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Review

Naturally occurring *seco*- and *nor*-polycyclic polyprenylated acylphloroglucinols: distribution, structural diversity, and biological activityYulin Duan^{a,b}, Ying Tang^{d,*}, Changxing Qi^{c,*}, Yonghui Zhang^{c,*}^a Department of Pharmacy, Traditional Chinese and Western Medicine Hospital of Wuhan, Tongji Medical College, Huazhong University of Science and Technology, Wuhan 430030, China^b Department of Pharmacy, Wuhan No. 1 Hospital, Wuhan 430022, China^c Hubei Key Laboratory of Natural Medicinal Chemistry and Resource Evaluation, School of Pharmacy, Tongji Medical College, Huazhong University of Science and Technology, Wuhan 430030, China^d Department of Pharmacy, Tongji Hospital, Tongji Medical College, Huazhong University of Science and Technology, Wuhan 430033, China

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ABSTRACT

Polycyclic polyprenylated acylphloroglucinols (PPAPs) represent a distinct subclass of specialized metabolites predominantly found in the plant kingdom, particularly within the Guttiferae (Clusiaceae) family. These compounds exhibit remarkable structural diversity and a wide range of biological activities. *Seco*- and *nor*-PPAPs, two unique variants of PPAPs with diverse skeletal structures, have been extensively investigated. As of June 2023, 200 compounds have been isolated from four genera, with *Hypericum* being the primary source. Notably, 115 of these compounds were identified in the past four years, indicating a significant increase in research activity. *Seco*- and *nor*-PPAPs can be categorized into six main subgroups based on the original PPAP scaffolds. Biological studies have revealed their potential in various therapeutic applications, including anti-cancer, anti-inflammatory, hepatoprotective, anti-Alzheimer's disease (anti-AD), multidrug resistance (MDR) reversal, anti-depressant, neuroprotective, and immunosuppressive effects. This review provides a comprehensive overview of the occurrence, structures, and bioactivities of natural *seco*- and *nor*-PPAPs, offering valuable insights for the further development of PPAPs.

1. Introduction

Polycyclic polyprenylated acylphloroglucinols (PPAPs), characteristic secondary metabolites of plants, are primarily derived from the genera *Hypericum* and *Garcinia*. These compounds consist of highly oxygenated acylphloroglucinol-derived cores substituted with multiple isoprene units¹⁻³. Over 1000 naturally occurring PPAPs have been reported and can be structurally categorized into three subgroups based on their chemical skeleton characteristics, which are directly related to their biogenesis: (i) bicyclic polyprenylated acylphloroglucinols (BPAPs), (ii) caged PPAPs with adamantane and homoadamantane cores, and (iii) other PPAPs, including spirocyclic PPAPs with an octahydro-spiro[cyclohexan-1,5'-indene]-2,4,6-trione core and complex PPAPs derived from intramolecular [4 + 2] cycloaddition between the acylphloroglucinol core and side chains (Fig. 1)¹⁻⁴. The diverse structures and intriguing biological activities of these compounds have garnered significant interest from pharmacologists and synthetic chemists^{1, 5-15}. Hyperforin, first isolated from *H. perforatum*, exhibits profound biological applications, notably its anti-depressant activity, which is responsible for the use of *H.*

perforatum in treating depression¹⁶⁻²².

Seco- and *nor*-PPAPs, distinct categories of PPAPs, exhibit unique oxygen or carbon ring structures, which serve as crucial sources of diverse molecular frameworks. These PPAPs demonstrate significant structural variations derived from conventional PPAPs through varying degrees of cleavage or degradation of acylphloroglucinol cores or side chains, followed by rearrangement (Fig. 1). Such structural diversity underpins their wide range of biological activities, including anti-cancer, anti-inflammatory, hepatoprotective, anti-Alzheimer's disease (anti-AD), multidrug resistance (MDR) reversal, anti-depressant, neuroprotective, and immunosuppressive properties²³⁻²⁶. As of June 2023, approximately 200 naturally occurring *seco*- and *nor*-PPAPs with multiple skeletal structures have been documented in the literature.

While several general reviews have explored PPAPs with respect to their structural characteristics, bioactivities, and biosynthetic and synthetic methods, these works have predominantly focused on representative PPAPs, with limited attention given to the unique *seco*- and *nor*-PPAP subclasses^{1, 3, 5, 6, 27, 28}. In response to this gap, the present review aims to systematically profile the recent advances in the distribution, structural diversity, biological activities, and biosynthetic pathways of naturally occurring *seco*- and *nor*-PPAPs, emphasizing novel skeletons reported between January 1988 and June 2023. For clarity and concise-

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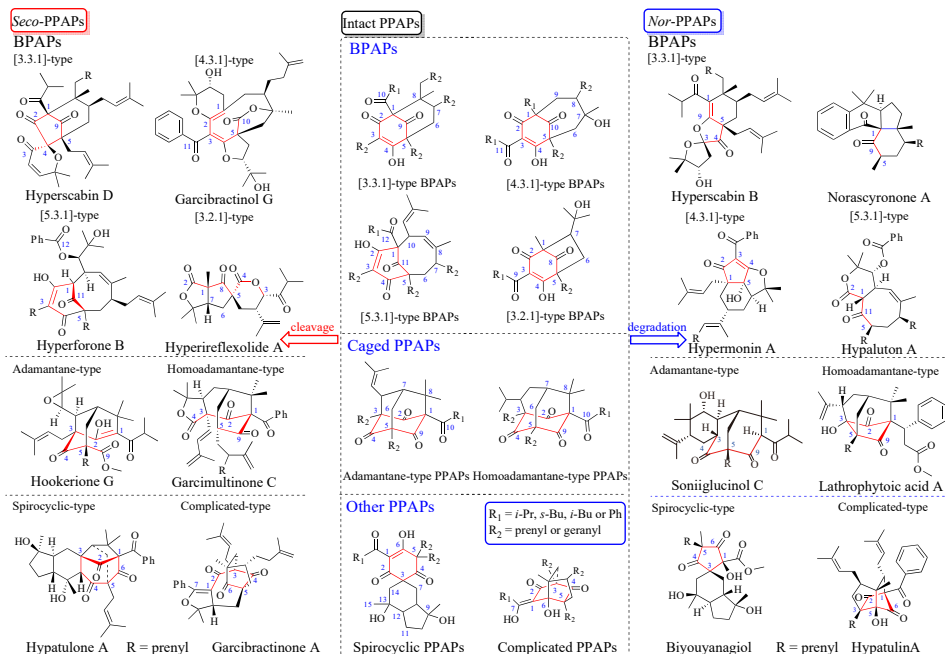


Fig. 1 Classification of PPAPs.

ness, only selected representative structures are highlighted within the main text, while detailed information on all identified structures, their sources, and associated bioactivities is provided in the Supplementary information. This review is intended to serve as a comprehensive reference and to facilitate further research and development efforts focused on these structurally and biologically intriguing PPAP subclasses.

2. Occurrence, classifications, and structures

As of June 2023, a total of 88 *seco*-PPAPs and 112 *nor*-PPAPs, exhibiting diverse skeletal frameworks, have been isolated from 30 plant species belonging to four genera (*Garcinia*, *Hypericum*, *Calophyllum*, and *Kielmeyera*) and two plant families (Clusiaceae and Calophyllaceae) as of June 2023. The majority of these compounds were isolated from the family Clusiaceae, with the exception of lathrophytoic acid A (**182**), which was obtained from the family Calophyllaceae²⁹. The genus *Hypericum* was the predominant source, accounting for approximately 90% of the compounds. The aerial parts of plants serve as rich sources of these PPAPs. Notably, 115 of these compounds were discovered within the past four years, reflecting the rapid advancement of phytochemical research. The distribution of these compounds across different plant genera and parts, along with their reported years, is illustrated in Fig. 2 and detailed in Tables S1–S7.

Seco- and *nor*-PPAPs are biosynthetically derived from normal PPAPs (Fig. 1). Their extensive structural variations stem from varying degrees of cleavage or degradation, rearrange-

ments, and cyclization, which have significantly contributed to a diverse and complex array of novel skeletons. These compounds can be categorized into six groups based on the skeletons of normal PPAPs (Fig. 1): *seco*- and *nor*-BPAPs, *seco*- and *nor*-caged PPAPs with adamantane and homoadamantane cores, and *seco*- and *nor*-other (spirocyclic and complicated types) PPAPs. *Seco*- and *nor*-BPAPs and *seco*-spirocyclic PPAPs represent the most common structural types, comprising 24%, 45%, and 14% of the total, respectively (Fig. 3).

Moreover, the proposed biosynthetic pathways suggest that the highly oxygenated acylphloroglucinol-derived cores of typical PPAPs are susceptible to oxidative cleavage or degradation. Additionally, isoprenyl or geranyl side chains may undergo intramolecular cyclization, forming new oxygen or carbon rings. In BPAPs, for instance, bonds such as C-1/C-2, C-1/C-9, C-1/C-10, C-1/C-12, C-2/C-3, C-3/C-4, C-3/C-10, C-4/C-5, or C-5/C-8 in acylphloroglucinols are prone to cleavage and cyclization with side chains, resulting in diverse structures. These can be categorized based on the location of ring splitting and carbon loss, such as 1,2-*seco*, 1,9-*seco*, and 2,3-*seco*-BPAPs. Typically, the C-1/C-2 and C-1/C-9 bonds of the acylphloroglucinol core are common cleavage sites. Further degradation, rearrangement, and cyclization can lead to other novel skeletons, including 2-*nor*, 2,3-*nor*, 2,3,4-*nor*, or 9-*nor* PPAPs. Considering the varying degrees and locations of cleavage, as well as ring formation, it is anticipated that more novel *seco*- and *nor*-PPAPs will be discovered in the future. The hypothetical biogenetic pathways of representative *seco*- and *nor*-PPAPs are summarized (Fig. 4).

