they named deacetylphomoxanthone B, 101 (which is in fact a monodeacetylphomoxanthone and is identical to penexanthone A). This compound exhibits a strong antimicrobial activity against gammaproteobacteria *Xanthomonas oryzae* with a MIC of 32 µg mL⁻¹, compared to dicerandrols A–C which showed 8, 16 and 16 µg mL⁻¹ respectively.¹⁴⁰ Phomoxanthone A has also been found to strongly inhibit the proliferation of the murine lymphoma cell line L5178Y, as well as other cell lines, including some cisplatin resistant cancer cell lines.¹³⁸ Remarkably, the cytotoxicity of phomoxanthone A towards the lymphoma cells was found to be increased over 100-fold with respect to healthy blood cells. A semisynthetic fully deacetylated phomoxanthone A showed no activity in this assay. Proksch *et al.* also analysed the apoptosis inducing potential of phomoxanthone A by flow cytometric analyses of hypodiploid nuclei in Jurkat T cells.¹³⁸ The presence of these hypodiploid nuclei is indicative of aspartate-directed cysteine protease (caspase) induced DNA fragmentation, which is an indicator for apoptosis. They found that among all tested substances the 4,4'-linked phomoxanthone A and deacetylphomoxanthone A scored highest on these tests and actively induce apoptosis. On top of that, they also found that they activate immune effector functions in murine immune cells, such as primary T lymphocytes, NK cells and macrophages, as was observed by the upregulation of cell-type specific activation markers. During a preliminary structure–activity assessment the authors conclude that the presence of the acetyl groups and the 4,4'-linked system in xanthone dimers 106 and 107 are important to maintain the interesting biological properties, possibly because of an underlying mechanism where there is a necessity to pass the cellular membrane. The deacetylated xanthones are less lipophilic and therefore often score very low on biological tests.

When comparing the overall activity of the phomoxanthones with other similar xanthones, it must indeed be noted that the acetylated species, such as the dicerandrols and neosartorin,



113 Rugulotrosin A (relative configuration)



114 Rugulotrosin B (relative configuration)

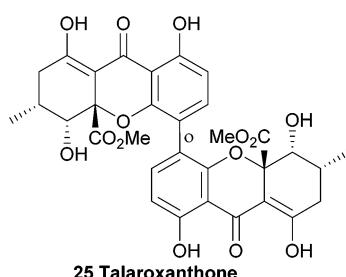
often show quite interesting biological properties, whereas more often the non-acetylated compounds either were not tested, or did not show promising activities. One could speculate that this is possibly due to enhanced cell penetration, however, in the case of the dicerandrols, the more acetylated dicerandrol C was found to be much less cytotoxic to cancer cells than the dicerandrols A and B that have free hydroxyl groups.

4.10 Rugulotrosins

Rugulotrosins A and B (113, 114) were isolated from cultures of *Penicillium* sp. derived from Australian soil samples. Rugulotrosin A is symmetrically 2,2'-linked while rugulotrosin B is a 2,4' linked dimer. The structures of these compounds were determined by NMR and MS techniques, and supported by single crystal X-ray analysis of rugulotrosin A. The compounds were of considerable activity against several microbial species (*Bacillus subtilis*, *Enterococcus faecalis* and *Bacillus cereus*).¹⁷³

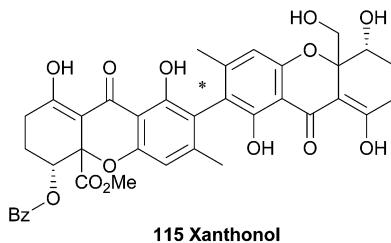
4.11 Talaroxanthone

The identity of talaroxanthone (25) was reported by Koolen and co-workers in 2013. This compound, featuring a 4,4'-linked diaryl motif, was isolated from the culture broth of DgCr22.1b, an endophytic *Talaromyces* sp. fungi isolated from the healthy tissue of the medicinal plant *Duguetia stelechantha* in the Amazonian rainforest. The yellow powder that was subsequently isolated by chromatographic techniques was determined to be the structure shown by HRMS and several 2D NMR techniques. Whether the medicinal qualities of the plant material are due to the presence of this fungal-derived compound is as yet unclear.⁵⁷



4.12 Xanthonol

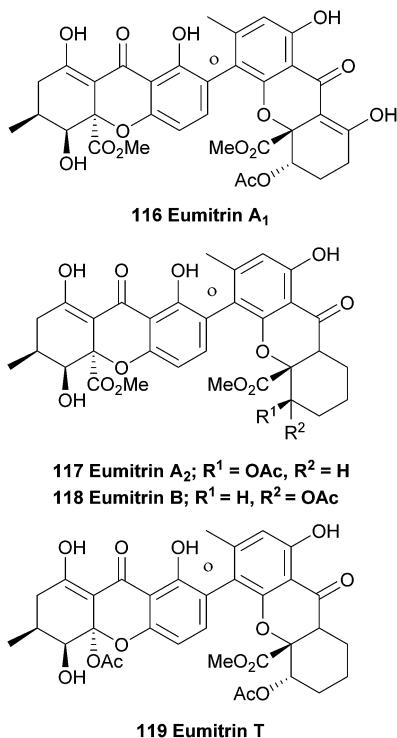
The unsymmetrical 2,2'-biaryl-linked dimeric xanthone, xanthonol (115), was isolated from the fermentation broth of a non-sporulating fungi found in the leaf litter of the plant *Manikara bidentata*. Xanthonol bears the methyl-substituents in the aryl positions, rather than the 6,6'-methylation of the ergochromes etc. Also interesting is the benzoylated alcohol at the C5' position. Xanthonol is an anthelmintic, as revealed by tests upon the larvae of *Lucilia sericata*, *Aedes aegypti*, and *Haemonchus contortus*.¹⁷⁴



