

Structural diversity and biological activities of caged *Garcinia* xanthones: recent updates

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ABSTRACT

Caged xanthones are a class of natural compounds with approximately 200 members that are commonly isolated from the *Garcinia* genus in the Clusiaceae (formerly Guttiferae) family. They are often characterized by a notable 4-oxa-tricyclo[4.3.1.0^{3,7}]dec-2-one (caged) architecture with a common xanthone backbone. Because most caged xanthones have potent anticancer properties, they have become a target of interest in natural product chemistry. The unique chemical architectures and increasingly identified biological importance of these compounds have stimulated many studies and intense interest in their isolation, biological evaluation and mechanistic studies. This review summarizes recent progress and development in the chemistry and biological activity of caged *Garcinia* xanthones and of several compounds of non-*Garcinia* origin, from the years 2008 to 2021, providing an in-depth discussion of their structural diversity and medicinal potential. A preliminary discussion on structure-activity relationships is also provided.

Keywords: caged xanthone, gambogenic acid, isomorellin, neobractatin, anticancer, structure-activity relationship

1. INTRODUCTION

Garcinia is a pantropical genus of plants consisting of more than 400 species native to South America, Africa, India and East Asia [1, 2]. For their long history, the fruits, pericarp, epicarp and seeds of Garcinia species have been used in traditional medicine for treating ulcers, wounds, suppuration, dysentery and diarrhea [3, 4]. The chemical constituents in the extract of this genus are rich in polyphenols, flavonoids, polyprenylated acylphloroglucinols and xanthones [5]. Among them, the xanthones, particularly caged xanthones, have become a subject of intense interest among the scientific community because of their medicinal potential, particularly in cancer treatment. For instance, the most representative caged xanthone, gambogic acid, has been approved by the China Food and Drug Administration for phase II clinical trials involving intravenous treatment of lung cancer [6, 7].

Until 2008, approximately 100 natural caged Garcinia xanthones with diverse molecular connectivity had been discovered and isolated, mainly from Garcinia hanburyi [8], G. morella [9], G. gaudichaudii [10], G. scortechinii [11], and G. bracteata [12], according to our previous review [13]. Considerable progress on the phytochemistry of caged xanthones has since been made, and 96 new compounds have been isolated in the past decade

(Table 1). Most of the caged xanthones feature an unusual 4-oxa-tricyclo[$4.3.1.0^{3,7}$]dec-2-one (caged) scaffold [40] with a polyprenylated xanthone backbone, and show versatile pharmacological properties including anticancer [41, 42], antiviral [43], antibacterial [8], antiatherosclerosis [44] and neurotrophic activities [45].

Three key reviews exploiting the chemistry and biology of caged Garcinia xanthones have described [13, 46, 47], the most recent of which was published by the Reutrakul group in 2012 [47]. Numerous reviews by several research teams have focused on xanthone compounds, a small subset of which are caged xanthones [48-56]. In addition, the therapeutic potential of gambogic acid has recently been highlighted in several elegant reviews [57-60] and a book section [61]. Herein, this review does not attempt to overlap with those excellent published works but instead focuses on the structural diversity and biological investigation of newly isolated caged xanthone compounds from the years 2008 to 2021. Reports with keywords "caged xanthone," "caged polyprenylated xanthonoid," "caged Garcinia xanthone" and the names of several bioactive caged xanthones, such as "gambogic acid," "gambogenic acid," "isomorellin" and "neobractatin" found in major scientific databases such as Google Scholar, Web of Science, PubMed and ScienceDirect were compiled and analyzed.

 Table 1
 Isolated caged xanthones reported from 2008 to 2021 and associated cytotoxicity*

	Compound	Source	Cell lines	IC ₅₀ (μM)	Ref.
1	Gambogic aldehyde	G. hanburyi	P388	0.24	[14]
2	7-Methoxyisomorellinol	G. hanburyi	HeLa	1.2	[15]
3	7-Methoxygambogic acid	G. hanburyi	HeLa	0.93	[15]
4	7-Methoxyepigambogic acid	G. hanburyi	HeLa	0.85	[15]
5	Oxygambogic acid	G. hanburyi	HeLa	1.9	[15]
6	11,12-Dihydro-12-hydroxymorellic acid	G. lateriflora	HT-29	2.9 ^a	[16]
7	Isogaudichaudiic acid E	G. lateriflora	HT-29	2.6 ^a	[16]
8	Wightiic acid	G. wightii	A-375	4.7	[17]
9	16- O -methyl wightiic acid	G. wightii	_	_	[17]
10	Garcioligantone G	G. oligantha	A549 HeLa PC-3	3.7 3.6 4.3	[18]
11	Garcioligantone H	G. oligantha	_	_	[18]
12	Oliganthone B	G. oligantha	А549 НерG2 НТ-29 РС-3	3.9 4.5 4.8 4.6	[19]
13	Epiisobractatin	G. bracteata	A549 HL-60 HT-29 K562 WPMY-1	4.7 1.2 ^b 3.9 2.1 ^b 2.7	[20, 21]
14	13-Hydroxyepiisobractatin	G. bracteata	HL-60 K562	1.1 ^b 2.1 ^b	[20]
15	13-Hydroxyisobractatin	G. bracteata	HL-60 K562	1.1 ^b 2.1 ^b	[20]
16	Epicantleyanone B	G. oligantha	A549 HeLa PC-3	2.6 1.9 2.1	[18]
17	11-Methoxyepicantleyanone B	G. oligantha	-	_	[18]
18	Garcioligantone C	G. oligantha	HeLa	3.6	[18]
19	Garcioligantone D	G. oligantha	A549 HeLa K562 PC-3	4.4 2.6 4.6 2.7	[18]
20	Garcioligantone E	G. oligantha	_	-	[18]
21	Garcioligantone F or oliganthone E	G. oligantha	PC-3	4.6	[18, 22]
22	Gambogenific acid	G. hanburyi	HCT-116 HeLa HUVEC	4.5 1.8 2.5	[15, 23]
23	Gambogefic acid A	G. hanburyi	HeLa	2.1	[24]
24	12-Hydroxygambogefic acid A	G. hanburyi	HCT-116 MDA-MB-231	2.1 2.2	[25]
25	Gambogefic acid	G. hanburyi	HeLa	0.5	[15]

