Epipolythiodioxopiperazines from Fungi: Chemistry and Bioactivities

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Abstract: Epipolythiodioxopiperazines (ETPs), characterized by a unique bridged disulfide or polysulfide dioxopiperazine six-membered ring, occur in many fungi. Due to its broad spectra of bioactivities, ETPs have drawn wide attention in recent years. This review covers diverse natural sources that produce ETPs, the synthetic chemistry of ETPs and an overview of promising bioactivities exhibited by some well studied ETPs. The plausible biosynthetic hypotheses of ETPs and some new results on antitumor activity of ETPs are also reviewed.

Keywords: Epipolythiodioxopiperazines, fungus, bioactivities, biosynthesis, antitumor.

1. INTRODUCTION

Epipolythiodioxopiperazines (ETPs), a group of important secondary metabolites from fungi, characterized by the bridged disulfide or polysulfide dioxopiperazine sixmembered ring, have been found to have wide range of bioactivities, such as antitumor, antimicrobial, antiviral immunosuppressive and enzyme inhibitory activities. It has been well recognized that disulfide or polysulfide functionalities are the key moiety for their bioactivities, and removal/cleavage of the sulfur bridge or addition of the reducing agent, e.g. dithiothreitol, usually will lead to complete loss of bioactivities. Gliotoxin (1), the first reported ETP, was originally isolated from Gliocladium fimbriatum in 1932 [1]. The chemistry of 35 ETPs up to the year 1976 had been reviewed by Taylor and Leigh [2]. With the rapid development of new techniques for structural identification and bioactivity screening, ETPs have attracted more intensive attention, and an updated overview of recent progress on ETPs appears necessary. This review covers the source, chemistry and bioactivities of reported ETPs, and SAR studies of their analogues. All ETPs are classified into eight groups according to their structural characteristics. The biosynthetic hypotheses of ETPs and some new results on antitumor activities of ETPs are summarized as well.

2. EPIPOLYTHIOPIPERAZINES

2.1. Gliotoxins and Analogues

Gliotoxin (1), the first reported ETP, was originally isolated from *Gliocladium fimbriatum* in 1932 [1] and lately isolated from *Thermascus* [3], *Candida* [4], *Penicillum terlikowskii* and *Aspergillus fumigatus* [5]. The structure of 1 was proposed by Woodward *et al.* in 1958 without specification of configurational detail [6]. Its absolute configuration was later unambiguously confirmed by X-ray crystallographic analysis in 1966 [7] and its first total synthesis using a novel solvent-dependent Michael reaction as a key step was achieved by Fukuyama et al. [8]. Due to its antibiotic, antiviral, immunosuppressive, platelet aggregation inhibitory and antitumor activities [9-14], to date, 1 has been being one of the most studied ETPs in pharmacology. The mechanism of toxicity of 1 was reviewed by Waring and Gardiner, respectively [15, 16]. Gliotoxin-E (2). characterized by a same framework of 1 containing a trithio bridge between C-3 and C-10a, was isolated from P. terlikowskii, A. fumigatus and T. crustaceus. Compounds 1 and 2 both were found to have ED_{50} values of 20 ± 10 ng/ml in the macrophage adherence assay [17]. The absolute configuration of 2 was determined by comparison of its CD profile with that of 1 and by chemical convertion to 1 in warm absolute ethanol in about 20% yield. Gliotoxin-G (3), characterized by containing a tetrathiodioxopiperazine moiety, was isolated from A. fumigatus. Compound 3 exhibited immunosuppressive activity in the macrophage adherence assay, in contrast to its inactive dithiomethylether analogues [18]. The absolute configuration of **3** was determined by comparison of its ORD curve with that of 1 [18], and its partial synthesis was completed by Kirby et al. [19]. Dehydrogliotoxin (4), isolated from P. terlikowskii, was shown to inhibit the growth of Bacillus subtilis at similar concentrations of 1 [20] and its first total synthesis was achieved by Kishi et al. [21]. Gliontrin A (5), a unique ETP containing a nitro-substituted aromatic ring, was competitive obtained from interaction between Sphingomonas bacterial strain and A. fumgatus fungual strain, and its absolute configuration was assigned by comparison of its CD profile with that of 1, and was further confirmed by X-ray crystallographic examination. Compound 5 displayed promising antibacterial activity against three strains of MRSA with significant MIC values of 0.78 µg/ml. It also shows potent cytotoxic activity against HCT-116, A549, AGS, DU145, MCF-7 and HepG2 cancer cell lines with IC₅₀ values ranging from 0.24 to 2.3 μ M in *in vitro* MTT assay [22]. It is worth to note that **5** could not be detected in monoculture broths. This result demonstrated that microbial co-culture could produce novel biologically relevant molecules (compounds 1-5 are shown in Table 1).

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Table 1.Structures of 1-5



No.	Name	Name x Species					
1	Gliotoxin	2	G. fimbriatum, T. sp., C. sp., P. terlikowskii, A. fumigatus	[1, 3, 4, 5]			
2	Gliotoxin-E	3	P. terlikowskii, A. fumigaus, T. crustaceus	[17]			
3	Gliotoxin-G	4	A. fumigatus	[18]			
4	Dehydrogliotoxin		Dehydrogliotoxin P. terlikowskii				
5	Glionitrin A		Coculture of Sphingomonas bacteria and A. fumigatus				