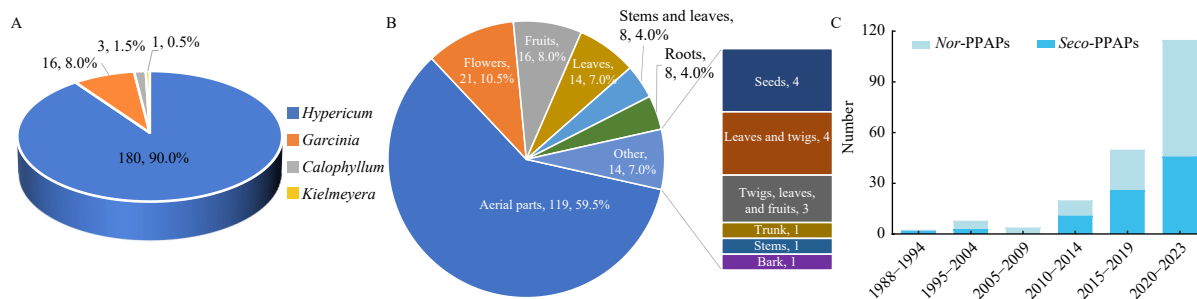


Fig. 2 Distributions of *seco*- and *nor*-PPAPs (A) in genera (B) and plant parts. (C) Number of *seco*- and *nor*-PPAPs reported in the literature up to June 2023.

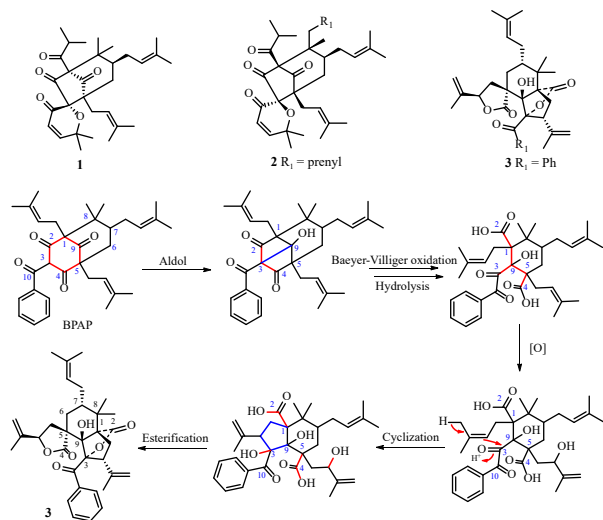


Fig. 5 Structures of 1–3 and possible biosynthetic pathway of 3.

ones A (**24**) and F (**25**)^{33,43}, hyperacrosin M (**26**)⁴⁴, hyperforinol D (**27**)⁴¹, and hyperacrosins C (**28**) and D (**29**) share an octahydrobenzofuran system skeleton⁴⁵, extracted from *H. sampsonii*, *H. acmosepalum*, or *H. forrestii*. Hyperbenzones A (**30**) and B (**31**), isolated from *H. beanii* roots, exhibit a 6/5/5 spirocyclic system and an unusual 3-oxabicyclo[3.1.0]hexan-2-one moiety, respectively. These structures are hypothesized to form via retro-Claisen reaction and cyclization (Fig. 7). The structure of **30** was conclusively determined through DP4+ calculation, crystallographic analysis, and ECD studies⁴⁶.

Thoreliolides A (**32**) and B (**33**) were isolated from the fruits

of *Calophyllum thorelii*, exhibiting a distinctive hexacyclic-fused six-membered lactone ring system⁴⁷. Hypsampsionone A (**34**), characterized by an octahydro-2*H*-chromen-2-one framework, was elucidated via comprehensive spectroscopic analyses and quantum chemistry calculations⁴⁸.

The 3,10-*seco*-BPAPs are proposed to arise from BPAPs via the Baeyer–Villiger reaction at the C-3 side chain, leading to structural cleavage. To date, four novel 3,10-*seco*-BPAPs (**37**–**40**) have been identified. Paucinone C (**37**), the first member of this subclass, possesses a cyclohexane-spiro-tetrahydrofuran moiety and was isolated from the leaves of *G. paucineris* in 2010⁴⁹. Oblongifolin M (**38**)⁵⁰, hyperascyrin N (**39**)⁵¹, and compound **40**, all exhibiting a tricyclo-[4.3.1.1⁴]-undecane skeleton⁵², were obtained from *G. oblongifolia*, *H. ascyron*, and *H. scabrum*, respectively.

Three novel compounds (**41**–**43**) derived from [3.2.1]-type BPAPs have been identified. Hyperireflexolides A (**41**) and B (**42**), a pair of epimers characterized by a 5/5/6 system with dilactone rings, were initially isolated from *H. reflexum*. The structure of **41** was confirmed through X-ray analysis, while **42** was subsequently revised and defined via a bioinspired total synthesis^{30,53}. Furthermore, George's group reported that **41** and **42** were derived from the significant precursor enaimeone A⁵⁴, a bicyclo[3.2.1]octane PPAP natural product, through the cleavages of C-2/C-3 and C-3/C-4 bonds, and rearrangement based on their biosynthetic pathway (Fig. 8)⁷. This work represents the first discovery of *seco*-[3.2.1]-type BPAPs. Compound **43**, isolated from *H. scabrum*⁵², possesses a cycloheptane ring fused to furan and γ -lactone rings and was generated from BPAP through the cleavage of the C-5/C-8 bond (Fig. 8). The structure of **43** was successfully elucidated by X-ray analysis³⁰.

Garcibractinols G (**44**) and H (**45**), two uncommon *seco*-

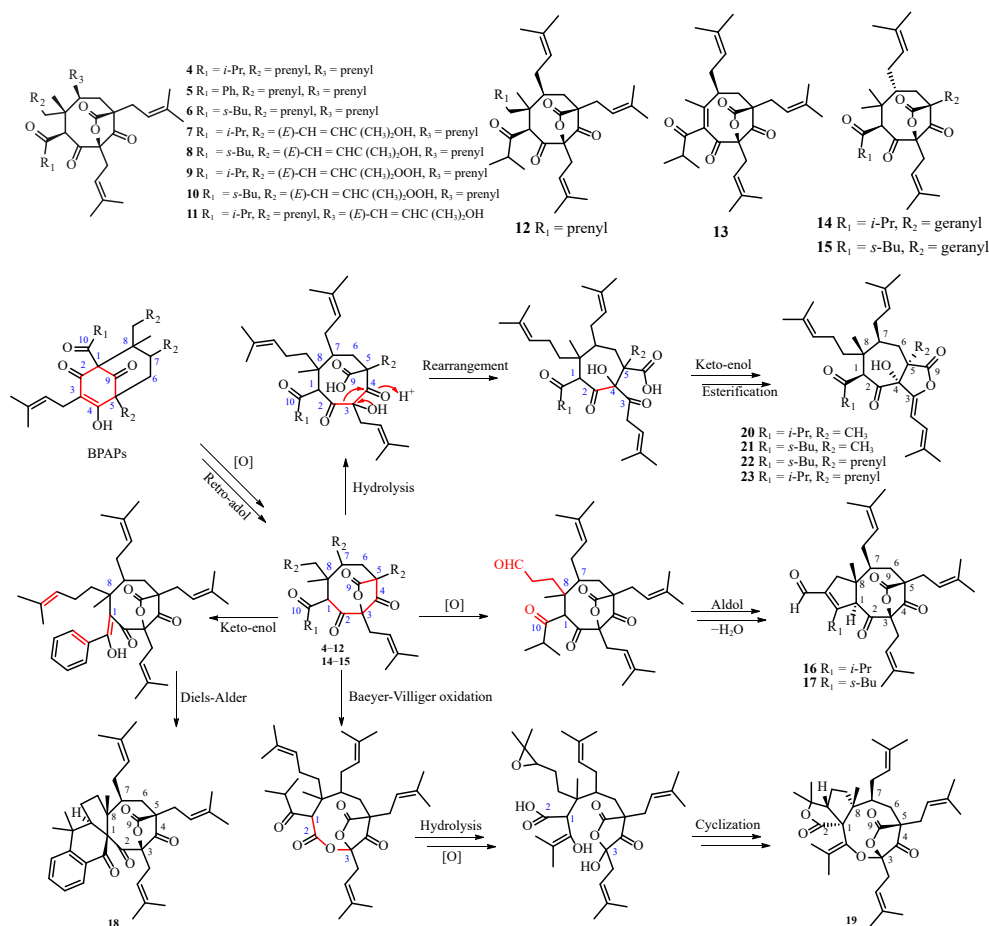


Fig. 6 Structures of 4–15 and possible biosynthetic pathway of 4–12, 14–23.