Table 1 | Continued

	Compound	Source	Cell lines	IC ₅₀ (μΜ)	Ref.
26	7-Methoxygambogellic acid	G. hanburyi	HeLa	0.9	[15]
27	Gambogellic acid A	G. hanburyi	HeLa	1.7	[24]
28	Gambogollic acid	G. hanburyi	A549 SMMC-7721	1.1 1.5	[26]
29	Epigambogollic acid	G. hanburyi	A549 SMMC-7721	1.6 1.8	[26]
30	Doitunggarcinone K	G. propinqua, G. bracteata	T98	3.5	[27, 28]
31	3-O-Geranylforbesione	G. hanburyi	HeLa	4.1	[15]
32	3- 0- Methylbractatin	G. propinqua, G. bracteata	A549 HCT-116 HL-60 MCF-7 SMMC-7721 SW480	0.7 (-)-isomer: 1.5 (+)-isomer: 3.4 0.2 0.3 0.2 0.5	[29-31]
33	Isogaudichaudiic acid B	G. lateriflora	_	-	[16]
34	Isogaudichaudiic acid	G. lateriflora	HT-29	3.2 ^a	[16]
35	Garcibractatin A	G. bracteata	A549 HeLa HT-29 PC-3 WPMY-1	2.0 2.6 1.1 2.9 0.8	[21]
36	16,17-Dihydroxygambogenic acid	G. hanburyi ^c	HeLa MDA-MB-231	4.3 2.1	[32]
37	22,23-Dihydroxydihydrogambogenic acid	G. hanburyi	A549 HCT-116 MDA-MB-231	5.0 0.7 2.7	[25]
38	Garcioligantone A or oliganthone C	G. oligantha	A549 HeLa PC-3	5.3 2.7 2.5	[18, 22]
39	Garcioligantone B	G. oligantha	-	-	[18]
40	Garcioligantone I	G. oligantha	_	-	[18]
41	Garcioligantone J	G. oligantha	A549 HeLa PC-3	5.9 3.7 3.4	[18]
42	Garcioligantone K	G. oligantha	HeLa PC-3	2.6 2.8	[18]
43	Garcioligantone L	G. oligantha	A549 HeLa K562	4.5 3.0 2.5	[18]
44	Oliganthone D	G. oligantha	_	_	[22]
45	Oliganthone F	G. oligantha	-	_	[22]
46	Gamboketanol	G. hanburyi	HeLa	3.8	[24]
47	8,8a-Dihydro-8-hydroxymorellic acid	G. hanburyi	HeLa	1.7	[15]

Table 1 | Continued

	Compound	Source	Cell lines	IC ₅₀ (μM)	Ref.
48	10α-Ethoxy-9,10-dihydromorellic acid	G. hanburyi	A549 HCT-116 HepG2 SK-BR-3	4.9 1.7 2.1 2.2	[33]
49	8,8a-Dihydro-8-hydroxygambogic acid	G. hanburyi	HeLa	0.64	[15]
50	8,8a-Dihydro-8-hydroxygambogic acid epimer	G. hanburyi	HeLa	0.63	[15]
51	Gambogic acid A	G. hanburyi	_	_	[34]
52	Epigambogic acid A	G. hanburyi	_	_	[34]
53	Gambogic acid B	G. hanburyi	_	_	[34]
54	Epigambogic acid B	G. hanburyi	_	-	[34]
55	10α-Butoxy gambogic acid	G. hanburyi	_	-	[34]
56	Isomoreollic acid	G. lateriflora	HT-29	1.9 ^a	[16]
57	Methyl 8,8a-dihydromorellate	G. hanburyi	_	-	[15]
58	Garcinolic acid	G. hanburyi	A549 HCT-116 HepG2 SK-BR-3	2.8 1.8 1.8 1.9	[33]
59	Doitunggarcinone J	G. propinqua	_	_	[27]
60	Gambogic acid C	G. hanburyi	_	-	[34]
61	Gambogic acid C epimer	G. hanburyi	_	-	[34]
62	Doitunggarcinone F	G. propinqua	_	-	[27]
63	Doitunggarcinone G	G. propinqua	_	-	[27]
64	Doitunggarcinone H	G. propinqua	_	-	[27]
65	Doitunggaricinone I	G. propinqua	_	-	[27]
66′	Garcibracteatone'	G. bracteata	K562	3.8 ^b	[20]
67	Garcibractone A	G. bracteata	_	_	[21]
68	Doitunggarcinone L	G. propinqua	_	_	[30]
69	Garcibractone B	G. bracteata	_	_	[21]
70	8,8a-Dihydro-8-hydroxygambogenic acid	G. hanburyi	A549 HCT-116 MDA-MB-231	3.2 2.0 0.5	[15]
71	10α -Ethoxy-9,10-dihydrogambogenic acid	G. hanburyi	HCT-116	3.6	[33]
72	8,8a-Dihydrogaudichaudione H	G. oligantha	A549	4.7	[18]
73	15-Acetoxy-8,8a-dihydrogaudichaudione H	G. oligantha	_	-	[18]
74	8-Hydroxy-8,8a-dihydrogaudichaudione H	G. oligantha	_	-	[18]
75	8-Ethoxy-8,8a-dihydrogaudichaudione H	G. oligantha	_	-	[18]
76	Garcibracteamone F	G. bracteata	_	_	[28]
77	8-Methoxy-8,8a-dihydrobractatin	G. bracteata	K562	3.4 ^b	[20]
78	8-Ethoxy-8,8a-dihydrobractatin	G. bracteata	HL-60 K562	4.5 ^b 2.1 ^b	[20]
79	Garcibracteamone G	G. bracteata	_	_	[28]

 Table 1 | Continued

	Compound	Source	Cell lines	IC ₅₀ (μM)	Ref.
80	Doitunggarcinone E	G. propinqua	-	-	[27]
81	Gambospiroene	G. hanburyi	_	_	[15]
82	Neobractatin	G. propinqua, G. bracteata	A549 HeLa HL-60 HT-29 K562 MCF-7 PC-3 SMMC-7721 SW480 WPMY-1	1.5, 3.5 0.9 0.8 ^b , 3.3 3.8 1.3 ^b 2.2 2.9 4.3 2.8 2.1	[29-31, 35]
83	3- <i>O</i> -methylneobractatin	G. propinqua, G. bracteata	HL-60 HT-29 MCF-7 PC-3 SMMC-7721 SW480 WPMY-1	2.9 3.4 1.9 4.5 3.9 2.8 1.8	[29-31]
84	8-Methoxy-8,8a-dihydroneobractatin	G. bracteata	K562	3.5 ^b	[20]
85	Neogaudichaudione H	G. oligantha	_	_	[18]
86	Garcineobractatin A	G. bracteata	_	_	[21]
87	Neobraclactone A	G. bracteata	HL-60 K562	0.50 ^b 0.40 ^b	[36]
88	Neobraclactone B	G. bracteata	HL-60 K562	0.50 ^b 0.86 ^b	[36]
89	Neobraclactone C	G. bracteata	_	_	[36]
90	Cochinchinoxanthone A	C. cochinchinense	_	_	[37]
91	Cochinchinoxanthone B	C. cochinchinense	-	-	[37]
92	Cochinchinoxanthone D	C. cochinchinense	-	-	[37]
93	Cochinchinoxanthone	C. cochinchinense, G. bracteata	HT-29 PC-3 WPMY-1	3.1 4.8 3.1	[21, 38]
94	Cochinchinoxanthone C	C. cochinchinense	-	-	[37]
95	Pruniflorone U	C. formosum	-	-	[39]
96	Pruniflorone T	C. formosum	-	-	[39]

*Cytotoxicities with $IC_{50} \ge 5 \ \mu M$ are not listed.

 $^{a}ED_{50}$ (in μ M) was recorded.

 ${}^{b}\text{GI}_{50}$ (in μ M) was recorded.

^cFrom microbial transformation of gambogenic acid (isolated from *G. hanburyi*).