2.2. Emestrins and Analogues

Emestrins A (6) [23] and B (7) [24] were originally isolated from *Emericella striata*. Compound 6, the first reported example of 15-membered macrocyclic ETP with strong antifungal activity, was formally derived from two molecules of phenylalanine and one molecule of benzoic acid. The structure of 6, including the absolute configuration, was determined based on a spectroscopic investigation of some derivatives and X-ray crystallography of emestrin methanol solvate. The absolute configuration of 7 was determined by comparison of its CD profile with that of 6. As determined by the cylinder-agar plate assay, the MIC of 6 against Gliierella zeae and P. expansum were 10 and 2.5 µg/ml, respectively [23]. Compound 7 showed effective inhibition against Tricophyton sp. at 0.78-1.56 µg/ml and Microsporum sp. at 3.12-6.25 µg/ml in the in vitro assay. Compound 7 also showed inhibitory activity against Bacillus subtilis and Escherichia coli at concentration of 25 µg/disc [24]. Emestrins C-E (8-10), Secoemetrin C_1 (17) and MPC1001B-H (11-15) were produced by fungi from Musk Ox dung and C. sp. KY4922, respectively [25, 26]. The absolute configuration of 8 was determined on the basis of NMR and CD profile by comparison with those of 7, and the conclusion was further supported by the similarity of optical rotation [26]. Compounds 6, 8-10 and 17 were evaluated for their activity in the CCR2 membrane binding assay using CCR2 receptor CHO cell membrane and ¹²⁵I-MCP [25]. Compounds 6, 8 and 9, each with two to three sulfur atoms, and with or without a free phenolic group, exhibited (IC₅₀) value $\sim 0.8 \mu$ M) equal activities in CCR2 membrane binding assay. Compound 10, with four-sulfur atoms, was slightly less active (IC₅₀ = 1.1 μ M) than 6, 8 and 9. Compound 17 exhibited IC₅₀ = 9.3 μ M and was ten times less active than 6. In a whole cell assay using human monocyte harvested from Leukopacks, 17 inhibited the ¹²⁵I-hMCP-1 binding to CCR2 receptor with IC₅₀ value of 7.9 μ M, which is about two fold lower than macrocyclic **6** (IC₅₀ = 5.4 μ M), **8** (IC₅₀ = 4.1 μ M) and 9 (IC₅₀ = 4.2μ M). These results indicate the importance of the macrocycle for inhibitory activity. Compounds **8**, **9**, **11** and **12**, all containing both macrocyclic skeletons and polysulfide bridge, showed antiproliferative activities against DU145 human prostate cancer cell line with IC₅₀ values of 9.3, 16, 39 and 12 nM, respectively. However, the decrease of the antiproliferative activities in **13** (IC₅₀ = 83 nM), **14** (IC₅₀ = 350 nM) and **15** (IC₅₀ = 83 nM) indicates that the macrocyclic ring and polysulfide structures are critical for the antitumor activity [26]. Secoemestrin C (**16**), an epitetrathiodioxopiperazine derivative isolated from an emestrin-producing fungus *E. foveolata*, may be the key intermediate from **6** and/or **7** to **14** and **15** [27] (compounds **6-17** are shown in Table **2**).

2.3. Sporidesmins

Sporidesmins A-J (18-26, shown in Table 3) are toxic metabolites isolated from *Pithomyces chartarum*, a fungus causing farm animals susceptible to a seasonal hepatogenous photosensitization disease known as facial eczema in New Zealand. Successful isolation and structure determination of **18-26** were carried out by Taylor and his coworkers [28-34]. Sporidesmin A (18), principal toxin of *Pithomyce* chartarum, was totally synthesized by Kishi et al. [35] and its absolute configuration was determined in the early 1960s [7]. Sporidesmin D (21) can be obtained by treating 18 with NaBH₄ and MeI. The CD profile for the **21** was identical to **18**, indicating no change in absolute configurations [30]. Sporidesmin E (22) can be converted to 21 by reduction and methylation, and to 18 when treated with PPh3. When 18 was sulfurated with P_4S_{10} and S, it was possible to isolate 22 from the reaction products [32]. When the tail factions from chromatogram of 22 were re-chromatographed, sporidesmin G (24) was obtained. Compound 24 can be converted to 18 on irradiation or treatment with PPh_3 , and to 21 after reduction and methylation. Similarly, 18 and 22, when treated with H_2S_2 , can be converted to 24 [33]. The dimethyl thioether analogues of 18, sporidesmins D (21) and F (23), are both inactive metabolites, which indicates that the polysulfide structure is vital for their activity and toxicity

Table 2. Structures of 6-17



H₃CO

СНО

15

H₃CO





No.	Name	R ₁	\mathbf{R}_2	R ₃	x	Species	Refs.
6	Emestrin	ОН	ОН	ОН	2	E. striata	[23]
7	Emestrin B	ОН	ОН	ОН	3	E. striata, E. foveolata	[24, 27]
8	Emestrin C (MPC1001)	ОН	ОН	OCH ₃	2	SM ^a , C. sp. KY4922	[25, 26]
9	Emestrin D (MPC1001D)	ОН	ОН	OCH ₃	3	SM, C. sp. KY4922	[25, 26]
10	Emestrin E	ОН	ОН	OCH ₃	4	SM	[25]
11	MPC1001B	Н	ОН	OCH ₃	2	C. sp. KY4922	[26]
12	MPC1001C	ОН	Н	OCH ₃	2	C. sp. KY4922	[26]
13	MPC1	001E				C. sp. KY4922	[25]
14	MPC1	001G				C. sp. KY4922	[26]
15	MPC1	001H				C. sp. KY4922	[26]
16	Secoeme	estrin C				E. foveolata	[27]
17	Secoeme	strin C1				SM ^a	[25]

^aSterile mycelia.

[30]. Compound **22** was an unstable, but highly cytotoxic metabolite among the above spirodesmins with an end-point of 0.1 ng/ml against HeLa cancer cells [31, 32]. Compound **24** inhibited the growth of HeLa cells at 2 ng/ml, which is being slightly less active than **18** (1 ng/ml) and considerably less toxic than **22**. The toxicity of **24** may be due, in part, to undetected trace of **22** [33].