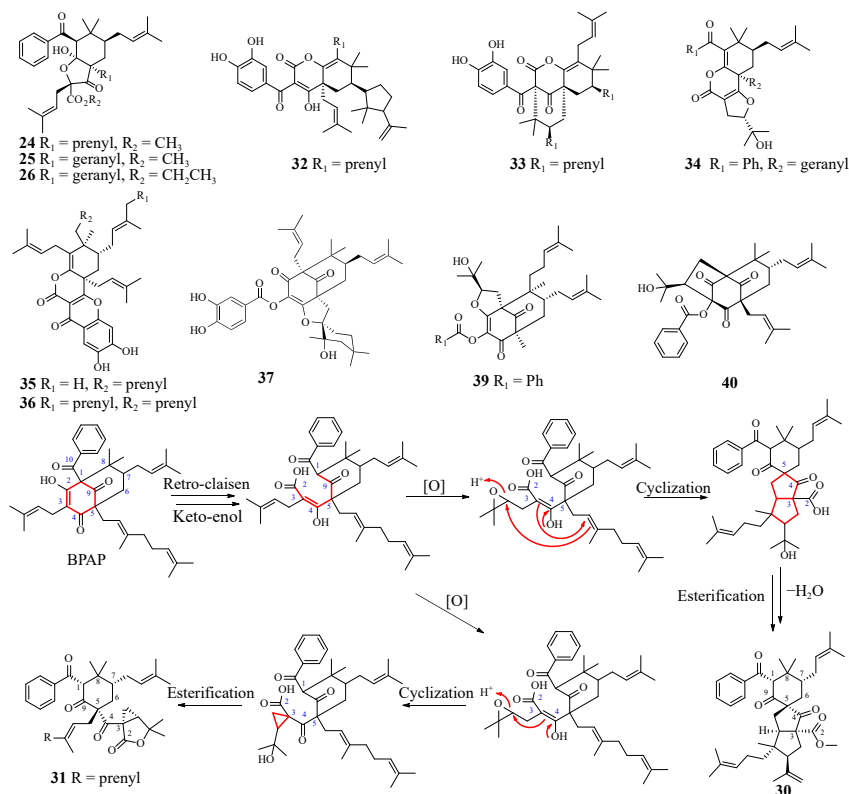


Fig. 7 Structures of 24–26, 32–37, 39–40 and possible biosynthetic pathways of 30–31.

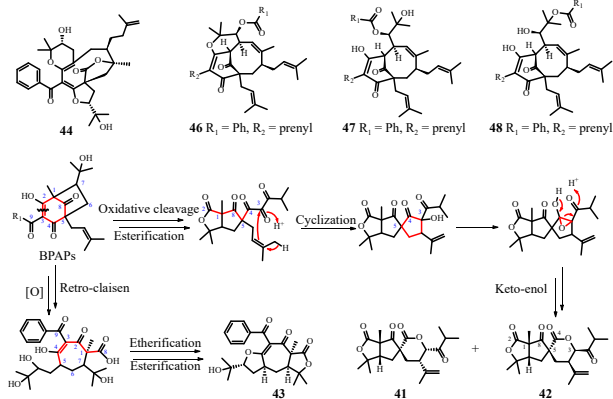


Fig. 8 Structures of 44, 46–48 and possible biosynthetic pathways of 41–43.

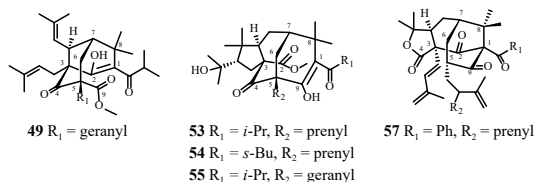


Fig. 9 Structures of 49, 53–55, and 57.

[4.3.1]-type BPAPs, were isolated from the fruits of *G. bracteata*. These compounds result from the cleavage of the C-1/C-10 bond, which opens the acylphloroglucinol ring through a retro-Claisen reaction (Fig. 4)⁵⁵.

Hyperforones A–C (46–48), three novel compounds obtained from *H. forrestii*, represent the first examples of benzoyl-migrated [5.3.1]-type BPAPs, classified as 1,12-*seco*-[5.3.1]-type BPAPs (Fig. 4)⁵⁶. From a biosynthetic perspective, 47 is hypothesized to be the precursor of 46 and 48⁵⁶.

2.3. *Seco*-caged PPAPs with adamantane and homoadamantane cores

Seco-caged PPAPs were hypothesized to originate from PPAPs with adamantane (tricyclo[3.3.1.1]decane) and homoadamantane (tricyclo[4.3.1.1]undecane) cores (Fig. 4). Four novel 1,9-*seco*-adamantane-type PPAPs, namely hypersubone A (49) (Fig. 9)⁵⁷, and hookeriones E–G (50–52), were isolated from *H. subsessile* or *H. hookerianum*⁵⁸. 49 represents the first identified member of 1,9-*seco*-adamantane-type PPAPs, featuring a tetracyclo-[6.3.1.1^{3,10}.0^{4,8}]-tridecane core. Additionally, biomimetic synthesis was employed to determine the absolute configurations of 50–52⁵⁸.

Four 1,2-*seco*-homoadamantane-type PPAPs, hypersonins A–D (53–56) (Fig. 9), were isolated from *H. wilsonii*. These compounds feature a novel bicyclo[4.3.1]decane-3-methoxycarbonyl structure⁵⁹. The absolute configuration of 53 was elucidated through X-ray crystallography⁵⁹. Additionally, garcimultinone C (57) (Fig. 9), the first reported 4,5-*seco*-homoadamantane-type PPAPs, exhibiting a 5-oxatricyclo[7.3.1.0^{3,7}]tridecane skeleton, was extracted from *G. multiflora*⁶⁰.

2.4. *Seco*-other PPAPs

Twenty-nine *seco*-spirocyclic PPAPs (58–86) and two *seco*-complicated PPAPs (87 and 88) are described. From a biogenetic perspective, *seco*-spirocyclic PPAPs likely originate from spirocyclic PPAPs with an octahydrospiro[cyclohexan-1,5'-indene]-2,4,6-trione core through the cleavage of C-1/C-2, C-1/C-6, C-3/C-4, C-4/C-5, C-12/C-13, or C-13/C-14 bonds, followed by cyclization and rearrangement to form new skeletons (Fig. 4). Complicated PPAPs, resulting from intramolecular [4 + 2] cycloaddition between the acylphloroglucinol core and side chains, potentially undergo oxidative cleavage and reconstruction to form new oxygen or carbon rings (Fig. 4).

Six 3,4-*seco*-spirocyclic PPAPs have been identified. Hypatunone A (58), featuring a novel skeleton, was isolated from *H. patu-*

lum⁶¹. Its structure was subsequently revised and confirmed through NMR analysis, biogenetic pathway examination, and quantum chemical calculations by Grossman's research team⁶². **58** represents the first 3,4-*seco*-spirocyclic PPAP, characterized by a unique 5/6/6/6/5 ring system⁶². The proposed biosynthetic pathway suggests that Paternò-Büchi reaction, nucleophilic attack, and aldol reactions could be key steps, as illustrated in Fig. 10. Hymoins C (**59**) and D (**60**), sharing the same skeletal structure, were isolated from *H. monogynum*⁶³. Hyperberlone A (**61**), isolated from *H. beanii*, features a rare tricyclo[4.3.1.0^{3,8}]decane core. It possessed an interesting caged tricyclic 6/6/5 ring system similar to **58–60** (Fig. 10)⁶⁴. Spihyperglucinols C (**62**) and D (**63**) contain an unexpected bicyclo[3.2.2]nonane core, with a proposed biosynthetic pathway depicted in Fig. 10⁶⁵.

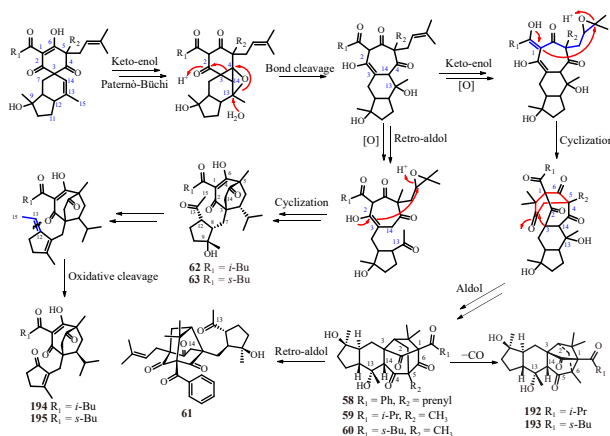


Fig. 10 Possible biosynthetic pathways of **58–63** and **192–195**.

To date, eleven novel 1,2-*seco*-spirocyclic PPAPs (**64–76**) with four distinct skeletons have been reported, and their proposed biosynthetic pathways are illustrated in Fig. 11. Compounds **64–68** exhibit a characteristic five-membered dilactone motif, originating from the cleavage of the C-1/C-2 bond of typical spirocyclic PPAPs via retro-Claisen reaction, followed by cyclization, keto-enol tautomerization, esterification, and stereospecific photoisomerization reactions. Biyoulactones A–C (**64–66**), isolated from *H. chinense*, represent the first discovered 1,2-*seco*-spirocyclic PPAPs⁶⁶, while hypemoins A (**67**) and B (**68**) were isolated from *H. monogynum*⁶⁷. Hypermonones A–D (**69–72**), four new 1,2-*seco*-spirocyclic PPAPs featuring single δ-lactone and tricyclic γ-lactone moieties⁶⁸, and hybeanones A (**73**) and B (**74**), containing a cyclopentanone unit fused to a tricyclic γ-lactone unit via a ketone carbonyl, were isolated from *H. beanii*. Their proposed biosynthetic pathway involves key photoisomerization or Baeyer-Villiger oxidation steps (Fig. 11)⁶⁹. The structures of **64**, **67**, **69**, and **73** were confirmed through X-ray diffraction analysis. Furthermore, hunascynols A (**75**) and B (**76**) represent the first instances of direct acylphloroglucinol core-opening spirocyclic PPAPs without subsequent cyclization (Fig. 11)⁷⁰.

Five 4,5-*seco*-spirocyclic PPAPs (**77–80**) are examined (Fig. 4). Furanmonogones A (**77**) and B (**78**), featuring a 4,5-*seco*-3(2*H*)-furanone motif, were isolated from *H. monogynum*⁷¹. Subsequently, Yang's group reported the first total synthesis of their deacylation products, further validating their skeletal structure⁷². Hyperhenone M (**79**), obtained from *H. henryi*, exhibited the same skeletal structure as **77** and **78**⁷³. Hyperbeanone A (**80**), characterized by a benz[*f*]indene-1,9(4*H*)-dione ring system, was isolated from *H. beanii*⁷⁴. Its carbon skeleton is hypothesized to form through the cleavage of the C-4/C-5 bond and a Diels-Alder reaction based on the intact spirocyclic PPAP (Fig. 11).