2. STRUCTURAL DIVERSITY OF CAGED *GARCINIA* XANTHONES

The complex and unusual chemical architectures of caged xanthones have been attractive to chemists worldwide. Salient features of compounds belonging to this group of natural products include a common xanthone backbone with a D ring transformed into a unique 4-oxa-tricyclo[4.3.1.0^{3,7}]dec-2-one (caged) scaffold embedded with a highly substituted tetrahydro-furan core bearing three quaternary carbon centers (**Figure 1**). The B ring is typically a benzene ring substituted with hydroxyl and R groups (H, methyl or substituents shown in **Figure 2**) in various positions. Prenylation



3-hydroxy-3-methyl

but-1-envl

Figure 2 | Structures and abbreviations of side chains

isopreny

or geranylation of B rings produces bi- or tricyclic ring systems upon cyclization reaction. The D ring, in contrast, also has variable substituents with or without a double bond at the C-8/8a position. Together, the structural variability in A, B and D ring systems leads to diverse skeletons of caged xanthones, as shown by the chemical structure of gambogic acid.

homopreny

aeranv

Although most caged xanthones possess the 4-oxatricyclo[4.3.1.0^{3,7}]dec-2-one skeleton, few compounds with a rearranged caged backbone, known as neocaged compounds, have been reported (Figure 1). A representative compound from this class is neobractatin. Similarly, different substituents on B and D rings contribute to the structural diversity of these compounds. The structures of side chains and the abbreviations used herein are illustrated in Figure 2.

2.1 Caged xanthones with $\Delta^{8/8a}$ on the D ring 2.1.1 Presence of the A ring

Caged xanthones with an A ring and $\Delta^{8/8a}$ on the D ring comprise 29 new members (1–29, Figure 3) from the year 2008 to the present. Compounds 1 and 3–5 are four derivatives of gambogic acid isolated from the resin of *G. hanburyi* [14, 15]. Among them, gambogic aldehyde (1) has the same skeleton as gambogic acid (Figure 1), except that the -COOH group of gambogic acid is substituted by -CHO. A pair of epimers, 7-methoxygambogic acid (3, 13*R*) and 7-methoxyepigambogic acid (4, 13*S*),

possess a methoxy group at C-7, whereas oxygambogic acid (5) is characterized by a 3-hydroxy-3-metylbut-1enyl instead of prenyl R¹ substituent. The compound 7-methoxyisomorellinol (2), bearing an additional C-7 methoxy group, as compared with its parent compound, isomorellinol, has also been found in G. hanburyi [15]. Isolated from G. lateriflora, 11,12-dihydro-12-hydroxymorellic acid (6) features a hydroxy group at C-12 whose absolute configuration remains undefined, because of the limited quantity of the compound available, whereas isogaudichaudiic acid E (7) exhibits a 2-hydroxy-3-methylbut-3-enyl group at C-4 and an (E)-C-22/C-23 double bond [16]. Investigations on G. wightii, a species native to India, have led to the isolation of wightiic acid (8) and 16-O-methyl wightiic acid (9) from its leaves [17]. Both bear an unusual C-4 epoxy butyl group and C-5 cycloprop-2-envl moiety.

1.1-dimethyl-

prop-2-env

2-hydroxy-3-methyl

but-3-envl

In addition, 13 caged xanthones (10–22) bear an A ring on C-3/C-4 instead of C-2/C-3. Garcioligantones G (10), H (11) and oliganthone B (12) from twigs of G. oligantha are three examples of C-7 methoxylated caged xanthones with a pyran A ring [18, 19]. Another six compounds, epiisobractatin (13), 13-hydroxyepi-isobractatin (14, 125) and 13-hydroxyisobractatin (15, 12*R*) from G. bracteata [20, 21]; epicantleyanone B (16) and 11-methoxyepicantleyanone B (17) from G. oligantha [18]; and gambogenific acid (22) from G. hanburyi [15, 23], represent novel caged xanthones with



Figure 3 | Structures of caged xanthones with $\Delta^{8/8a}$ and an A ring.

a multisubstituted dihydrofuran unit as the A ring. Compounds **16** and **17** are 7-methoxylated, whereas **22** bears a geranyl moiety at C-2 and carboxylic acid functionality at C-5. Related compounds with a furan skeleton instead of dihydrofuran have been observed in garcioligantones C-F (**18-21**) [18]. Notably, garcioligantone F (**21**) has also been reported by another research group but given a different name: oliganthone E [22]. The other compounds, **23–25** from *G. hanburyi*, present a fused bicyclic ring attached to the caged xanthone skeleton in which gambogefic acid A (**23**) [24] and 12-hydroxygambogefic acid A (**24**) [25] are the hydrated and dihydroxylated version of gambogefic acid (**25**) [15], respectively. Beyond gambogellic acid,

which was discovered and isolated in 1996 [62], another three compounds (26–29) bearing a rare bridged bicyclic A ring system have been found. The compound 7-methoxygambogellic acid (26) has a methoxy moiety substituted at C-7 [18]; gambogellic acid A (27) is 12-hydroxylated [24] and gambogollic acid (28) and its epimer (29) are oxidized at C-38, with 28 bearing β -H at C-37 and 29 bearing α -H [26].

2.1.2 Absence of the A ring

Since 2008, 17 caged xanthones (**30–46**, Figure 4) with $\Delta^{8/8a}$ but not an A ring have been isolated and identified. The structural diversity of these compounds lies mainly in the different substitutions at the C-2, C-4 and C-5 positions. Doitunggarcinone K (**30**) [27, 28], 3-O-geranylforbesione (**31**) [15], 3-O-methylbractatin (**32**) [29-31] and garcibractatin A (**35**) [21] are alkoxylated

at C-3. Compounds 32 and 35 possess exactly the same chemical structure, except that the C-1 hydroxyl is replaced by -OMe in 35. Isogaudichaudiic acid B (33) and isogaudichaudiic acid (34), obtained from G. lateriflora, have a C-2 prenyl unit and two phenolic hydroxyls [16]. Two dihydroxylated derivatives of gambogenic acid, 16,17-dihydroxygambogenic acid (36) [32] and 22,23-dihydroxydihydrogambogenic acid (37) [25], display a characteristic C-2 dihydroxylated geranyl and C-4 dihydroxylated prenyl group, respectively. Another eight 7-methoxylated caged xanthones (38-45) from G. oligantha contain oxidized C-4 or C-5 substituents [18]. Structurally, the 3-hydroxy-3-methylbut-1-enyl group at C-5 is observed for garcioligantones A (38, repeatedly reported but named oliganthone C [22]) and B (39), whereas the prenyl moiety is shown for garcioligantones I-L (40-43) and oliganthones D (44), F (45).



Figure 4 | Structures of caged xanthones with $\triangle^{8/8a}$ but without an A ring.

Gamboketanol (46) from G. hanburyi is a rarely found pentaprenylxanthonoid likely to arise from decarboxvlation and rearrangement of gambogenic acid [24]. Additional cyclization forms between the C-5 side chain and C-6 carbonyl group.

2.2 Caged xanthones without $\Delta^{8/8a}$ on the D ring 2.2.1 Presence of an A ring

Detailed phytochemical investigations of Garcinia plants have identified several caged xanthones (47-61, Figure 5) with a hydroxyl or alkoxy group attached at the C-8 position. The compounds 8,8a-dihydro-8-hydroxymorellic acid (47) [15] and 10α -ethoxy-9,10-dihydromorellic acid (48) [33] exhibit a dimethyl substitution at the C-13 and C-8 hydroxyl in 47 but at the C-8 ethoxy group in 48. Instead of a dimethyl, one of the methyl groups at C-13 is substituted with a homoprenyl side chain in compounds 49-55, namely 8,8a-dihydro-8-hydroxygambogic acid and epimer (49/50) [15], gambogic acid A and epimer (51/52), gambogic acid B and epimer (53/54), as well as 10α -butoxygambogic acid (55) [34]. Accordingly, their C-8 positions are substituted by hydroxyl, methoxy,



80

Figure 5 | Structures of caged xanthones without $\Delta^{8/8a}$.

ethoxy or butoxy groups. Isomoreollic acid (**56**) features a dihydropyran A ring, C-8 methoxy moiety and (*Z*)-2methylbut-2-enoic acid at C-5 [16]. Compared with **56**, C-8 of methyl-8,8a-dihydromorellate (**57**) is not substituted, and the carboxylic acid is replaced by methyl ester functionality [15]. Isolated from the ethyl acetate extract of the resin of *G. hanburyi*, garcinolic acid (**58**) is highly similar to **51/52**, except for an oxidized C-5 group [33]. The structure of doitunggarcinone J (**59**) bears a trimethylated dihydrofuran A ring, C-8 methoxy group and C-5 3-hydroxy-3-metylbut-1-enyl side chain [27]. The novel gambogic acid C (**60**, α-Me) and epimer (**61**, β-Me) from *G. hanburyi* have two additional ring systems on the molecular scaffold [34].