2.4. Aranotin and Analogues

Aranotin (27), acetylaranotin (LL-S88, 28) [36] and apoaranotin (29) [37] were all produced by a fungus

designated as *Arachniotus aureus*. Compound **28** was also isolated from *A. terreus* [38]. The absolute configuration of the key metabolite, **28**, was confirmed by X-ray diffraction analysis [39]. Compound **28** was found to be a viral RNA synthesis inhibitor against strains of rhino-, Coxasckie, polio- and parainfluenza viruses, and to protect mice against lethal infections produced with Coxsackie A21 or influenza S/Md viruses. However, no antiviral activity was observed for LL-S88 β , which is a dithiomethyl ether derivative of **28**. The specific action of **28**, completely blocking viral RNA synthesis at levels which are without effect on cellular RNA synthesis, was believed to be the basis for its antiviral

Table 3. Structures of 18-26





No.	Name	R ₁	R ₂	R ₃	\mathbf{R}_4	R 5	x	Species	Refs.
18	Sporidesmin A	Cl	CH ₃ O	CH ₃ O	OH	OH	2	Pithomyces. chartarum	[28]
19	Sporidesmin B	Cl	CH ₃ O	CH ₃ O	ОН	Н	2	Pithomyces. chartarum	[28]
20	Sporidesmin C	Cl	CH ₃ O	CH ₃ O	OAc	OAc		Pithomyces. chartarum	[29]
21	Sporidesmin D	Cl	CH ₃ O	CH ₃ O	ОН	ОН	-	Pithomyces. chartarum	[30]
22	Sporidesmin E	Cl	CH ₃ O	CH ₃ O	ОН	ОН	3	Pithomyces. chartarum	[31, 32]
23	Sporidesmin F	Cl	CH ₃ O	CH ₃ O	ОН	ОН	-	Pithomyces. chartarum	[30]
24	Sporidesmin G	Cl	CH ₃ O	CH ₃ O	ОН	ОН	4	Pithomyces. chartarum	[33]
25	Sporidesmin H	Н	CH ₃ O	CH ₃ O	Cl	Н	2	Pithomyces. chartarum	[34]
26	Sporidesmin J	Cl	CH ₃ O	Н	ОН	ОН	2	Pithomyces. chartarum	[34]

activity [38]. It was also found that 28 inhibited influenza RNA polymerase in the absence of dithiothreitol [40]. Emethallicins A-F (30-35) were isolated from E. heterothallica and their absolute configurations were established by Kawi and his coworkers [41-44]. Compounds 30-32, 34 and 35 showed a potent inhibitory activity upon compound 48/80-induced histamine release from mast cell with IC₅₀ values of 0.03, 0.08, 1.0, 0.1 and 0.2 µM, respectively, and the IC₅₀ values for inhibition of 5lipoxygenase were determined as 1.7, 1.3 and 2.6 µM for 30-32, respectively. The inhibitory activity, toward histamine release, of the acetates of 30-32 was usually weaker than those of the original emethallicins, except for emethallicin D monoacetate whose activity was as strong as that of 30. SCH 64874, SCH 64875 and SCH 64877 (36-38), isolated from an unidentified fungus, were found to be potent EGF receptor antagonists, with IC₅₀ values of 1.0, 1.0, 1.25 μ g/ml, respectively [45] (compounds 27-38 are shown in Table 4).

2.5. Epicorzines and Analogues

Epicorazines A (**39**) and B (**40**) both exhibiting antibiotic activities were isolated from *Epicoccum nigrum* and *Stereum hirutum* [46-49], and epicorazine C (**41**) was isolated from *Stereum hirutum* [49]. The absolute stereochemistries of **39**

and 40 were determined by X-ray analysis and interpretation of their CD and NMR data [47, 48], while the absolute stereochemistry of 41 was derived from NMR data and CD comparison with those of 40 [49]. Moreover, the antibacterial activity assay showed that 39 and 41 were more sensitive to Gram-positive bacteria than Gram-negative bacteria and both compounds also displayed inhibitory effect to Candida albicans [49]. Ambewelamides A (42) and B (43) were isolated from the lichen Usnea sp. [50], representing rare members of ETP family characterized by containing ring A/E epoxides and a diketopiperazine epidisulfide bridge in their molecules. Their structures were elucidated via a combination of X-ray diffraction and spectroscopic analyses. Compound 42 exhibited potent in *vitro* cytotoxicity (murine leukemia P388: $IC_{50} = 8.6$ ng/ml) and significant in vivo antineoplastic activity (P388: %T/C 140 @ 160 µg/Kg). Rostratins A-D (44-47) were isolated and characterized from Exserohilum rostratum, and their absolute configurations were determined by the modified Mosher method [51]. Compounds 44-47 showed in vitro cytotoxicity against HCT-116 with IC₅₀ values of 9, 4.4, 1.6 and 35 µM, respectively. Epicoccins A-H (48-55), all possessing common unusual sulfur bridges, were obtained from cultures of a Cordyceps-colonizing fungus E. nigrum

Table 4.Structures of 27-38



No.	Name	R ₁	R ₂	x	Species	Refs.
27	Aranotin		Ac	2	Arachniotus aureus	[36]
28	Acetylaranotin (LL-S88)	Ac	Ac	2	Arachniotus aureus, A. terreus	[36, 38]
29	Apoaranotin	Н	Ac	2	Arachniotus	[37]
30	Emethallicin A	а	b	2	E. heterothallica	[41]
31	Emethallicin B	а	b	4	E. heterothallica	[42]
32	Emethallicin C		а	4	E. heterothallica	[42]
33	Emethallicin D	а	b	3	E. heterothallica	[43]
34	Emethallicin E	b	b	2	E. heterothallica	[44]
35	Emethallicin F	b	b	2	E. heterothallica	[44]
36	SCH 64874	с	с	2	NF ^a	[45]
37	SCH 64875		с	3	NF	[45]
38	SCH 64877	с	с	4	NF	[45]

^aUnidentified fungus.