Biyoulactones D (**81**) and E (**82**), two novel 1,6-*seco*-spirocyclic PPAPs, featuring a unique γ-lactone ring, were initially isolated from *H. chinense* in 2012⁷⁵. Hyperilongenols A–C (**83–85**),

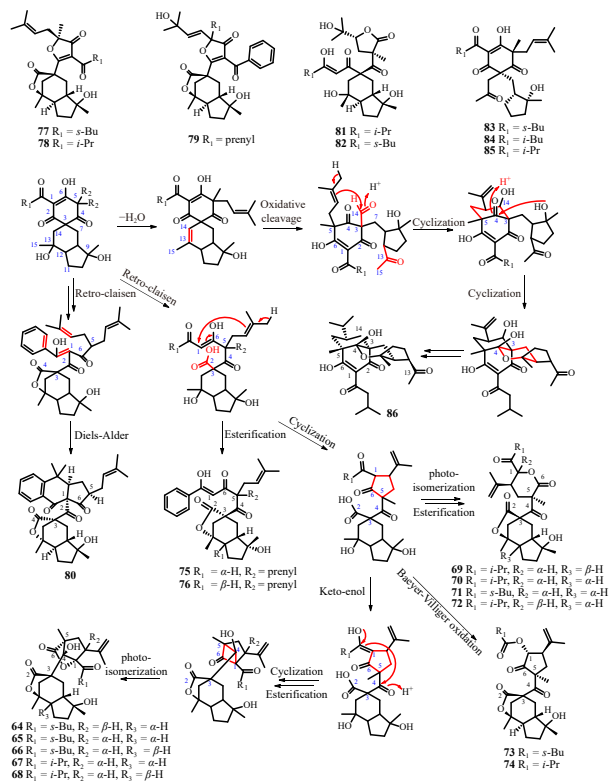


Fig. 11 Structures of **77–79**, **81–85** and possible biosynthetic pathways of **64–76**, **80**, **86**.

the first naturally occurring 12,13-*seco*-spirocyclic PPAPs with an enolizable β,β'-tricarbonyl system, were extracted from *H. longistylum*. **83** existed as a mixture of inseparable configurational isomers of *iso*-butyryl in solid crystals, while a suitable crystal of **83** confirmed its structure⁷⁶. Further analysis of *H. longistylum* yielded three unusual 13,14-*seco*-spirocyclic PPAPs, including longisglucinol A (**86**)⁷⁷. The structure of **86**, possessing an unprecedented 6/6/6/5 core as the first representative of this subgroup, was verified by X-ray analysis. Its potential biosynthetic pathway, involving oxidative cleavage and a series of cyclization reactions, is illustrated in Fig. 11.

Garcibractinones A (**87**) and B (**88**), two rare *seco*-complicated PPAPs, were isolated from *G. bracteata*, exhibiting a caged tricyclo-[4.4.1.1^{1,4}]dodecane skeleton. These compounds likely originate from nemorosonol or doitunggarcinone B via C-1/C-6 bond cleavage, followed by intramolecular cyclization of the side chains at C-1 and C-5 (Fig. 12)^{14, 78–81}. The structure of **87** was determined through comprehensive spectroscopic analysis and single-crystal X-ray diffraction⁷⁸.

2.5. Nor-PPAPs

Nor-PPAPs are derived through various degrees of carbon loss occurring in the side chain or *seco*-rings^{1, 23, 25}. Since the ini-

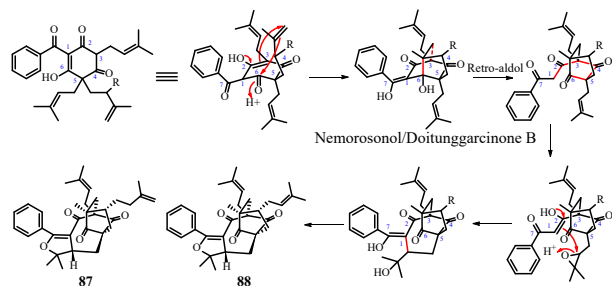


Fig. 12 Possible biosynthetic pathways of **87** and **88**.

tial discovery of soulatrone A (**117**) from the bark of *C. soulattri* in 1988⁸², 112 *nor*-PPAPs (**89–200**) have been identified. These compounds can be categorized into three groups: *nor*-BPAPs (**89–178**), *nor*-caged PPAPs with adamantane and homoadamantane cores (**179–186**), and *nor*-other PPAPs (**187–200**). *Nor*-BPAPs constitute the primary category of these compounds, exhibiting diverse structures and representing 45% of the summarized review (Fig. 3). Consequently, their proposed biosynthetic pathways have been the most extensively described in the literature.

2.6. *Nor*-BPAPs

Ninety *nor*-BPAPs, primarily derived from [3.3.1]-type BPAPs through degradation of C-2, C-2/C-3, C-2/C-3/C-4, or C-9 (including the side chain at C-3) (**89–174**), are described (Fig. 4). Compounds **175–177** originate from [4.3.1]-type BPAPs, while hypaluton A (**178**) stems from [5.3.1]-type BPAP.

Twenty-eight new 2-*nor*-BPAPs (**89–116**) exhibiting eleven distinct skeletal structures were identified through the absence of the C-2 carbonyl group in the acylphloroglucinol ring of conventional BPAPs (Fig. 4).

Norsampsones A–D (**89–92**) were the first identified 2-*nor*-BPAPs²⁶. Xu's group initially reported the total synthesis pathways of **89** and **90** to further confirm their structures⁸³. **94–104** share an identical skeleton with a unique hexacyclic-fused 1,6-dioxaspiro[4.4]nonane core and were isolated from *H. scabrum*, *H. patulum*, or *H. acmosepalum* in recent years. X-ray crystallographic data identified norhyperpalums B (**99**) and A (**104**)²⁴. Hyperacmosin N (**105**)⁸⁴, hyperpatone A (**106**)⁸⁵, and hyperacmosin R (**107**) exhibit a similar 6/5 skeleton, formed through key Baeyer-Villiger oxidative, decarbonylation, and intramolecular cyclization of the side chains at C-1 and C-8. Their oxygen ring systems were generated *via* a series of oxidation and intramolecular cyclization reactions (Fig. 13). Notably, **106** contains an intramolecular peroxy bridge with a rare 8/6/5/6/5 system, while **107** includes a 5,8-spiroketal subunit⁸⁶. Norgarmultinone B (**108**) and norwilsonnol B (**109**) feature a cyclohexane ring fused to a pyran or furan ring^{87,88}. Extensive phytochemical study of *H. perforatum* revealed three novel hyperforones F–H (**110–112**), possessing a rare macrocyclic lactone ring, presumed to form *via* the key oxidative cleavage of the C-3/C-4 bond

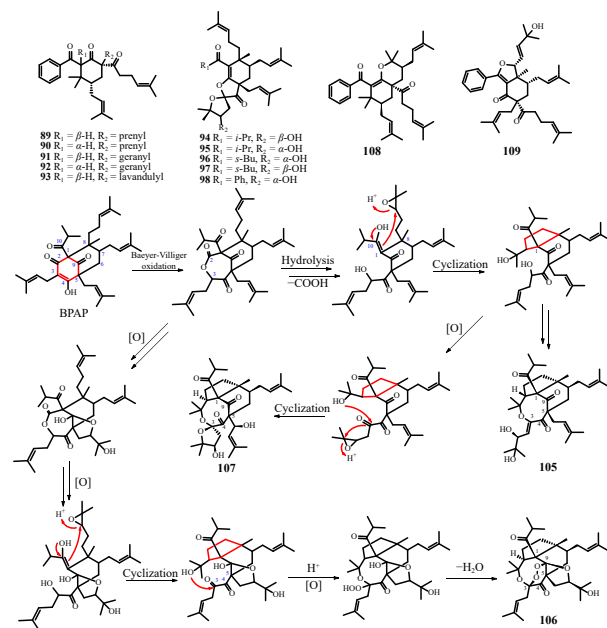


Fig. 13 Structures of **89–98**, **108**, **109** and possible biosynthetic pathways of **105–107**.

to open the ring (Fig. 14). X-ray diffraction analysis confirmed the absolute configuration of **110**⁸⁹. Investigation of *H. przewalskii* led to the isolation and characterization of norprzewalsone A (**113**), featuring a new 5/6/5/6/6 pentacyclic ring system⁹⁰, potentially originating from typical BPAP through C-2 degradation and secondary cyclization, including a Diels-Alder reaction between the side chains at C-1 and C-8 (Fig. 14).

Hypersampones A–C (**114–116**) represent the first *nor*-PPAPs exhibiting a tetracyclic 6/5/5/6 ring system. This unique skeletal structure is hypothesized to result from isopentenyl cyclase catalysis of intact BPAPs, followed by retro-Dieckmann reaction and decarboxylation (Fig. 14). Compounds **114** and **115** were presumably formed through C-1/C-10 bond cleavage and rearrangement from **116**. The structural elucidation of **114** was confirmed *via* X-ray crystallography⁹¹.