2.2.2 Absence of an A ring

Twenty caged xanthones (62–81, Figure 5) without $\Delta^{8/8a}$ and an A ring have been obtained from Garcinia genus since 2008. Doitunggarcinones F-I (62-65), obtained from the stem bark extract of G. propingua, possess two methoxy groups and a phenolic hydroxyl group with a different C-5 moiety [27]. In contrast, garcibracteatone' (66') [20] and doitunggarcinone L (68) [30] have only one methoxy group but two phenolic hydroxyl groups. The name of garcibracteatone is somewhat confusing. because it represents two different compounds with entirely different molecular skeletons (Figure 6) [20, 63]. Garcibracteatone (66) was first isolated from the bark of G. bracteata in 2005 [63] and from the fruits of the same plant in 2020 [64]. Its biomimetic total synthesis was reported by the George research group in 2012 [65], and enantioselective total synthesis was reported in 2014 [66]. An unprecedented cage xanthone, compound 66', isolated from the leaves of G. bracteata, was also named garcibracteatone (designated garcibracteatone' herein) by Li and Hua in 2018 [20]. Compared with 68, garcibractone A (67) bears a hydroxyl group at C-8 instead of -OMe [21], whereas garcibractone B (69) has two methoxy groups at C-1 and C-3 and an ethoxy side chain at C-8.

Compounds **70** and **71** are two C-8 derivatives of gambogenic acid with a hydroxyl or ethoxy side chain at C-8 [15, 33]. The compound 8,8a-dihydrogaudichaudione



Figure 6 | Chemical structures of two garcibracteatones.

H (72), isolated from G. oligantha, is methoxylated at the C-7 position [18]. Xanthones 73-75 are the acetoxylated, hydroxylated or ethoxylated derivatives of 72, respectively, and were found from the same species [18]. Isolated from G. bracteata, garcibracteamone F (76) is reduced at C-8/C-8a, whereas garcibracteamone G (79) is epoxidized at C-8/C-8a [28]. The compounds 8-methoxy-8,8a-dihydrobractatin and 8-ethoxy-8,8a-dihydrobractatin (77 and 78) differ from 76 only in the substituent at C-8 [20]. The chemical structure of doitunggarcinone E (80) is presumably formed from dihydroxylation of the C-5 prenyl moiety, followed by intramolecular ketal formation to vield additional fused 5/6 rings [27]. A degraded and rearranged caged xanthone scaffold has been observed in gambospiroene (81) from G. hanburyi [15].

2.3 Neocaged xanthones

Xanthones with the neo scaffold have the ketone carbonyl group at C-5 instead of C-6, as observed in the classic caged xanthones. Only five natural compounds (82-86) from this classification have been isolated in the described period, along with three rearranged structures (87-89) (Figure 7). In continued efforts to search for bioactive components from tropical plants, two novel compounds (82-83) with a neocaged structure have been found from twigs of G. bracteata and G. propingua [29-31, 35]. As the name implies, 3-O-methylneobractatin (83) bears an -OMe substituent at C-3, as compared with neobractatin (82); whereas 8-methoxy-8,8a-dihydroneobractatin (84) has the -OMe at the C-8 position instead [20]. Neogaudichaudione H (85) bears a prenyl group at C-4 and a methoxy group at C-7 [18]. The structure of garcineobractatin A (86) is highly similar to that of neobractatin, except for an unsubstituted C-4 and a prenyl side chain at C-2 [21]. The unusual rearrangement of the neo skeleton leads to neobraclactones A-C (87-89) from G. bracteata [36]. They share a common octahydro-2H-1,3-dioxacyclopenta[c,d]inden-2-one framework; however, additional cyclization is observed in 89.

2.4 Caged xanthones of non-Garcinia origin

Beyond compounds from Garcinia, caged and neocaged xanthones (90-96, Figure 7) are also found in the Cratoxylum plant. Cochinchinoxanthone A (90) from Cratoxylum cochinchinense contains a simple caged skeleton with a methoxy group at C-7 [37]. Cochinchinoxanthones B (91) and D (92) are also substituted with 8a-OH and 8-OMe, respectively. Compared with 90, cochinchinoxanthone (93) has an unsubstituted C-7, a double bond between C8 and C-8a and an additional C-3 hydroxyl group [38]. Cochinchinoxanthone C (94) contains a C-8 -OMe group with a different relative configuration from that of 92. Moreover, pruniflorone U (95) is an uncommon example of a caged xanthone isolated from the roots of C. formosum ssp. pruniflorum [39]. It is an unprecedented ring-cleaved, rearranged caged xanthone with bond breakage at C-6/C-7. In



Figure 7 | Structures of neocaged and non-Garcinia caged xanthones.

addition, pruniflorone T (96), reported from *C. formo-sum*, bears a neocaged scaffold similar to that of 85 but does not have substituents at C-3 and C-4 [39].

3. BIOLOGICAL ACTIVITIES AND MECHANISMS

Beyond structural diversity, the promising biological activities of caged xanthones have made these compounds a prominent subject in the discovery and development of lead compounds, particularly as anticancer drugs. Over the past two decades, a vast number of caged xanthones have been subjected to cancer cell line screening. Close attention has been paid to several compounds with potent anticancer effects, such as neobractatin (82), gambogic acid (97), gambogenic acid (100), isomorellin (101) and forbesione (103) (Figure 8). Herein, the bioactivities and the mechanisms of action of these natural compounds are summarized and reviewed, along with the noteworthy derivatives of gambogic acid cluvenone (98) and DDO6101 (99, also known as MAD28).

3.1 Anticancer activities

3.1.1 General properties

The cytotoxicity and effects of many caged xanthones have been tested against multiple human cancer cell lines in the past decade. Frequently, the cytotoxicity of these compounds has been assessed against A549 human lung carcinoma and HeLa human cervical carcinoma cell lines. Very recently, a total of nine 7-methoxylated caged xanthones (10, 16, 19, 21, 38, 41-43 and 72) have demonstrated promising cytotoxicity (IC₅₀ < 8 μ M) against A549, HeLa and human prostate cancer PC-3 cell lines [18]. The other two compounds, 18 and 20, also display good activity against HeLa cells, at IC₅₀ = 3.6 and 5.9 μ M, respectively. Many caged xanthones including 35, 67-69, (-)-78, 82, 83, 86, 93, bractatin, 1-O-methylbractatin, isobractatin, 1-O-methylisobractatin, epiisobractatin and isoforbesione have also been tested against A549, HeLa, human colon adenocarcinoma HT-29, human prostate cancer PC-3 and WPMY-1 cell lines [21, 67]. The results have indicated that 67, 69 and 86 are inactive against all cell lines (IC $_{50}$ > 10 μM); 68, (-)-78 and 1-O-methylbractatin are active against only two or three cell lines, and the rest are cytotoxic (IC₅₀ < 10 μ M) in all cell lines. Mechanistically, the antiproliferative effect of isobractatin against PC-3 is caused by G0/G1 cell cycle arrest and apoptosis induction [67].