[52, 53]. The structurally most related known compound 47 could be the biosynthetic precursor for 48-55 [52]. Epicoccin A (48) showed modest antimicrobial activity against Bacilus subtilis, affording a zone of inhibition of 12 mm at 100 μ g/disk, whereas **49-51** were inactive at the same level (ciproflozaxin: 28 mm zone of inhibition at 100 µg/disk) [52]. Compounds 52-55 were tested for in vitro anti-HIV-1 activity. However, only the dithiomethyl ethers, 54 and 55, showed inhibitory effects on HIV-1 replication in C8166 cells with EC₅₀ values of 13.5 and 42.2 μ M, respectively [53]. Gliovirin (56), isolated from G. virens, is possibly derived from two molecules of phenylalanine. Compound 56 showed antibiotic activity against Pyhtium ultimum and selective activity against members of the Oomycetes [54]. The N-methlygliovirin, FA-2097 (57), isolated and characterized from Eupenicillium abidjanum IFO 8939, exhibited highly antibiotic activity against anaerobic bacteria, especially Fusobacterium sp. and Bacteroides sp. [55, 56]. Pertrichodermamide A (58), isolated from Trichoderma sp. BCC 5926, exhibited inhibitory activity against Mycobacterium tuberculosis H₃₇Ra with a MIC value of 12.5 µg/ml, and its absolute configuration was established by X-ray crystallographic analysis [57] (compounds **39-58** are shown in Table **5**).

2.6. Chaetocins, Verticillins, and Analogues

The ETPs 59, 61-67 were isolated from *Chaetomium* sp. [58, 60-63], and their structures including absolute configuration were determined by chemical, spectroscopic or X-ray analyses, and 11,11'-dihydroxychaetocin (60) was originally isolated from the fungus Verticillium tenerum [59]. Chaetocin (63) was a compound with antibacterial and cytostatic activity [58], and its asymmetric total synthesis was achieved by Sodeoka et al. [64]. The bioassay against the growth of Hela cells showed that 59-63 were all cytotoxic with IC₅₀ values ranging from 0.02 to 0.04 μ g/ml, much stronger than three derivatives of 59 including monosulfide, dethiomethylthio, dethioacetylthio, and the similar SAR were observed in the antibacterial bioassay against S. aureus [60]. Compounds 64 and 65, processing ETP moieties, both displayed comparatively high immunosuppressive activity against Con A-induced (T-cells) and LPS-induced (B-cells) proliferations of mouse splenic

Table 5.Structures of 39-58



No.	Name	Species	Refs.
39	Epicorazine A	Epococcum Nigrum, Stereum hirutum	[46, 49]
40	Epicorazine B	Epococcum Nigrum, Stereum hirutum	[46, 49]
41	Epicorazine C	Stereum hirutum	[49]
42	Ambewelaminde A	Lichen usnea sp.	[50]
43	Ambewelaminde B	Lichen usnea sp.	[50]
44	Rostratin A	Exserohilum rostratum	[51]

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No.	Name	Species	Refs.
45	Rostratin B	Exserohilum rostratum	[51]
46	Rostratin C	Exserohilum rostratum	[51]
47	Rostratin D	Exserohilum rostratum	[51]
48	Epicoccin A	Epicoccum. Nigrum	[52]
49	Epicoccin B	Epicoccum. Nigrum	[52]
50	Epicoccin C	Epicoccum. Nigrum	[52]
51	Epicoccin D	Epicoccum. Nigrum	[53]
52	Epicoccin E	Epicoccum. Nigrum	[53]
53	Epicoccin F	Epicoccum. Nigrum	[53]
54	Epicoccin G	Epicoccum. Nigrum	[53]
55	Epicoccin H	Epicoccum. Nigrum	[53]
56	Gliovirin	G. virens	[54]
57	FA-2097	Eupenicillium abidyamum	[55]
58	Pertrichodermamide A	Trichoderma sp.	[57]