2,3-*Nor*-BPAPs may undergo loss of C-2, C-3, and the side chain at C-3, followed by cyclization and rearrangement to form diverse skeletons (Figs. 4 and 15). Thirteen novel 2,3-*nor*-BPAPs (**117–129**) have been reported. Soulatrone A (**117**), the first member of this subgroup isolated from *C. soulattri*, features a cyclohexane fused to γ -lactone, as determined *via* NMR data and X-ray crystallographic evidence⁸². Garcinielliptone J (**118**)³¹, and norhyperpalums F (**119**) and G (**120**) share the same skeleton as **117**²⁴. Norhyperpalum H (**121**)²⁴, and spirohypolactones A

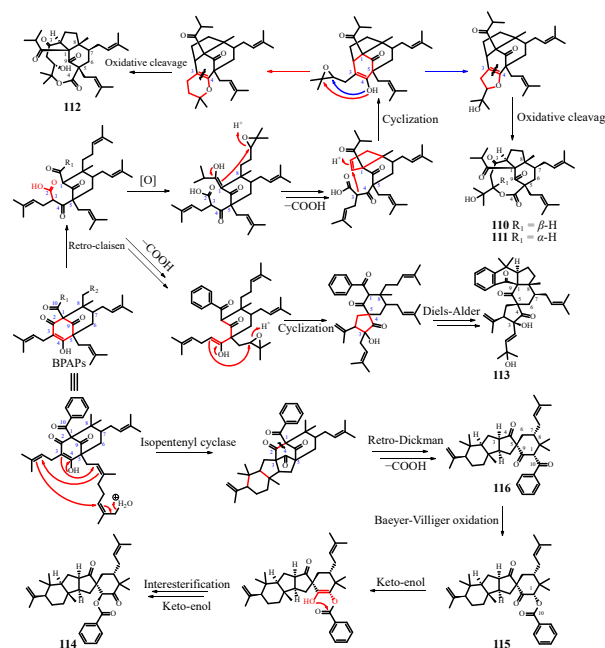


Fig. 14 Possible biosynthetic pathways of **110–116**.

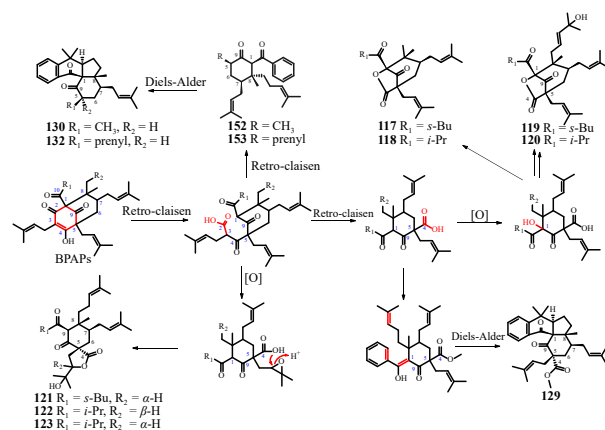


Fig. 15 Possible biosynthetic pathways of **117–123**, **129**, **130**, **132**, **152**, and **153**.

(122) and B (123)⁹², isolated from *H. patulum* or *H. perforatum*, possess the 2-oxaspiro[4.5]decane spirocyclic skeleton. Compounds 124–128 exhibit the same cyclohexane monocyclic system, and the absolute structure of garcinielliptone O (127) was established through total synthesis⁸³. A chemical investigation of *H. pseudohenryi* identified hypseudohenrin A (129) with a unique 6/5/6/6 ring system⁹³, which likely formed biogenetically via key retro-Claisen and Diels-Alder reactions (Fig. 15).

Forty-one 2,3,4-*nor*-BPAPs (130–170) exhibiting diverse skeletal structures have been identified. In comparison to 2,3-*nor*-BPAPs, the defining structural characteristic of this subgroup is the degradation at the C-4 position (Fig. 4).

Norascyrones A (130) and B (131) represent the first instances of 2,3,4-*nor*-BPAPs exhibiting the same carbon skeleton as 129. Their structures were confirmed through X-ray diffraction data and total or biomimetic synthesis^{23, 94-96}. Hypseudohenrin B (132) shares a similar skeleton to 130⁹³. Seven novel compounds (138–144) were isolated from *H. yezeense* or *H. monogynum*, comprising 6/5 carbon rings fused to a furan ring^{97, 98}. These compounds were formed through the degradation of C-2/C-3/C-4 and intramolecular cyclization (Fig. 16). The structures of yezo'otogirin A (138) and C (140), and hypermogin D (144) were elucidated via biomimetic synthesis^{95, 99-101}.

Since 151 was obtained from *H. perforatum* in 2001 as the first 2,3,4-*nor*-BPAP with a cyclohexane monocyclic ring^{102, 103}, eighteen analogous compounds (145–162) have been reported. The structures of hyperibrin B (146), and hyperscabrones H (148) and I (149) were revised through biosynthetic methods, NMR analysis, and chemical synthesis^{68, 104-106}. X-ray crystallography confirmed the structure of hypermonone H (147)⁶⁸. Hyperibrin A (145), garcinielliptone N (150), and hyperscabrin A (157) were successfully synthesized, validating their absolute structures^{83, 95, 107}. Phytochemical investigation of *H. perforatum* by Zhang's group yielded a series of 2,3,4-*nor*-BPAPs: hyperhexanones C–E (134–136)⁹², hyperforones A–E (163–167), I (168) and J (170) with nine novel skeletons⁸⁹. A plausible biosynthetic pathway for 163–170 was proposed, with key steps potentially involving retro-Claisen, intramolecular cyclization, oxidative cleavage, and aldol reactions (Fig. 16). Notably, hyperforones A–C (163–165 and 46–48) were reported with identical names but different structures. Norhyperpalum I (133)²⁴, norwilsonnol A (137)⁸⁸, and norprzewalsone B (169)⁹⁰ were obtained from *H. patulum*, *H. wilsonii*, or *H. przewalskii*, respectively, by the same research group. X-ray diffraction analysis confirmed the structures of 137, 163, 168, and 170.

The distinctive feature of 9-*nor*-BPAP (171–174) is the absence of C-9 in comparison to [3.3.1]-type BPAPs (Fig. 4). Hyphenrone F (171), the inaugural 9-*nor*-BPAP, contains an eight-membered carbon ring, potentially derived from 4 via decarbonylation (Fig. 17). Its structure was confirmed through X-ray diffraction analysis³⁴. Ascyrone C (172), isolated from *H. ascyron*, shares the same skeletal structure as 171⁴⁰. A bioassay-guided investigation of *H. perforatum* using UPLC-Q-Orbitrap-MS/MS led to the isolation of hyperforen A (173), the first PPAP exhibiting a bicyclo[7.3.0]dodecane core and the largest carbon ring¹⁰⁸. Uralin B (174), extracted from *H. uralum*, features a five-membered carbon ring. From a biogenetic perspective, 171, serving as a precursor, underwent a crucial Westphalen rearrangement for ring expansion, followed by Wagner-Meerwein rearrangement and cyclization between the two prenyl groups to form the 9/5 ring system. Conversely, 174 may have evolved from 171 through the loss of C-2/C-3/C-4 and the side chain at C-3, followed by oxidation and cyclization processes (Fig. 17)¹⁰⁹.

Hypermonins A (175) and B (176), a pair of isomers at C-5, exhibit a decahydroindeno[1,7-*bc*]furan ring system, derived from the [4.3.1]-type BPAPs through the loss of C-10 and subsequent cyclization (Fig. 18). These compounds represent the first examples of *nor*-[4.3.1]-type BPAPs¹¹⁰. Garcyunnanin G (177), isolated from *G. yunnanensis*, also belongs to this group¹¹¹. Hypaluton A (178), obtained from *H. patulum*, is the first known degradation product of [5.3.1]-type BPAP. Its biosynthesis occurs through the degradation of C-3 and C-4 based on the [5.3.1]-type BPAP structure (Fig. 18)²⁵.

2.7. *Nor*-caged PPAPs with adamantane and homoadamantane cores

These classes of *nor*-PPAPs originate from caged PPAPs with adamantane and homoadamantane cores via processes of decarbonylation and structural reconstruction (Fig. 4).

The eight reported compounds (179–186) are categorized into two distinct subgroups (Fig. 19): *nor*-adamantane-type PPAPs (179–181) and *nor*-homoadamantane-type PPAPs (182–186).

Hookerone H (179), isolated from *H. hookerianum*, represents the first characterized compound in this group exhibiting the adamantane-type PPAP skeleton through the loss of C-9 (Fig. 4)⁵⁸. Soniiglucinols C (180) and D (181) (Fig. 19), two 2-*nor*-adamantane-type PPAPs, were isolated from *H. wilsonii*¹¹².

Lathrophytoic acid A (182)²⁹, norsampsone E (183)¹¹³, hy-

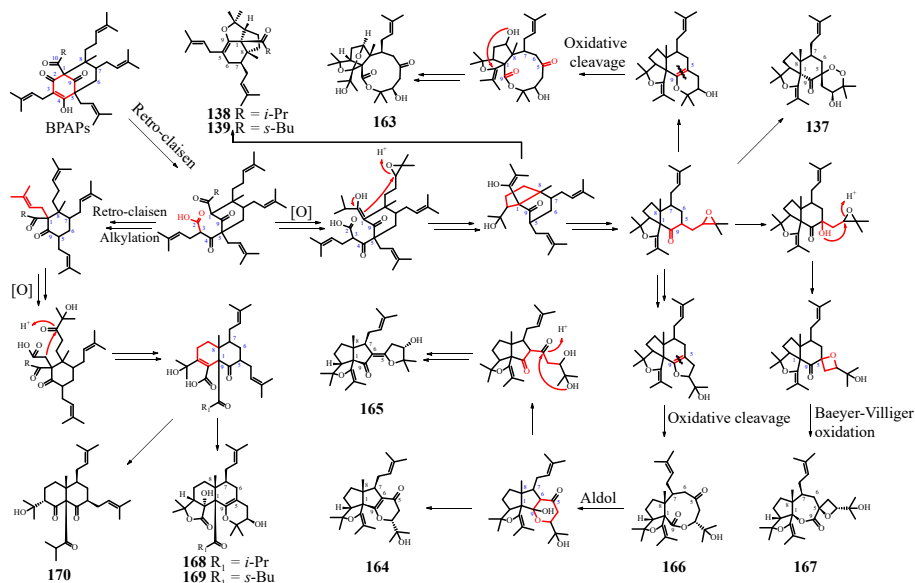


Fig. 16 Possible biosynthetic pathways of 137–139 and 163–170.