Furthermore, ten compounds (22, desoxymorellin, desoxygambogenin, gambogenin, isogambogenin, isogambogic acid, isogambogenic acid, morellic acid, a mixture of morellin and an inseparable regioisomer) are cytotoxic against four cancer cell lines (A549, HeLa, human colorectal carcinoma HCT116 and human hepatocellular carcinoma HepG2 cells), with IC₅₀ ranging from 0.64 to 14.23 μ M; moreover, all show potent



Figure 8 | Structures of several bioactive caged xanthones.

antiproliferative activity (IC₅₀ < 7 μ M) in human umbilical vein endothelial cells (HUVECs) [23]. Among them, remarkable antiangiogenic activity has been found for isogambogic acid, gambogenin and morellic acid tested in an in vivo zebrafish model. These compounds have lower toxicity than gambogic acid, according to death and heart rates. Thus, gambogenin has been strongly suggested to be a potential angiogenesis inhibitor, given its potent activity and absence of toxicity at 8–16 μ M.

In addition, six compounds (32, 82, 83, bractatin, isobractatin and 1-O-methylisobractatin) are active (IC₅₀ \leq 5.10 µM) against A549, human leukemia HL-60, human hepatocarcinoma SMMC-7721, human breast adenocarcinoma MCF-7 and human colon cancer SW480 cell lines [31]. Natural products 37, 70, gambogoic acid B, 30-hydroxygambogic acid, 30-hydroxyepigambogic acid, formoxanthone J, epiformoxanthone J, isomorellic acid, isogambogenic acid, gambogenin and gambogellic acid have also been found to be cytotoxic, with $IC_{50} < 8 \ \mu M$ against A549, HCT116 and triple-negative breast cancer MDA-MB-231 cell lines [25]. Caged xanthones 21, 38 and 44 have been shown to decrease the viability of A549 cells by inducing apoptosis [22]. Compound 24 has an IC_{50} = 2.05 and 2.20 μ M against HCT116 and MDA-MB-231 cell lines, respectively; whereas 36 has activity at 2.09 µM against the triple-negative breast cancer MDA-MB-231 cell line [25]. Meanwhile, (+)-/(-)-82, (+)-/(-)-83 and (-)-bractatin have potent activity (IC₅₀ = 1.47-7.02 μ M) against HCT116 cells [30], whereas 30 and (-)-59 have only weak activity (IC₅₀ = 23.95 and 14.23 μ M, respectively), as compared with that of the positive control doxorubicin, with IC_{50} value of 9.74 μ M [27].

All eight compounds (48, 58, 71, deoxygaudichaudione A, gambogenic acid, desoxygambogenin, hanburin and desoxymorellin) exhibit good activity against A549, HCT116, HepG2 and human breast cancer SK-BR-3 cell lines, with an IC₅₀ < 10 μ M [33]. Compound **12**, gaudichaudione H and cantleyanone A display IC₅₀ values of less than 9 μ M in A549, HepG2, PC-3, human colorectal adenocarcinoma HT-29 and human colorectal adenocarcinoma HL-7702 cell lines [19]. Compounds **28**, **29**, gambogellic acid and its epimer have been reported to show excellent cytotoxicity against A549 and human hepatocarcinoma SMMC-7221 cells, with IC₅₀ values ranging from 1.07 to 2.10 μ M, as compared with that of the positive control cisplatin, with IC₅₀ values of 9.38 and 11.4 μ M, respectively [26].

Cytotoxic studies in HeLa human cervical cell lines have revealed several bioactive compounds with $IC_{50} < 9 \ \mu$ M, including 2-5, 22-23, 25-27, 31, 36, 46-47, 49-50, 70, 81-82, isobractatin, gaudichaudione H and gambogenic acid [15, 24, 32, 68, 69]. Caspase 3 activation, as evidenced by poly(ADP-ribose) polymerase (PARP) cleavage, has been observed in HeLa cells treated with 82 [69], gaudichaudione H [68], and isobractatin [69], thus suggesting that apoptosis induction is involved in their antiproliferative effect. Gaudichaudione H and isobractatin have also been reported to increase the sub-G1 fraction in A549 lung carcinoma cells [69]. Furthermore, 82 and isobractatin have been shown to suppress autophagic flux in both cell lines [69]. These compounds convert light chain 3B-I (LC3B-I) to LC3B-II, increase production of p62 proteins at high concentration (4 μ M) and induce puncta formation.

Moreover, **13–15**, **32**, **66**['], **77–78**, **82–84**, **87–88**, bractatin, isobractatin, neoisobractatins A and B, as well as 1-O-methyl-77, have been shown to possess growth-in-hibitory activities against HL-60 (promyelocytic leukemia) and K562 (chronic myeloid leukemia) cell lines, with GI_{50} values between 0.2 and 8.8 μ M [20, 63]. When

tested against K562 cells, potent activity has also been observed for 43, with an IC₅₀ = 2.5 μ M [18]. Compound 1 inhibits the growth of murine leukemia P388 and P388/ADR cells, with IC₅₀ values of 0.243 and 7.60 μ M, respectively [14]. Compounds 8, 9, gaudichaudic acid E and isogaudichaudic acid E show strong inhibitory activity, with IC₅₀ values from 4.7 to 9.7 μ M against MCF-7, human melanoma A-375 and human breast carcinoma SKBR-3 cells [17]. Moreover, investigations of 30 and 82 have shown marked inhibitory activity against HepG2, MCF-7 and T98 (human glioblastoma multiforme) cell lines, with IC_{50} between 3.21 and 6.27 μM [28]. Cochinchinone C and a mixture of 95/96 have excellent activity against the MCF-7 cell line [39]. The compounds (-)-6, (-)-7, (-)-34, (-)-56, 93 and (-)-morellic acid exhibit significant cytotoxicity toward the HT-29 colon cancer cell line, with an ED_{50} less than 6 μM [16, 38]. Compound 93 also displays good activity (IC₅₀ = 3.3 μ M) in mitochondrial transmembrane potential assays [38].

In 2015, 64 Garcinia compounds including 13 caged xanthones were screened against a wide spectrum of cancer cell lines [70]. Although all compounds demonstrated excellent activity toward all cell lines (except H1573 cells), with IC_{50} values less than 5 $\mu M,$ the lowest IC₅₀ was observed toward the NCI-H1650 cell line. Further study revealed that 33-hydroxyepigambogic acid and 35-hydroxyepigambogic acid were the most potent inhibitors of NCI-H1650 cell growth and colony formation [70]. Both compounds dose-dependently increased the activity of caspase 3 and 7 in NCI-H1650 cells, and cell cycle analysis showed significant accumulation of S or G2/M phase cells. Several proapoptotic BH3-only genes, such as PUMA, Noxa and cell death involved p53-target (CDIP), were upregulated. Inhibition of Janus kinase (JAK) activity, particularly that of JAK2, was observed, thus affecting the expression and phosphorylation of signal transducer and activator of transcription 3 (STAT3), as shown by western blot analysis. Phosphorylation of extracellular signal-regulated kinase (ERK) and protein kinase B (Akt) was also inhibited. The findings suggest that both compounds exert cytotoxicity by targeting the JAK-STAT signaling pathway.