lymphocytes, while 66 and 67 processing dethiodimethylthiodioxopiperazine moieties displayed very weak activity in the above mentioned bioassay, indicating the immunosuppressive activity of 64 and 65 might attribute to their ETP moieties in their molecules [63]. This seems to be similar to the result already obtained on other fungal ETPs, sporidesmin A (18) and gliotoxin (1), by Mullbacher *et al.* [65]. Verticillins A-C (68-70), D-F (71-73) and G (74) were originally isolated from Verticillium sp., G. catenulatum and Bionectra byssicola F120, respectively [66-68]. The absolute configurations of 68 and 69 were elucidated by chemical and physicochemical methods [66]. The stereochemistries for 71-73 were proposed by analogy to other ETPs in this class based on close agreement of relevant ¹H and ¹³C NMR chemical shifts [67]. Compounds 68-70 exhibited inhibitory activity against Gram-positive bacteria and mycobacteria but no inhibitory activity against Gram-negative bacteria and fungi, and the cytotoxic effects (ED₅₀) of **68** and **69** against HeLa cells were both 0.2 µg/ml [66]. Compounds 71-73 showed antibacterial activity at 100 µg/ml, affording inhibitory zone sizes of 23 to 26 mm against Bacillus subtilis, and of 11 to 14 mm against S. aureus [67]. Compound 74 inhibited the growth of S. aureus including MRSA and QRSA with MIC values of 3 to 10 µg/ml [68]. 11,11'-Dideoxyverticillin A (75) and 11'-deoxyverticillin A (76) were originally isolated from *Penicillium* sp. [69], and 75 was also isolated from Shiraia bambusicola [70]. Their absolute configurations were assigned on the basis of NMR and CD experiments. Compounds 75 and 76 were found to exhibit potent in vitro cytotoxicity against HCT-116 with IC₅₀ values of 30 ng/ml [69], and 75 was found to be a structurally novel antiangiogenesis inhibitor [70]. The concise enantioselective total synthesis of 75 was achieved via a strategy inspired by the biosynthetic hypothesis by Movassaghi et al. [71]. The strategy should be applicable to other members of ETP family for a much thorough investigation of their synthesis, provides valuable insight into the function of enzymes involved in the biosynthesis of these natural products, and provides an access to obtain analogues which may have more promising pharmacological activity. Compounds 77-84 were isolated and characterized from fungus G. sp. [72-74]. The stereochemistries for Sch52900 (77) and Sch 52901 (78) were assumed to be the same as 68 by a direct comparison of their optical rotations and CD profile [72]. Chu et al. found 77 and 78 to be inhibitors of *c-fos* protp-oncogene induction and suggested that ETPs exerted antitumor activity by blocking a signal transduction pathway that was common to and necessary for the induction of at least a subset of immediate early genes involved in cell proliferation [72]. In vitro immersion tests showed that 68, 76-83 exhibited antinematodal activity against Caenorhabditis elegans (ED₅₀ = 30, 10, 50, 50, 25, 30, 200, 200, 200 µg/ml, respectively) and Panagrellus redicicus (ED₅₀ = 80, 40, 80, 50, 50, 25, 250, 250, 250 respectively), suggesting that $\mu g/ml$, monomeric epipolysulfanydioxopiperzaines (81-83), with indole moiety, are less active than the dimeric ones (68, 76-80) [73]. Gliocladicillins A (85) and B (86), as cancer cell proliferation inhibitors and apoptosis inducers, were identified from Cordyceps-colonizing fungus using an ecology-based approach [75]. Bionectines A-C (87-89) were isolated from B. byssicola F120 [76], and T988 A-C (90-92) were isolated from *Tilachlidium* sp. [77]. Compounds 87 and 88 exhibited antibacterial activity against S. aureus including MRSA and QRSA with MIC values of 10-30 µg/ml, and the dimeric ETP (71) showed stronger antibacterial activity with MIC values of $3-10 \,\mu\text{g/ml}$, while **89** showed no antibacterial activity even at 100 µg/ml [76]. 90-92 all showed potent cytotoxicities against cultured P388 leukemia cells with ID₅₀ values of 0.25, 2.18, and 0.56 µM, respectively [77]. The

Table 6.Structures of 59-95









81-84, 87, 88, 90, 92, 95

89, 91, 93, 94

No.	Name	\mathbf{R}_1	\mathbf{R}_2	R ₃	\mathbf{R}_4	x	у	Species	Refs.
59	Chaetocin	CH ₂ OH	Н	Н	CH ₂ OH	2	2	C. minutum	[58]
60	11,11'- Dihydroxychaeto cin	CH ₂ OH	ОН	ОН	CH ₂ OH	2	2	V. tenerum	[59]
61	Chaetocin B	CH_2OH	Н	Н	CH_2OH	3	2	C. virescens var. thielavioideum	[60]
62	Chaetocin C	CH_2OH	Н	Н	CH ₂ OH	3	3	C. virescens var. thielavioideum	[60]
63	Chetracin A	CH_2OH	ОН	ОН	CH_2OH	4	4	C. nigricolor, C. retardatum.	[60]
64	Chetomin	-	-	-	-	2	2	C. cochliodes, C. globosum, C. seminudum	[61, 62]
65	Chetoseminudin A	-	-	-	-	2	3	C. seminudum	[63]
66	Chetoseminudin B	Н	-	-	-	-	-	C. seminudum	[63]
67	Chetoseminudin C	CH ₃	-	-	-	-	-	C. seminudum	[63]
68	Verticillin A	CH ₃	ОН	ОН	CH ₃	2	2	V. sp., P. sp., G. sp., G. roseum	[66,69,72, 73]
69	Verticillin B	CH ₃	OH	ОН	CH ₂ OH	2	2	<i>V. sp.</i>	[66]
70	Verticillin C	CH ₃	OH	ОН	CH ₂ OH	2(3)	3(2)	<i>V. sp.</i>	[66]
71	Verticillin D	CH(OH)CH ₃	OH	OH	CH(OH)CH ₃	2	2	G. catenulatum, B. byssicola	[67, 68]