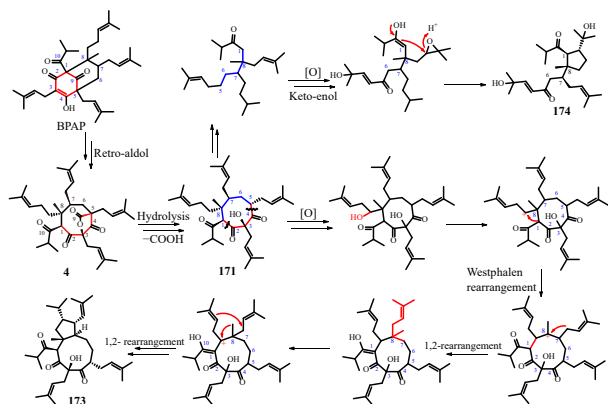


Fig. 17 Possible biosynthetic pathways of **4**, **171**, **173**, and **174**.

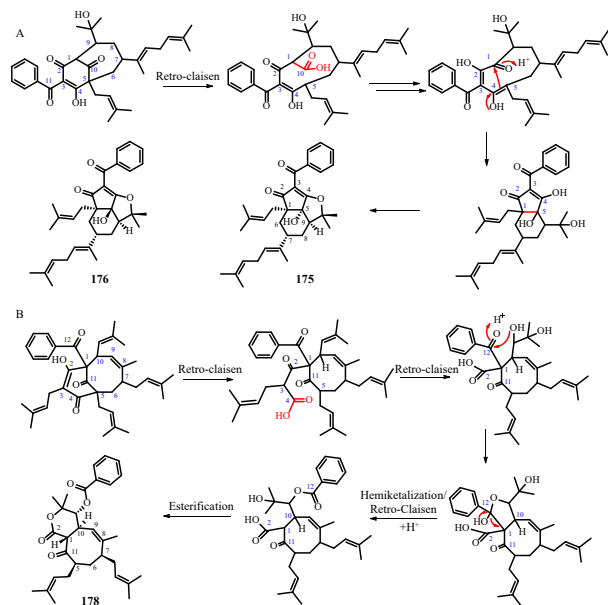


Fig. 18 Possible biosynthetic pathways of **175**, **176**, and **178**.

peracmosin E (**184**)¹¹⁴, and soniiglucinol A (**185**)¹¹², were formed *via* the loss of C-4 based on homoadamantane-type PPAPs. **182–184** exhibit an unusual caged carbon skeleton featuring a 1,3-dione-4-cyclopentanol moiety, while soniiglucinol A (**185**) presents an atypical 5/7/6 carbon skeleton. Furthermore, soniiglucinol B (**186**) represents the first identified 2-*nor*-homoadamantane-type PPAP¹¹². The structures of **180** and **185** were confirmed *via* single-crystal X-ray diffraction analysis.

2.8. *Nor*-other PPAPs

Nor-spirocyclic PPAPs (**187–195**) and *nor*-complicated

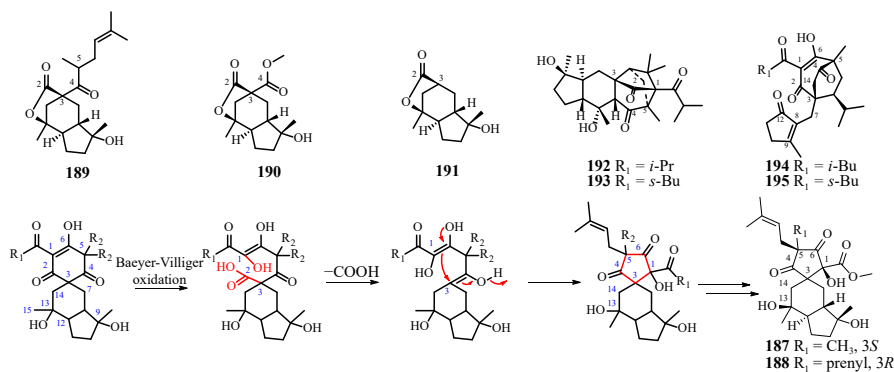


Fig. 20 Structures of **189–195**, and possible biosynthetic pathways of **187** and **188**.

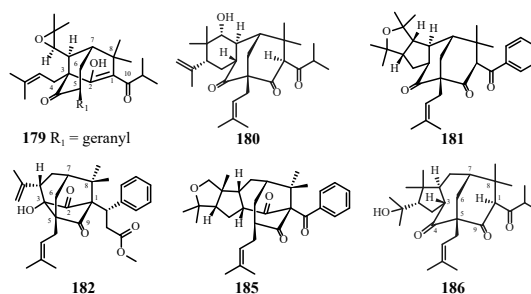


Fig. 19 Structures of **179–182**, **185**, and **186**.

PPAPs (**196–200**) are categorized according to their structural features and hypothesized biosynthetic routes (Fig. 1).

Biyouyanagiol (**187**), the first example of *nor*-spirocyclic PPAPs, likely originates from typical spirocyclic PPAP through the loss of C-2 *via* Baeyer-Villiger oxidation and decarboxylation, followed by cyclization, forming a unique cyclopenta-1,3-dione moiety (Fig. 20)¹¹⁵. Another novel 2-*nor*-spirocyclic PPAPs, uralin D (**188**)¹⁰⁹, was isolated from *H. uralum*.

Investigations into the chemical composition of *H. monogynum* flowers yielded three previously undescribed oxidative degradation spirocyclic PPAPs, hypemoins C–E (**189–191**)⁶⁷, as well as hymoins A (**192**) and B (**193**), which possess a rare caged pentacyclic 5/6/6/5/5 structure⁶³. Compounds **192** and **193** are hypothesized to be C-6 degradation products of **59** and **60**, respectively, potentially resulting from light-induced transformation in the proposed biosynthetic pathway (Fig. 10). The structural configuration of **192** was further confirmed through X-ray crystallography analysis.

Spihyperglucinols A (**194**) and B (**195**), two novel 13,15-*nor*-spirocyclic-type PPAPs isolated from *H. longistylum*⁶⁵, are likely derived from the loss of C-13 and C-15 in two 13,14-*seco*-spirocyclic PPAPs, **62** and **63** (Fig. 10). The absolute configuration of **194** was determined through single-crystal X-ray diffraction analysis⁶⁵.

Five novel *nor*-complicated PPAPs (**196–200**) were isolated from *H. patulum* or *H. forrestii*. Three 4-*nor*-complicated PPAPs, hypatulins A (**196**) and B (**197**)¹¹⁶, and hyperforinol C (**198**)⁴¹, were identified. Structurally, **196** and **198**, a pair of isomers, feature a unique tricyclic octahydro-1,5-methanopentalene core. Biogenetically, **196** may be formed through intramolecular cyclization, oxidation, ring cleavage of the C-3/C-4 bond, and cyclization associated with decarboxylation of a plausible biogenetic precursor, a monocyclic polyprenylated acylphloroglucinol (MPAP). Compound **197**, possessing a bicyclo[3.2.1]octane moiety, might be produced by oxidative cleavage of the C-1/C-6 bond based on precursor **196**, followed by methylation (Fig. 21). Additionally, Christmann's group reported a total synthesis of 3-*epi*-hypatulins B, highlighting the use of flow chemistry to enable challenging late-stage transformations in natural product synthesis¹¹⁷. An extensive chemical investigation of *H. patulum* yielded

two 6-*nor*-complicated PPAPs with an unprecedented tetracyclic system, hyperinoids A (**199**) and B (**200**)¹¹⁸. These compounds were proposed to biogenetically originate from an MPAP precursor through key intramolecular cyclization, oxidative cleavage of the C-5/C-6 bond, decarboxylation, and cyclization (Fig. 21). The structure of **199** was further confirmed by X-ray crystallographic data.

3. Biological activities

Seco- and *nor*-PPAPs, with their novel structural skeletons, have garnered significant attention due to their potent biological activities. These activities encompass anti-cancer, anti-inflammatory, hepatoprotective, anti-AD, MDR reversal, anti-depressant, neuroprotective, and immunosuppressive effects (Fig. 22). Among the reported compounds, twenty-nine *seco*- and *nor*-PPAPs demonstrated anti-cancer and MDR reversal activities. Additionally, 14 and 9 *seco*- and *nor*-PPAPs exhibited notable anti-AD and anti-depressant activities, respectively. Specifically, 10 *nor*-PPAPs showed potent anti-AD effects, while 6 displayed significant anti-depressant properties. This section provides a comprehensive summary and discussion of the diverse biological activities of *seco*- and *nor*-PPAPs (Tables S8–S16).

3.1. Anti-cancer activity

Approximately one-third of the compounds among 200 *seco*- and *nor*-PPAPs exhibited cytotoxic activity against various human cancer cell lines (Table S8). Several of these compounds demonstrated potent anti-cancer activity [half maximal inhibitory concentration (IC₅₀) < 10 μmol·L⁻¹], suggesting their potential as promising antitumor agents. Consequently, anti-cancer activity stands out as a notable characteristic of these PPAPs.