3.1.2 Gambogic acid and derivatives

Gambogic acid (97, Figure 8), extracted from the resin of *G. hanburyi*, demonstrates significant cytotoxicity against several cancer types including but not limited to breast, lung, liver, colorectal and prostate cancers, and melanoma [71-76]. This compound exerts antiproliferative, antimetastatic and antiangiogenic properties through apoptosis induction [77, 78], reactive oxygen species (ROS)-induced endoplasmic reticulum (ER) stress [72, 79], autophagy [80, 81] and modulation of various cellular pathways, such as nuclear factor κ -light-chainenhancer of activated B cells (NF- κ B) [82, 83], Akt/mammalian target of rapamycin (mTOR) [80, 84], phosphoinositide 3-kinase (PI3K)/Akt [75, 76], c-Jun N-terminal kinase (JNK) [79, 85] and mitogen-activated protein

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kinase (MAPK) [86, 87] pathways. Moreover, it enhances chemosensitivity or synergistically potentiates the anticancer effects of various chemotherapeutic agents, such as docetaxel [88, 89], doxorubicin [85, 90], chloroquine [91], cisplatin [86], vorinostat [92], 5-fluorouracil [93] and others [94, 95].

Despite having promising anticancer properties and high selectivity for cancer cells over normal cells [96], the low aqueous solubility of gambogic acid hinders its potential clinical applications. Continual efforts to elucidate the minimum pharmacophore of gambogic acid for subsequent enhancement of drug-like properties has revealed promising candidates worthy of further study. In 2007, the Theodorakis group discovered that cluvenone (98, Figure 8), a synthetic gambogic acid derivative, is a potential inhibitor of angiogenesis [97]. It exerts cytotoxicity against HUVECs, with $IC_{50} = 1.38 \mu M$, through apoptosis induction [97]. The antiproliferative properties of cluvenone against cancer and multidrug resistant cells have been supported by evidence from a NCI-60 human tumor cell line screen indicating GI₅₀ values between 0.1 and 2.7 µM [98]. This compound also induces apoptosis in human T-cell acute lymphoblastic leukemia (ALL) CEM cells (EC₅₀ = 0.25 μ M) and has demonstrated selective toxicity against primary B-ALL cells (IC₅₀ = 1.1 μ M) compared with peripheral blood mononuclear cells from normal donors (IC₅₀ = 5.2 μ M) [98]. The antiproliferative effect of cluvenone bypasses the mechanisms of multidrug resistance conferred by P-glycoprotein, and its inhibitory activity against HL-60 and HL-60/ADR cell lines does not significantly differ [97-99]. Gene expression profiling has suggested that cluvenone activates the MAPK pathway along with its downstream nuclear factor-erythroid factor 2-related factor 2 (Nrf2)-mediated oxidative stress response signaling pathway, on the basis of observations of upregulation of heat shock protein family A member 1A (HSPA1A), HSPA8, p38 and other stress-associated genes [98]. Cluvenone has also been demonstrated to induce apoptosis by dissipating the mitochondrial membrane potential in HeLa cells and promoting the release of cytochrome C from mitochondria to the cytoplasm in CEM cells [100]. The resultant activity of caspase 9 and caspase 3 is therefore enhanced [100, 101]. At low micromolar concentrations, this compound induces degradation of Hsp90 client proteins and PARP cleavage [101]. In addition, cluvenone induces formation of ROS in the PC3 human prostate cancer cell line, thus inducing cell stress and apoptosis [98].

DDO6101 (102), also known as MAD28, is another gambogic acid derivative that has attracted wide-spread attention because its activity is comparable to that of gambogic acid. The antiproliferative activity of DDO6101 against HepG2 cell lines is highly similar to that of gambogic acid ($IC_{50} = 2.37$ and 2.40 μ M, respectively) [102]. Mechanistic studies have revealed that DDO6101 induces G2/M phase cell cycle arrest and promotes apoptosis by decreasing the expression

of Bcl-2 and procaspase 3 while increasing the activity of caspase 3. Screening of (+)-, (-)- and (±)-DDO6101 against a panel of breast cancer cell lines (SKBR3, BT474, MDA-MB-231, MCF-7, MDA-MB-468 and MARY-X) has demonstrated potent activity with IC₅₀ ranging from 0.6 to 3.8 μ M, with a dose-dependent increase in cleaved-PARP, caspase 3 and caspase 7 [103]. Studies of other bioactive cluvenone derivatives are underway [104-109].

3.1.3 Gambogenic acid

Gambogenic acid (100, Figure 8) was first discovered and isolated from G. hanburyi in 1996 [62]. Extensive research has focused on the mechanistic study of this natural compound, because it demonstrates potent anticancer properties by modulating various cellular signaling pathways in cancer cells (Figure 9). In vitro studies against many cell lines, including lung cancer, nasopharyngeal cancer, colon cancer, glioblastoma, breast cancer, gastric cancer and melanoma, have shown that gambogenic acid inhibits cancer cell proliferation mainly through mitochondria-mediated apoptosis induction and cell cycle arrest at G0/G1 phase [110-118]. The apoptotic effect of gambogenic acid is also associated with decreased protein levels of p38, MAPK and p-ERK1/2 [119, 120], and the inactivation of Akt signaling pathway [113, 115, 117].

In the B16 mouse melanoma cell line, gambogenic acid not only induces apoptosis by inhibiting the expression of p-PI3K, p-Akt and p-mTOR [121], but also promotes autophagy via upregulation of the ROS/ sirtuin 3 (SIRT3)/adenosine monophosphate-activated protein kinase (AMPK) pathway [122]. It also suppresses

metastasis of melanoma cells by targeting proteins associated with epithelial-mesenchymal transition [118] and promoting ferroptosis through activation of p53/solute carrier family 7 member 11 (SLC7A11)/glutathione peroxidase 4 (GPX4) [123]. Moreover, gambogenic acid degrades cancerous inhibitor of protein phosphatase 2A (CIP2A) in hepatocellular carcinoma via the ubiquitin-proteasome pathway [124]. Downregulation of its associated downstream molecules, that is, c-Myc and p-Akt, has been observed. Furthermore, gambogenic acid exerts anti-inflammatory and antiapoptotic effects against in vivo acute hepatotoxicity by regulating the PI3K/Akt and NF-κB signaling pathways [125]. Its inhibitory effect on NF-kB signaling has also been found to suppress growth and metastasis of bladder cancer cells [126]. In HepG2/Adr cells, gambogenic acid downregulates P-glycoprotein expression, possibly via inhibition of NF-κB and MAPK pathways [127].

In addition, gambogenic acid has been found to promote autophagy of lung cancer cells through glycogen synthase kinase 3 beta (GSK3 β) activation and suppression of Akt/mTOR [128]. Gambogenic acid suppresses the acidification of lysosomes, and hinders the fusion of autophagosomes and lysosomes, and consequently lysosomal degradation and autophagy induction [129]. This compound exerts antiproliferative effects against erlotinib-resistant non-small-cell lung cancer cell lines and patient-derived xenografts by downregulating the fibroblast growth factor receptor (FGFR) signaling pathway [130]. Synergistic growth inhibition of lung cancer cells has also been observed after combined treatment with gambogenic acid and



Figure 9 | Molecular targets of gambogenic acid.