CH₃

(Table	6).	Contd
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No.	Name	\mathbf{R}_1	\mathbf{R}_2	R ₃	R ₄	x	у	Species	Refs.
72	Verticillin E	Ac	ОН	ОН	Ac	2	2	G. catenulatum	[67]
73	Verticillin F	CH(OH)CH ₃	ОН	ОН	Ac	2	2	G. catenulatum	[67]
74	Verticillin G	-	-	-	-	-	-	B. byssicola	[68]
75	11,11'- Dideoxyverticilli n A	CH ₃	Н	Н	CH ₃	2	2	P. sp., S. bambusicola,	[69,70]
76	11'- Deoxyverticillin A	CH ₃	ОН	Н	CH ₃	2	2	P. sp., G. roseum	[69, 73]
77	Sch 52900	CH(OH)CH ₃	ОН	ОН	CH_3	2	2	G. sp., G. roseum	[72,73]
78	Sch 52901	CH ₂ CH ₃	ОН	ОН	CH ₃	2	2	G. sp., G. roseum	[72,73]
79	Gliocladine A	CH ₃	ОН	ОН	CH ₃	3	2	G. roseum	[73]
80	Gliocladine B	CH ₃	ОН	ОН	CH ₃	4	2	G. roseum	[73]
81	Gliocladine C	CH ₃	ОН	Н	-	2	-	G. roseum	[73]
82	Gliocladine D	CH ₃	ОН	Н	-	3	-	G. roseum	[73]
83	Gliocladine E	CH ₃	ОН	Н	-	4	-	G. roseum	[73]
84	Glioclatine	CH ₃	Н	Н	-	2	-	G. roseum	[74]
85	Gliocladicillin A	CH ₃	Н	Н	CH(OH)CH ₃	2	2	Cordyceps-colonizing fungus	[75]
86	Gliocladicillin B	CH ₃	Н	Н	CH ₂ CH ₃	2	2	Cordyceps-colonizing fungus	[75]
87	Bionectin A	Н	ОН	Н	-	2	-	B. byssicola	[76]
88	Bionectin B	CH(OH)CH ₃	ОН	Н	-	2	-	B. byssicola	[76]
89	Bionectin C	Н	ОН	Н	-	-	-	B. byssicola	[76]
90	T988 A	CH ₂ OH	ОН	Н	-	3	-	Tilachlidium. sp, Plectosphaerella cucumerina	[77, 78]
91	T988 B	CH ₂ OH	ОН	Н	-	-	-	Tilachlidium. sp	[77]
92	T988 C	CH ₂ OH	ОН	Н	-	2	-	Tilachlidium. sp	[77]
93	Plectosphaeroic acid A	CH ₂ OH	ОН	a	-	-	-	Plectosphaerella cucumerina	[78]
94	Plectosphaeroic acid B	CH ₃	ОН	a	-	-	-	Plectosphaerella cucumerina	[78]
95	Plectosphaeroic acid C	CH ₂ OH	ОН	a	-	3	-	Plectosphaerella cucumerina	[78]

results indicated that the cytotoxic and antibacterial activities of the ETP class of compounds were mediated in part due to the sulfide bridge in the ETP moiety, while the number of sulfur atoms in the bridge may not significant influence the activity. Plectosphaeroic acids A-C (**92-95**) were all isolated from the fungus *Plectosphaerella cucumerina* together with **90** [78]. Notablely, **93-95** all could inhibite purified recombinant human IDO, a promising molecular target for treating cancer, with identical IC₅₀ values of $\approx 0.2 \,\mu$ M, while **90** was completely inactive. Meanwhile, the synthetic cinnabarinic acid was also found to be active in this assay with an IC₅₀ value of $\approx 0.2 \ \mu$ M, indicating that some portion of the phenoxazineone fragment may play a key role for inhibitory effect to IDO (compounds **59-95** are shown in Table **6**).

2.7. Leptosins

Leptosins A-S (96-119), antitumor metabolites from *Leptoshaeria* sp., were discovered by Numata and his coworkers [79-84]. Compound 96 represented the first dimeric ETP with a different configuration at C-5'a and C-10'b respect to other related compounds, such as chaetoins

Table 7.Structures of 96-119











No.	Name	R ₁	\mathbf{R}_2	R ₃	\mathbf{R}_4	x	у	ED ₅₀	Species	Refs.
96	Leptosin A	CH ₂ OH	ОН	OH	CH(CH ₃) ₂	4	2	1.85 ng/ml	L. sp.	[79]
97	Leptosin B	CH ₂ OH	ОН	OH	CH(CH ₃) ₂	3	2	2.4 ng/ml	L. sp.	[79]
98	Leptosin C	CH ₂ OH	OH	OH	CH(CH ₃) ₂	2	2	1.75 ng/ml	L. sp.	[79]
99	Leptosin D	CH(CH ₃) ₂	ОН	Н	-	2	-	86 ng/ml	L. sp.	[79]
100	Leptosin E	CH(CH ₃) ₂	OH	Н	-	3	-	46 ng/ml	L. sp.	[79]
101	Leptosin F	CH(CH ₃) ₂	OH	Н	-	4	-	56 ng/ml	L. sp.	[79]
102	Leptosin G	CH ₂ OH	OH	OH	$CH(CH_3)_2$	4	3	4.6 ng/ml	L. sp.	[80]
103	Leptosin G ₁	CH ₂ OH	OH	OH	$CH(CH_3)_2$	3	3	4.3 ng/ml	L. sp.	[80]
104	Leptosin G ₂	CH ₂ OH	OH	OH	$CH(CH_3)_2$	2	3	4.4 ng/ml	L. sp.	[80]
105	Leptosin H	CH ₂ OH	OH	OH	CH(CH ₃) ₂	2	4	3.0 ng/ml	L. sp.	[80]
106	Leptosin I	CH ₂ OH	Н	Н	-	-	-	1.13 µg/ml	L. sp.	[81]
107	Leptosin J	Н	CH ₂ OH	Н	-	-	-	1.25 µg/ml	L. sp.	[81]
108	Leptosin K	CH(CH ₃) ₂	OH	OH	$CH(CH_3)_2$	2	2	3.8 ng/ml	L. sp.	[82]
109	Leptosin K ₁	CH(CH ₃) ₂	OH	OH	$CH(CH_3)_2$	2	3	2.2 ng/ml	L. sp.	[82]
110	Leptosin K ₂	CH(CH ₃) ₂	OH	OH	$CH(CH_3)_2$	2	4	2.1 ng/ml	L. sp.	[82]
111	Leptosin M	Н	OH	OH	Н	4	-	1.05 µg/ml	L. sp.	[83]
112	Leptosin M ₁	Н	OH	OH	Н	2	-	1.40 µg/ml	L. sp.	[83]
113	Leptosin N	OH	Н	Н	ОН	4	-	0.18 µg/ml	L. sp.	[83]
114	Leptosin N ₁	OH	Н	Н	ОН	3	-	0.19 µg/ml	L. sp.	[83]
115	Leptosin O	S_2CH_3	$CH(CH_3)_2$	-	-	-	-	1.1 µg/ml	L. sp.	[84]
116	Leptosin P	CH(CH ₃) ₂	S_2CH_3	-	-	-	-	0.1 µg/ml	L. sp.	[84]
117	Leptosin Q	SCH ₃	CH(CH ₃) ₂	-	-	-	-	14.8 µg/ml	L. sp.	[84]
118	Leptosin R	CH(CH ₃) ₂	SCH ₃	-	-	-	-	15.2 µg/ml	L. sp.	[84]
119	Leptosin S	NH ₂	$CH(CH_3)_2$		-	-	-	10.1 µg/ml	L. sp.	[84]