Hyperuralone E (**9**) and hyperfol A (**19**) exhibited activity against leukemia cell lines HEL and K562 with IC₅₀ values ran-

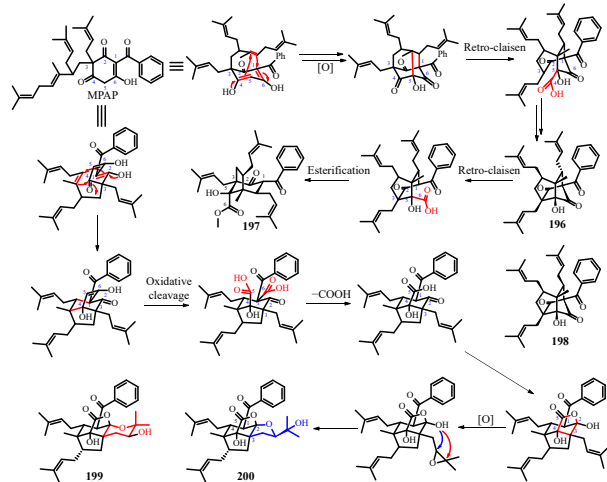


Fig. 21 Possible biosynthetic pathways of **196**–**200**.

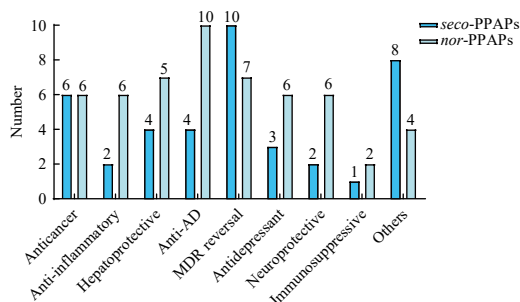


Fig. 22 Overview of potent biological activities of *seco*- and *nor*-PPAPs.

ging from 5.01 to 8.69 μmol·L⁻¹. These compounds induced apoptosis in HEL cells in a dose-dependent manner, with **9** increasing the expression of two marker proteins of apoptosis and cleavage of caspase-3 and PARP1³⁹. Thoreliolide B (**33**) demonstrated moderate cytotoxic activities against MCF-7, HeLa, and NCI-H460 tumor cell lines with IC₅₀ values of 9.7, 16.6, and 24.2 μmol·L⁻¹, respectively⁴⁷. Hunascynols A (**75**) (IC₅₀ 6.87 μmol·L⁻¹) and B (**76**) (IC₅₀ 9.86 μmol·L⁻¹) showed potent anti-proliferative effects against HCT116, surpassing the positive control 5-FU (IC₅₀ 22.12 μmol·L⁻¹). These compounds induced G₁ phase arrest and inhibited the expression of STAT3, Smad4, and Snail⁷⁰. Hyperbeanone A (**80**) exhibited remarkable cytotoxic activities by inducing G₁ phase cell cycle arrest and apoptosis in HL-60 and SU-DHL-4 cell lines with IC₅₀ values of 7.07 and 12.49 μmol·L⁻¹, respectively⁷⁴. Norhyperpalum B (**99**) showed good activity against hepatoma cell lines Hep3B, HepG2, SMMC-7721, and Huh-7 with IC₅₀ values ranging from 9.48 to 18.67 μmol·L⁻¹, while norhyperpalums C–E (**100**, **98**, and **97**) demonstrated weak activities. The structural comparison suggested that the benzene group and stereochemistry at the 1,6-dioxaspiro[4.4]nonane moiety might play crucial roles in cytotoxicity²⁴. Norascyronone A (**130**) exhibited cytotoxicity against SK-BR-3 (IC₅₀ 4.3 μmol·L⁻¹) and PANC-1 (IC₅₀ 8.4 μmol·L⁻¹) cell lines, while norascyronone B (**131**) showed a selective cytotoxic effect against SK-BR-3 (IC₅₀ 7.8 μmol·L⁻¹) cell lines²³. Additionally, hyperhexanone D (**135**) demonstrated mild anti-proliferative effects against HL-60 and NB4 cells with IC₅₀ values ranging from 8.1 to 11.7 μmol·L⁻¹⁹². Yezo'otogirin C (**140**) displayed potential cytotoxic effects against MGc80-3 with an IC₅₀ value of 9.54 μmol·L⁻¹¹⁰¹, and spihyperglucinol A (**194**) exhibited satisfactory cytotoxicity against HL60 with an IC₅₀ value of 9.12 μmol·L⁻¹⁶⁵.

3.2. Anti-inflammatory activity

The majority of *Hypericum* species are utilized in traditional medicine for treating inflammatory conditions, including wounds, bruises, hepatitis, and acute mastitis, suggesting that their secondary metabolites possess significant anti-inflammatory properties^{3, 119, 120}. A diverse range of structurally distinct *seco*- and *nor*-PPAPs have demonstrated anti-inflammatory activities by inhibiting the production of anti-inflammatory factors or inflammatory-related enzymes (Table S9).

Hypatulone A (**58**) demonstrated moderate NO inhibitory activity in an LPS-stimulated RAW264.7 cell model with an IC₅₀ value of 18.8 ± 1.75 μmol·L⁻¹, surpassing *N*-monomethyl-L-arginine (IC₅₀ 39.2 ± 0.36 μmol·L⁻¹, a positive control)^{61, 62}. Longisglucinol A (**86**) and spihyperglucinols A (**194**) and B (**195**) inhibited LPS-induced NO production with IC₅₀ values of 9.46 ± 1.21, 8.70 ± 1.18, and 9.23 ± 1.26 μmol·L⁻¹, respectively^{65, 77}. **86** suppressed proinflammatory factors nuclear factor κB (NF-κB), interleukin-6 (IL-6), IL-1β, TGF-β1, and enhanced anti-inflammatory factors ARG1 and cluster of differentiation 206 (CD206) expression at the messenger ribonucleic acid (mRNA) level, suggesting its potential as an anti-inflammatory agent⁷⁷. Soniiglucinols A (**185**) and B (**186**) inhibited NO production in RAW264.7 cells with IC₅₀ values of 4.23 and 7.58 μmol·L⁻¹, respectively, and COX-2 activities with IC₅₀ values of 5.27 and 8.32 μmol·L⁻¹, respectively, a characteristic of chronic inflammation¹¹². **185** decreased the production of various inflammatory factors in a dose-dependent manner, including PGE2, tumor necrosis factor α (TNF-α), and IL-1β, and significantly inhibited the LPS-induced nuclear translocation of NF-κB p65, supporting its anti-inflammatory and enzyme inhibitory properties¹¹². Additionally, NF-κB pathway luciferase assays revealed that hyperinoids A (**199**) and B (**200**) exhibited significant inhibitory activities in HEK293/NF-κB cells stimulated with TNF-α with IC₅₀ of 0.75 and 1.19 μmol·L⁻¹, respectively, compared to positive control bortezomib (IC₅₀ 0.07 μmol·L⁻¹). Concurrently, **199** and **200** suppressed the mRNA levels of certain proinflammatory genes, including IL-1β, IL-6,

and iNOS in RAW246.7 macrophages and primary mouse BMDM cells, indicating significant anti-inflammatory potential¹¹⁸.

3.3. Hepatoprotective activity

Seco- and *nor*-PPAPs demonstrated significant hepatoprotective properties (Table S10). Hyperascyrin N (**39**)⁵¹, hyperacmosin R (**107**)⁸⁶, hyperibrin G (**126**)¹²¹, hyperscabrone I (**149**)⁶⁸, and compound **161**¹²¹ exhibited hepatoprotective activities at 10 $\mu\text{mol}\cdot\text{L}^{-1}$ against paracetamol-induced HepG2 cell damage, with bicyclol or glutathione serving as positive controls. Hyphenrone A (**4**) and hyperscabin K (**159**) demonstrated hepatoprotective effects at 10 $\mu\text{mol}\cdot\text{L}^{-1}$ against DGalN-induced WB-F344 cell damage, with cell viabilities of 73% and 77%, respectively³². Hyperacmosins C (**28**) and D (**29**), as well as hyperacmosin E (**184**), exhibited potent hepatoprotective activity against paracetamol-induced HepG2 cell damage, showing cell viabilities of 56.31%, 55.10%, and 58.17%, respectively, compared to the hepatoprotective drug bicyclol as a positive control^{45, 114}. The structural comparison suggested that the *s*-butyl at C-10 might enhance hepatoprotective activity. Hypericumoxide N (**105**) displayed moderate hepatoprotective activity in the D-galactosamine-induced HL-7702 cell model⁸⁴.

3.4. Anti-AD activity

AD is a complex neurodegenerative disorder. Cholinesterase (ChE), including AChE and BChE, collectively regulate neurotransmitter levels, while protein phosphatase-2A (PP2A) and protein cleaving enzyme 1 (BACE1) influence β -amyloid ($A\beta$) production, representing significant biological targets in AD¹²²⁻¹²⁴. To date, nineteen *seco*- and *nor*-PPAPs have been evaluated for anti-AD activity (Table S11). Hyperuralones C (**7**) and D (**8**) demonstrated moderate AChE inhibitory activities with IC_{50} values of 9.6 and 7.1 $\mu\text{mol}\cdot\text{L}^{-1}$, respectively, compared to tacrine as a positive control (IC_{50} 0.33 $\mu\text{mol}\cdot\text{L}^{-1}$)³⁷. Two novel *seco*-PPAPs, hybeanones A (**73**) and B (**74**), exhibited potential AChE inhibitory potencies, with **74** showing superior binding affinity to **73** in molecular docking analysis⁶⁹. Hyperforones F-H (**110-112**), A-E (**163-167**), I (**168**) and J (**170**) enhanced PP2A activity with EC_{50} values ranging from 199.0–7761.0 $\text{nmol}\cdot\text{L}^{-1}$, and inhibited BACE1 activity with IC_{50} values ranging from 98.6–9553.0 $\text{nmol}\cdot\text{L}^{-1}$. Notably, **110** and **166** demonstrated remarkable PP2A activation with EC_{50} values of 199.0 and 258.8 $\text{nmol}\cdot\text{L}^{-1}$, respectively, and BACE1 inhibition with IC_{50} values of 98.6 and 136.2 $\text{nmol}\cdot\text{L}^{-1}$, respectively, surpassing the positive controls SCR1693 and LY2811376. Molecular docking analysis of **110** and **111** indicated the propan-2-ol group's significant role in PP2A and BACE1 interactions. **110** demonstrated superior efficacy in alleviating pathological and cognitive impairment in 3 × Tg AD mice compared to LY2811376, positioning it as a promising lead compound for AD treatment⁸⁹.