5-fluorouracil, in which the activation of cancer cell death is caused by both caspase-independent necroptosis and caspase-dependent apoptosis [131].

The Li research group has shown that gambogenic acid triggers ER stress by activating volume-sensitive outwardly rectifying chloride channels, thus leading to apoptosis of human nasopharyngeal carcinoma CNE-2Z cells [132]. The ER stress induced by gambogenic acid and caused by increased ROS production also activates inositol-requiring enzyme-1 α (IRE1 α), apoptosis signal regulating kinase 1 (ASK1) and downstream JNK, thus causing Noxa-mediated apoptosis [133]. In colorectal cancer, gambogenic acid shows good antiproliferative activity by inducing ER stress via downregulation of the Aurora A pathway, which plays a crucial role in cell division [134].

Furthermore, gambogenic acid has excellent antitumor activity against hypoxic multiple myeloma cells by decreasing hypoxia-inducible factor 1-alpha (HIF-1 α) accumulation and STAT3 phosphorylation, which play important roles in the modulation of miR-21/phosphatase and tensin homolog (PTEN) expression [135] Combination treatment with gambogenic acid/bortezomib has significant synergistic effects on apoptosis induction in MM.1S multiple myeloma cells through G2/M cell cycle arrest and modulation of p53/ROS/p38 MAPK signal transduction [136]. Moreover, gambogenic acid potentiates adriamycin-induced apoptosis in MCF-7/ADR cells via G0/G1 arrest and downregulation of the PTEN/PI3K/Akt pathway [137]. Unfortunately, gambogenic acid induces a thiol-dependent heat shock response by disrupting the interaction between heat shock protein 90 (HSP90) and heat shock factors (HSF) 1 or 2 [138], which are associated with cytoprotection of cancer cells [139], thus suggesting a need for further investigations.

3.1.4 Isomorellin and forbesione

The biological activities and mechanisms of action of isomorellin (101) and forbesione (103) against cholangiocarcinoma has been a major research area for Reutrakul's research team since 2010 (Figure 8) [140-144]. Four caged xanthones, gambogic acid, 101, isomorellinol (102) and 103, have been tested against two cholangiocarcinoma cell lines, KKU-100 and KKU-M156 [140]. All compounds displayed potent cytotoxicity, with IC_{50} from 0.02 to 2.64 μ M. The compounds induced apoptosis in both cell lines partly by downregulating Bcl-2 and survivin proteins while upregulating Bax protein and apoptosis-inducing factor (AIF) (Figure 10). Consequently, activation of caspase 3 and 9, and an increase in the Bax/Bcl2 ratio were observed, thus suggesting that the anticancer properties of these four compounds are mediated by mitochondrial-dependent apoptosis.

Further mechanistic study has shown that isomorellin induces cell cycle arrest at G0/G1 phase via regulation of the p53 and NF- κ B signaling pathways, thereby promoting apoptosis in both cells [141]. This compound increases the expression of p53 tumor suppressor



Figure 10 | Proposed anticancer mechanisms of isomorellin.

protein and cyclin-dependent kinase inhibitors p21 and p27, while decreasing the expression of cyclin D1, cyclin E, and the cyclin-dependent kinases Cdk2 and Cdk4. Moreover, isomorellin decreases cell viability, migration and invasion of KKU-100 cells [142] This compound also blocks the activity of focal adhesion kinase (FAK), protein kinase C (PKC) and the downstream p38-MAPK pathway, thereby upregulating the expression of IkB α , an inhibitor of NF- κ B. Consequently, the translocation of NF- κ B/p65 into the nucleus is inhibited, thus suppressing the expression of matrix metalloproteinase-2 (MMP-2), urokinase-type plasminogen activator (uPA) and cyclooxygenase-2, (COX-2) which are associated with invasion and metastasis of cancer cells.

Synergistic effects of isomorellin and forbesione with doxorubicin on growth inhibition and apoptosis induction have been assessed in KKU-100, KKU-M139, KKU-M156 human cholangiocarcinoma and Chang liver cell lines [143]. Either of the caged xanthones alone selectively inhibited the growth of cholangiocarcinoma cells but not Chang cells. A combination of isomorellin and doxorubicin exhibited enhanced bioactivity on KKU-M139 and KKU-M156 cells, whereas the forbesione/ doxorubicin pair displayed synergistic inhibitory activity on KKU-100 and KKU-M139 cells. An in-depth study has revealed that the combined treatments induce apoptosis by stimulating the expression of Bax/Bcl-2 ratio, caspase 9 and caspase 3, while downregulating the expression of survivin, procaspase 9 and procaspase 3. Inactivation of the NF-κB pathway—as evidenced by decreased NF-κB/ p65 expression and phosphorylated $I\kappa B\alpha$ levels, along with suppression of multidrug resistance-associated protein 1 (MRP1) protein expression—also plays a pivotal role in their synergistic growth-inhibitory effects.

Forbesione has been demonstrated to inhibit the growth of the Ham-1 hamster cholangiocarcinoma cell line and Ham-1 allografts in a hamster model, with no observed in vivo toxicity or adverse effects [144]. This

compound arrests the cell cycle at S phase through decreasing the protein expression of cyclin A, cyclin E and Cdk2, as well as promoting the upregulation of p21 and p27. Multiple key signaling pathways participate in forbesione-mediated apoptosis. Increased expression of Fas, Fas-associated death domain (FADD) and caspase 3, accompanied by decreased procaspase 8 and procaspase 3 levels, has indicated the involvement of the death receptor pathway. The mitochondrial pathway is also triggered by induction of Bax, $I\kappa B\alpha$ and caspase 9, with suppression of the protein levels of Bcl-2, procaspase 9 and NF- κ B/p65. A plausible ER pathway has been proposed in view of the increased expression of caspase 12.

3.1.5 Neobractatin

As demonstrated in Section 3.1.1, neobractatin (82) exhibits significant cytotoxicity against various cancer cell lines. Testing of this natural compound against seven cancer cell lines (HeLa, A549, MCF-7, SH-SY5Y, PC-3, K562 and U937) by our group has indicated potent cytotoxicity, with IC₅₀ values <4 µM within 24 h; moreover, an in vivo study on HeLa mouse xenograft models has exhibited decreased tumor growth and weight, with no observed toxicity [145]. A detailed mechanistic study has revealed that neobractatin induces G1/S phase arrest by decreasing E2F/DP heterodimeric transcription factor 1 (E2F1) activity (Figure 11). Simultaneously, neobractatin upregulates growth arrest and DNA damage-inducible protein 45 alpha (GADD45 α), decreases the expression of cyclin B1 and disrupts the mitotic spindle, thus arresting synchronized HeLa cells in G2/M phase.

Moreover, neobractatin has been demonstrated to be active against breast and lung cancer cell metastasis [146]. This compound inhibits metastasis both in vitro and in vivo, partly via upregulation of muscleblind-like protein 2 (MBNL2), which has diminished expression in breast or lung cancer tumor tissue. An increase in MBNL2



Figure 11 | Neobractatin-induced G1/S and G2/M cell cycle arrest.

in turn alters the modulation of p-Akt/epithelial-mesenchymal transition signaling pathway and the expression of related proteins such as vimentin, cofilin and MMP-2.