Table 8.Structures of 120-129



No.	Name		x	Species	Refs.
120	Sirodesmin A		2	Sirodesmium diversum	[85, 86]
121	Sirodesmin B	Ac	4	Sirodesmium diversum	[85, 86]
122	Sirodesmin C	Ac	3	Sirodesmium diversum	[85, 86]
123	Sirodesmin PL (Sirodesmin G)		2	Sirodesmium diversum, Phoma lingam	[85, 86, 87]
124	Sirodesmin H		1	Sirodesmium diversum, Phoma lingam	[86, 88]
125	Sirodesmin J		3	Sirodesmium diversum, Phoma lingam	[86, 89]
126	Sirodesmin K	Ac	4	Sirodesmium diversum, Phoma lingam	[86, 89]
127	Desirodesmin PL	Н	2	Phoma lingam	[87]
128	Phomalirazine		-	Phoma lingam	[90]
129	Dithiosilvation	-	-	A. silvaticus	[91]

and chetracin A. Compounds 96 and 98 exhibited significant antitumor activity against Sarcoma 180 ascites (%T/C 260 and 293, respectively) at doses of 0.5 mg/kg and 0.25 mg/kg, respectively [79]. Compounds 96-119 all showed potent cytotoxicity against cultured P388 cells (shown in Table 7). It was found that dimeric ETPs (96-98 and 102-105) showed more potent activity than the monoeric ETPs (99-101) with the indole moiety, and the number of sulfur atoms in dioxopiperazine rings appered no influence on the cytotoxiciy against P388 cells. Although the configuration of sulfur bridge in 108-110 differ from that in 96-98 and 102-105, their cytotoxic effects (ED₅₀) was almost identical, indicating that the configuration of sulfur bridge may not influence the cytotoxicity against P388. However, the cytotoxicities of 108-110 are much strong than those of 111-114, indicating the disulfide or polysulfide functionalities significantly influence their cytotocixities. Further, 113 and 114 are almost 10-fold potent than those of 111 and 112 in the cytotoxic bioassay against P388, indicating that the position of hydroxyl group may influence the cytotoxicity. It may be worthy to note that 111 showed appreciable cytotoxic activities against the 39 human cancer cell lines. Moreover, evaluation of the pattern of differential cytotoxicity using the COMPARE program suggested the possibility that the mode of action for 111 might be different from that shown by any other anticancer drug developed to date [83].

2.8. Sirodesmins and other ETPs

Sirodesmins A-C (**120-122**), D-F (without structure reported), PL (sirodesmin G, **123**) and H-K (**124-126**) were all originally isolated from *Sirodesmium diversum* [85, 86].

Compounds 123-126 were also isolated from Phoma lingam [87-89] as well as deacetylsirodesmin PL (127) [87] and phomalirazine (128) [90]. The structure of 120 was determined by X-ray analysis of its diacetate derivative, and the structures of 121-123 were determined by comparison of their chemical and spectroscopic properties with those of 120. The structure of 128, including absolute configuration, was unambiguously elucidated by X-ray diffraction analysis. In addition, the structure of 128 has important implications on the biogenetic pathway of this broad class of sulfur bridged dioxopiperazines [90]. Dithiosilvatin (129), a possible key intermediate in the biosynthesis of ETP, was isolated from A. silvaticus [91]. Compound 129 appeared to be biosynthesized biogenetically from a compound composed from tyrosine and alanine (or serine) as in the synthesis of gliotoxin (1) or from syrosine and another amino acid other than glycine, and its absolute configuration was determined by comparison of its CD profile with that of emestrin (6). Compounds 120-123 showed exceptionally high activity against small RNA-containing virus such as the rhinoviruses and the enteroviruses [85]. For example, 120 could reduce the growth of picornaviruses in cultures of human diploid lung cells by 50% at concentration of 8 ng/ml. In contrast, the concentration of 120 required to cause 50% cytotoxicity in uninfected cultures of the same cells is 1 μ g/ml. However, **120** had been shown to produce chromosome abnormalities in in vitro studies with human lymphocytes and in in vivo studies in Chinese hamster bone marrow cells. The phytotoxicity of 124, the first reported naturally occurring monosulfur bridged dioxopiperazine, was about ten times less than that of 123 [88]. Whether the lower toxicity was due to the absence of the disulfur bridge or was

due to the lability of the monosulfide remained to be determined. Pedras et al. had studied the effect of acetylation on the biological activity of 123, and found that acetylation at the 6,14-OH or 14-OH reduced the phytotoxicity considerably. whereas the phytotoxicity was not considerably affected by acetylation at the 6-OH, deducing that 14-OH may be involved in the mechanism of the toxin [89]. Rouxel et al. found that 123 can strongly and rapidly inhibit the incorporation $[^{14}C]$ uridine in RNAs by 5.2 μ M in liquid medium, and the toxicity, attributed to the reactivity of its disulphide bridge, could be reversed by the metals of the II B series (Zn, Hg and Cd). These results suggested an interaction of 123 with essential Zn from Zn-containing metalloenzymes such as RNA polymerases [92]. Boudart found Gram-positive bacteria were particularly susceptible to 123, whereas the growth of Gram-negative bacteria was more or less affected. Meanwhile, the antibacterial activity of 123 was stronger than its monosulfide derivative and dithiomethyl-related derivative (totally inactive), demonstrating the importance of disulfide bridge of the molecule in antibacterial activity [93] (compounds 120-129) are shown in Table 8).