3.5. MDR reversal activity

Seco- and *nor*-PPAPs from the genus *Hypericum* demonstrate potential multidrug reversal activities against adriamycin (ADR) resistant cancer cell lines, such as HepG2/ADR and MCF-7/ADR (Table S12). Hyperhexanone F (**24**) was identified as a potential modulator of MDR in MCF-7/ADR cells, exhibiting a higher reversal fold (RF = 38) than the positive control (verapamil, RF = 20) at the same concentration of 10 $\mu\text{mol}\cdot\text{L}^{-1}$ ³³. Four *seco*-PPAPs, biyoulactones A (**64**) and B (**65**)^{66, 67}, and hypemoins A (**67**) and B (**68**)⁶⁷, exhibited moderate reversal activities against MCF-7/ADR cells at a concentration of 20 $\mu\text{mol}\cdot\text{L}^{-1}$, with RFs ranging from 11 to 27. Hypermonones A-D (**69-72**), I (**146**), and H (**147**) displayed MDR reversal activities against MCF-7/ADR cells with RFs ranging from 11 to 153 at 20 $\mu\text{mol}\cdot\text{L}^{-1}$. **146** demonstrated higher activity than the positive control (verapamil, RF = 53)⁶⁸.

Hypermogins B (**142**) and C (**143**), **146**, and **147** exhibited effective reversal activities for HepG2/ADR cells. Notably, **146** displayed a stronger reversal effect than verapamil (RF = 124)^{68, 98}.

3.6. Anti-depressant activity

H. perforatum has been traditionally utilized for treating mild to moderate depression^{21, 22, 125}. Consequently, numerous pharmacological studies have examined the anti-depressant effects of secondary metabolites from the genera *Garcinia* and *Hypericum*. Hyperscabins D (**2**), J-L (**158-160**), hyphenrone A (**4**), and hyperuralone C (**7**) were assessed *in vitro* for their anti-depressant effects via the inhibition of ($[^3\text{H}]\text{-5-HT}$) and noradrenaline ($[^3\text{H}]\text{-NE}$) reuptake in rat brain synaptosomes, utilizing duloxetine as a positive control. The results indicated that these compounds exhibited significant inhibitory activities at 1 $\mu\text{mol}\cdot\text{L}^{-1}$ against $[^3\text{H}]\text{-NE}$ ³². Hyperscabins A (**94**) and B (**95**) demonstrated effective reuptake inhibition of $[^3\text{H}]\text{-5-HT}$ ¹²⁶. Additionally, hypericumoxide N (**105**) displayed moderate activity against the $[^3\text{H}]\text{-5-HT}$ mode (Table S13)⁸⁴.

3.7. Neuroprotective activity

Recent studies have revealed that certain *seco*- and *nor*-PPAPs exhibit significant neuroprotective properties. Hyphenrone A (**4**), compounds **128** and **151**, hyphenrone F (**171**), and hyperforen A (**173**) demonstrated moderate neuroprotective effects in corticosterone-induced PC12 cells (Table S14)¹⁰⁸. Hypermonin A (**175**) showed superior protection of PC12 cells against corticosterone-induced injury compared to hypermonin B (**176**), suggesting that hydroxyl orientation influences neuroprotective activity¹¹⁰. Furthermore, hyperascyrin N (**39**) and hyperibrin A (**145**) exhibited protective effects on SK-N-SH cells against glutamate-induced damage^{51, 104}. Hyperscabin A (**94**) enhanced cell viability in oxygen-glucose deprivation-induced damage in SH-SY5Y cells at a concentration of 10 $\mu\text{mol}\cdot\text{L}^{-1}$. Notably, **94** demonstrated more potent activity than the positive control drug potassium-(1-hydroxypentyl)-benzoate (47.4%)¹²⁶.

3.8. Immunosuppressive activity

Several *seco*- and *nor*-PPAPs have demonstrated immunosuppressive properties (Table S15). Hyperonin A (**53**) and norwilsonnol A (**137**) inhibited the proliferation of murine splenocytes induced by anti-CD3/anti-CD28 monoclonal antibodies^{59, 88}. Additionally, **137** exhibited inhibitory activity in anti-CD3/anti-CD30 mAb-induced human peripheral blood mononuclear cells with an IC_{50} of $8.21 \pm 0.98 \mu\text{mol}\cdot\text{L}^{-1}$, suppressing the production of cytokines such as IFN- γ , TNF- α , IL-4, and IL-10 in both models. Hypaluton A (**178**) displayed moderate inhibitory activity with an IC_{50} of $6.86 \pm 0.72 \mu\text{mol}\cdot\text{L}^{-1}$ in LPS-induced murine B cells²⁵.

3.9. Others

In addition to the aforementioned activities, *seco*- and *nor*-PPAPs exhibited other notable properties (Table S16). Oblongifolin M (**38**) demonstrated more potent anti-EV71 activity with an IC_{50} of $16.1 \pm 0.5 \mu\text{mol}\cdot\text{L}^{-1}$, surpassing the positive control, ribavirin (IC_{50} $253.1 \pm 13.0 \mu\text{mol}\cdot\text{L}^{-1}$). Compound **40** displayed anti-protozoal activity, inhibiting the proliferation of *Trypanosoma brucei rhodesiense* and *Plasmodium falciparum*, with IC_{50} values of 3.07 ± 0.77 and $2.25 \pm 0.10 \mu\text{mol}\cdot\text{L}^{-1}$, respectively. Hyperilongenol B (**84**) and hypatulins A (**196**) exhibited significant anti-bacterial activity against *Staphylococcus aureus* and *Bacillus subtilis*, respectively^{76, 116}.

The anti-coagulant activity of hymoin C (**59**) demonstrated its ability to inhibit rabbit platelet aggregation induced by the platelet-activating factor, with an IC_{50} value of $119.23 \pm 11.85 \mu\text{mol}\cdot\text{L}^{-1}$ ⁶³. Hyperbenzone A (**30**) exhibited dose-dependent anti-

lipid accumulation effects in a palmitic acid-induced HepG2 cell model⁴⁶, while hypersampone A (**114**) reduced lipid accumulation in an oleic acid-treated HepG2 cell model by suppressing FAS and ACACA protein expression⁹¹. Garcibractinols G (**44**) and H (**45**) displayed antihyperglycemic activity by enhancing glucose consumption in HepG2 cells, with **44** showing greater efficacy than the positive control, metformin⁵⁵. Hypsampsone A (**34**) exhibited moderate α -glucosidase inhibitory activity⁴⁸. Norsampone C (**91**) and norsampone E (**183**) demonstrated potent, dose-dependent RXRa transcriptional inhibitory activities^{26,113}.

4. Conclusions

This review comprehensively examines the occurrence, structural characteristics, potential biosynthetic pathways, and biological activities of naturally occurring *seco*- and *nor*-PPAPs. From January 1988 to June 2023, 88 *seco*-PPAPs and 112 *nor*-PPAPs with diverse skeletons have been documented, categorized into six groups. As distinctive subsets of PPAPs, *seco*- and *nor*-PPAPs exhibit novel oxygen or carbon rings, serving as crucial sources of structural diversity. Their significant structural variations stem from the degrees and locations of cleavage, ring formation, and substituent variety. Additionally, the presence of various double bonds or conjugated fragments enhances cyclization and rearrangement possibilities. Future discoveries of more *seco*- and *nor*-PPAPs are anticipated. These compounds' rearranged polycyclic systems typically feature multiple chiral centers and quaternary carbons, while polyisopentenyl groups increase molecular flexibility, complicating single crystal formation and thus structural determination, particularly stereochemistry. Given the structural revisions of some novel non-crystalline compounds, the proposed biosynthetic pathways of *seco*- and *nor*-PPAPs with new skeletons offer a valid approach to understanding and elucidating their structural features. This approach can be complemented by quantum chemical calculations such as DP4+, J-DP4+, and ANN to further confirm complex structures with multiple chiral centers, especially for non-crystalline compounds^{62,127,128}. Chemical and biomimetic synthesis also provide effective methods for determining and validating structures assigned via NMR data^{105,129}.

As previously mentioned, *seco*- and *nor*-PPAPs have garnered significant attention due to their diverse structures and intriguing biological activities. Research has primarily focused on their anti-cancer, anti-inflammatory, hepatoprotective, anti-AD, MDR reversal, anti-depressant, neuroprotective, and immunosuppressive properties. However, most studies have been limited to activity screening, with few exploring mechanisms and structure-activity relationships (SARs) analysis, largely due to the scarcity of available compounds. Consequently, the synthesis or modification of active *seco*- and *nor*-PPAPs could be crucial for a more comprehensive investigation of their bioactivities, presenting a valuable direction for future research. Concurrently, the majority of *seco*- and *nor*-PPAPs are derived from the Clusiaceae family, particularly the genus *Hypericum*, yet current studies have examined only a fraction of these plants. It has been reported that merely 10% of the 500 *Hypericum* species have been investigated³. Thus, the vast number of unexplored *Hypericum* species may serve as a promising resource pool for chemically diverse *seco*- and *nor*-PPAPs.

This review aims to offer a comprehensive overview of the structures, biosynthesis pathways, and bioactivities of *seco*- and *nor*-PPAPs. It will serve as a structural reference for future lead compounds and provide a theoretical foundation for exploring the medicinal potential of ethnomedicines in subsequent research and development efforts.

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Supporting information

Additional information supporting this research can be obtained by contacting the corresponding authors via E-mail.

Declaration of competing interest

These authors have no conflict of interest to declare.

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