The composition of bractatin and neobractatin has been found to have inhibitory activity against throat cancer cells (Hep-2 and FaDu) and Hep-2 xenograft models [147]. These compounds promote apoptosis by regulating multiple signaling pathways involved in ER stress-induced apoptosis, mitochondria-mediated apoptosis and inhibition of the Akt/GSK-3β/Bad pathway.

3.2 Anti-inflammatory activities

The compounds 8-methoxy-8,8a-dihydrobractatin (77), bractatin, isobractatin and neobractatin (82) inhibit nitric oxide production with a lower IC₅₀ (1.22–8.96 μ M) than that of the positive control dexamethasone [28] Furthermore, gambogenic acid and gaudichaudione H display good anti-inflammatory effects in macrophages via inhibition of the NF-kB and MAPK signaling pathways [148, 149]. The latter compound has also been tested in colitis mouse models induced by dextran sodium sulfate and found to inhibit the phosphorylation of $\text{AMPK}\alpha$ and proline-rich Akt substrate of 40 kDa (PRAS40) [149]. Pro-inflammatory mediators and cytokines such as interleukin-6, COX-2, tumor necrosis factor- α (TNF- α) and inducible nitric oxide synthase (iNOS) are downregulated in LPS-stimulated RAW 264.7 cells and colon tissues in mouse models. In addition, gambogenic acid has been shown to decrease hypertrophic scar formation by controlling the local inflammatory response, neoangiogenesis and growth factor expression during the wound healing process [150].

3.3 Miscellaneous activities

Only a small number of caged xanthones have been tested for potential bioactivity beyond anticancer effects. Moreollic acid, 10-methoxygambogenic acid, gambogoic acid, morellic acid, gambogenic acid and **93** have been found to competitively inhibit ($IC_{50} = 0.47-6.6 \mu M$) protein tyrosine phosphatase 1B (PTP1B) [37,

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151], an important modulator of insulin signaling, and a potential target for the treatment of cancer and obesity [152-154], These compounds, along with 51–54, 71, 10 α -hydroxygambogic acid, desoxymorellin, gambogin, gambogellic acid and desoxygambogenin, also exhibit inhibitory activity against α -glucosidase [34, 155].

Morellic acid has antibacterial effects against methicillin-resistant *Staphylococcus aureus* USA300, with an MIC value of 12.5 μ M, but its toxicity hinders its future applications and may warrant structural modification [156]. Two synthetic cluvenone derivatives (CR135 and CR142), which exist as triphenylphosphonium salts, exhibit antimalarial activity against *Plasmodium falciparum* [157].

4. STRUCTURE-ACTIVITY RELATIONSHIPS

The intact caged D ring plays an essential role in maintaining the cytotoxicity of caged xanthones against cancer cell lines (Figure 12) [97]. This scaffold has antiproliferative effects on cancer cells and inhibits IkB kinases (IKKs), the key regulators of NF-κB activation [158]. Replacing the oxa-caged D ring with an aza-caged ring is also feasible with inclusion of a hydrophobic moiety on the nitrogen atom; this modified compound shows significant increase in cytotoxicity and IKK^β inhibitory activity [105]. The unsaturated double bond at C-8/8a appears to significantly contribute to apoptosis induction and antitumor activity [99, 109, 159, 160]. Additionally, the Theodorakis and You research groups have independently reported that the gem-dimethyl groups at C-28 and C-33 are crucial for these compounds' cytotoxicity [97, 161]. However, the presence of a C-33 gem-dimethyl has been found to be inessential for cytotoxicity against HepG2 cells; substitution of this moiety by hydrogen atoms is tolerated but oxidation of C-33 to carbonyl have been found to decrease the activity [104]. In a later study, You has proposed that the C-28 geminal dimethyl group contributes to Hsp90 activity, but the C-33 geminal-dimethyl decreases its activity [109].



Figure 12 | General SARs of caged xanthones on antitumor activity

Modifications of the C-27/28 double bond to a dihydroxyl or epoxy group enhance cytotoxicity against the A549, MCF-7 and BGC-823 cell lines, among which the epoxy group confers better inhibition than the dihydroxyl group, except against A549 cells [162]. Functionalization of the carboxylic acid moiety at C-29 appears to be tolerated, thus indicating that this group could be modulated to improve physiochemical properties [99, 109, 159]. Introduction of an electron withdrawing group at C-29 improves the inhibitory activity against A549 cells [162]. Moreover, introduction of a hydrophilic amine side chain at C-31 maintains cytotoxicity while increasing water solubility and cell permeability, whereas a hydrophobic amine moiety diminishes the bioactivity [104].

In contrast, the aromaticity of the A ring must be maintained, because oxidation of the A ring to quinone leads to a complete loss of activity [163]. The presence of a hydroxyl group at the C-1 position is an important structural feature for antitumor activity [102, 161, 163] and has been found to enhance IKK β inhibitory activity [158]. Activity is preserved when the hydroxyl group is methylated or acylated [159], but replacing the hydroxyl group with a prenyl group is unfavorable [161]. The positions of hydroxyl, prenyl or fused pyran moieties on C-2, C-3 and C-4 have no significant correlation with antitumor activity [104, 158, 161]. Meanwhile, a free hydroxyl group at C-3 decreases the cytotoxicity [163], but converting it to methoxymethyl ether enhances the activity [162].

Furthermore, modification of the double bond at C-16/17 or C-22/23 to an epoxy group, and conversion of the methyl group at C-18 or C-25 into $-CH_2OH$ or C-18 methyl into -CHO provides better inhibitory activity against cancer cells [164, 165]. Introduction of hydrophilic amines at C-16, C-18 or C-25 increases the antitumor activity and improves drug-like properties [166-168].

Regarding antibacterial properties, Jiarpinitnun et al. have reported that the C-29 carboxylic acid moiety is crucial for growth inhibition against methicillin-resistant *Staphylococcus aureus* USA300. Switching this functionality to less polar substituents diminishes the antibacterial activity and may increase the cytotoxicity [156]. In contrast, the prenyl side chain at C-13 is not necessary for bioactivity.

5. CONCLUSIONS AND FUTURE PERSPECTIVES

Since the isolation of morellin in 1937, a total of 194 caged xanthones have been identified mostly from *Garcinia* species, and several have been identified from *Cratoxylum* plants. Modifications on the B and D rings of the caged xanthone scaffold by additional cyclization and/or side chain substitutions expand the structural diversity of this family of compounds. In general, caged xanthones can be characterized on the basis of the presence or absence of $\Delta^{8/8a}$ and the A ring. The study and analysis of caged xanthones according to

this classification is highly important, because available structure-activity relationship (SAR) studies have suggested that the double bond between C-8 and C-8a is essential for cytotoxicity, and the absence of an A ring does not affect biological activity [102, 159, 160]. On this basis, systematic SAR evaluations of other substitutions are warranted.

The unique architecture of caged xanthones results in potent bioactivities, particularly anticancer properties. Among them, gambogic acid has received the most attention, and its mechanism of action has been widely studied. The other natural compounds, although subjected to cancer cell line screening, have not been more deeply investigated. Only isomorellin and forbesione have been studied mechanistically against cholangiocarcinoma, and studies have also investigated gambogenic acid, neobractatin and gambogic acid derivative, i.e., cluvenone. In-depth research to discover optimal lead caged xanthones with excellent drug-like properties remains necessary. Discovery of other potential biological properties of caged xanthone beyond anticancer properties, and evaluation of toxicity should also be undertaken.

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