5 Xanthone dimers from lichens

5.1 Eumitrins

In 1967 Asahina first reported the occurrence of yellow pigments named eumitrins A₁, A₂, B and T (116–119) isolated from the lichen *Usnea baileyi*.¹⁷⁵ Eumitrins A and B were later isolated from benzene extracts of the dried lichen thalli.¹⁷⁶ Shibata *et al.* found that the fraction of eumitrin A could be separated into A₁ and A₂ by recrystallisation and elucidated the chemical structures of eumitrins A₁, A₂ and B by spectroscopic methods.¹⁷⁷ By measuring differences in chemical shifts in NMR experiments after acetylation of the compounds, they were able to assign the biphenyl junction in these dimers as a 4,2'-linkage. Using single-crystal X-ray analysis of a tribromo derivative of eumitrin B they confirmed the structures of eumitrins A₁, A₂ and B. Interestingly, these xanthones have been reduced down from their normal oxidation states to bear a non-oxygenated/non-aromatic cyclohexenyl C-ring.



In 2002 a Japanese patent was filed, claiming that eumitrins A₁ and A₂ could be used as inhibitors for nitric oxide formation by macrophages and thus be useful for a wide range of illnesses.¹⁷⁸ Elix *et al.* report the extraction of secalonic acid A and eumitrin T

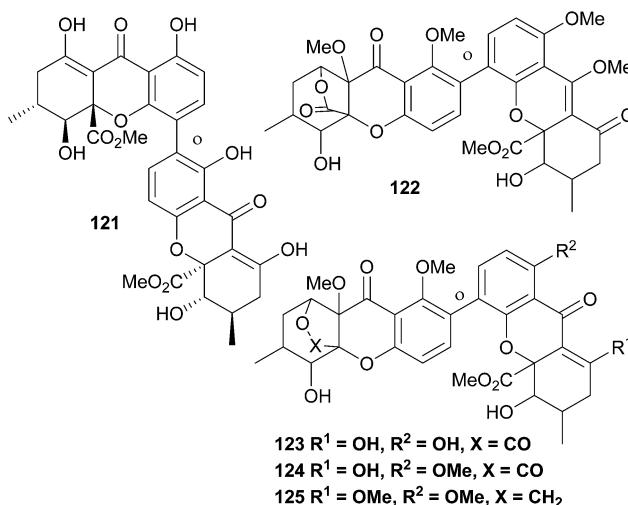
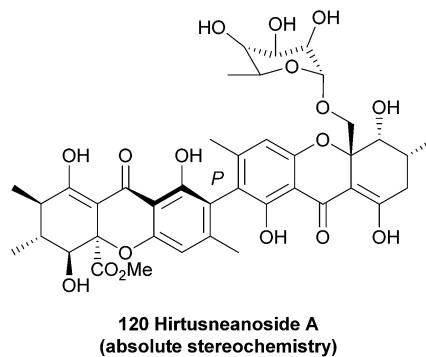


Fig. 4 Miscellaneous unnamed dimeric xanthones.

from the lichen-forming fungus *Physconia distorta*. Interestingly, they found that these dimeric xanthones were only found when intact lichen thalli were investigated and not in axenic cultures of the fungus, indicating a clear difference in metabolic pathways between the lichen and the axenic fungus.⁹² Recently, the same group investigated the contents of 18 lichen obtained from the mountainous areas of Malaysia using TLC and HPLC analyses. Amongst the secondary metabolites found in the lichen *Usnea baileyi* they found eumitrins A₁, A₃, B₁ and B₂. However, the structures of eumitrins A₃ and B₂ are not yet elucidated.¹⁷⁹

5.2 Hirtusneanoside

Hirtusneanoside A (120) was first isolated from the lichen *Usnea hirta* in 2007 by Řezanaka and Sigler, and found to be a L-rhamnose-O-deoxyglycoside of an unsymmetrical dimeric tetrahydroxanthone.¹⁸⁰ Fascinatingly, hirtusneanoside features an additional methyl in relation to the secalonic acids; the biosynthetic origin of this methyl group is as yet unknown. Hirtusneanoside A is effective against *Staphylococcus aureus* and *Bacillus subtilis*.



5.3 Unnamed dimeric xanthones

A number of unnamed xanthones have been reported, some due to the lack of full proof of their structure, some as parts of



patent applications (Fig. 4). In compound **125**, the lactone bridge appears to have been reductively transformed to a furan.

6 Conclusions

Dimeric xanthones are a structurally highly diverse group of compounds from various kingdoms of organisms, including plants, fungi and lichens. We have reviewed their origins, structural features, and, where applicable, bioactivities. Within the last five years, a number of monomeric xanthones have been prepared¹⁶ and the first synthesis of ergochromes has been achieved. We assume that in the next decade or so – with the further advent of arylation methods suitable for the unique requirements of these species – the synthesis of dimeric xanthones will be tackled with success.

7 Acknowledgements

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