3. BIOSYNTHESIS

In spite of almost of 50 years of research on ETPs since the structure of gliotoxin (1) was first described and its biosynthesis has been being an intensive research subject, very little is known about ETPs biosynthesis [16]. Labeling and feeding experiments, the most used techniques in the biosynthetic study, were carried out leading to uncover that amino acids and cyclic dipeptides were the precursors or intermediates for gliotoxin and sirodesmin [94-99]. In the pathway to biosynthesize ETPs from Phoma lingam, the introduction of sulfur atoms could occur immediately following the cyclodipeptide formation, and in addition to the cyclic dipeptide phomamide two other intermediates, deacetylsirodesmin PL (125) and phomarilazine (126), have been identified [89]. Hence, the core ETP structure is formed at an early stage in the biosynthetic pathway. On the other hand, inspection of the chemical structure of 126 seems to indicate that N-mehtylation of the dioxopiperazine would occur after the oxidative cyclization onto the phenyl ring [90]. Introduction of the sulfur atoms in to the core ETP moiety is poorly understood. Labeling experiments suggested that the sulfur atoms in ETPs may be derived from methionine, cysteine, and sodium sulfate, however, how the sulfur atoms are introduced is unknown [16]. Genes responsible for biosynthesis of fungal secondary metabolites were found to be usually tightly clustered in the genome and co-regulated with metabolite production. Recently, the availability of identification of the putative ETP biosynthetic gene cluster is a major step forward in understanding how these molecules are produced [16, 100-102]. Herein, according to the literature surveys [2, 16, 23, 37, 44, 50, 52, 71, 90-92, 94-99], the plausible biosynthetic hypotheses of some ETPs are summarized with our meager knowledge (Fig. 1)

4. ANTITUMOR ACTIVITY

The effect of ETPs on antitumor has been studied extensively and the mechanism of toxicity have been the

subject of a number of reviews in 1996 and 2005 [15, 16]. Gardiner *et al.* [16] considered that the toxicity was mediated in at least two ways: (1) conjugation to proteins with susceptible thiol residues and subsequent inactivation, and (2) generation of reactive oxygen species *via* redox cycling. However, most of research on toxicity limited only to gliotoxin (1). The mechanism of toxicity was remains to be investigated.

Recently, some new results on antitumor activity of ETPs were reported. For instance, 1 was found to be a dual inhibitor of farnesyltransferase and geranylgeranyltansferase I with pronounced antitumor activity and favorable toxicity profile in breast cancer in vitro and in vivo [14]. Axelsson et al. found that glutathione promoted gliotoxin (1)-induced cytotoxicity in human neuroblastoma SH-SY5Y cells by reducing the ETP disulfide bridge to the dithiol form [103]. Compound 1 can induce apoptosis of cancer cells in culture and human cancer xenografts in transplant SCID mice, and this anticancer activity was further confirmed by electromicroscopic observations [104]. Sporidesmin A (18) found inactivate human glutaredoxin was to (thioltransferase), a thiol-disulfide oxidoreductase, in a timeand concentration-dependent manner via a proposed initial reaction between the toxic sulfurs and cysteine 22 in the glutaredoxin active site [105]. This study implicates modification of sulfhydryls of target proteins in some of the cytotoxic effect of ETPs. Chaetocin (59) was found to be a promising compound with potent antimyeloma activity in IL-6-dependent and -independent myeloma cell lines in vitro and in vivo, and its antimyeloma activity was mediated primarily via imposition of oxidative stress and consequent apoptosis induction [106]. Recently, 59 and its enantiomer were found to inhibit H3K9 HMT G9a, a potential therapeutic target for treating human cancer. Both compounds were found equally effective with IC₅₀ values of 2.4 and 1.7 μ M. In contrast, its sulfur-deficient (±)-analogues were less active (IC₅₀ > 50 μ M) indicating that the sulfur functionality was crucial for the biological activity. Meanwhile, its enantiomer showed strong apoptosisinducing activity through caspase-8/caspase-3 activation [107]. Staab et al. found that chetomin (64) disrupted the interaction of HIF-1, an important therapeutic target for solid tumor therapy, with the transcriptional coactivator p300. HIF-1 inhibition by 64 effectively reduced hypoxiadependent transcription and radiosensitized hypoxic HT 1080 human fibrosarcoma cells in vitro [108]. The mechanistic studies revealed that 64 and other ETPs blocked the interaction between HIF-1 α and p300 by a zinc ejection mechanism and the structure-activity studies revealed that the structurally unique ETP core was required and was sufficient to block the interaction of HIF-1 α and p300 [109]. Gliocladicillins A (85) and B (86) were found to inhibit proliferation of HeLa, HepG2, MCF-7 tumor cells and in treated HeLa cells, to induce cell cycle arrest in the G₂/M phase, through activation of the p53 pathway and causing cvclin B accumulation. Compounds 85 and 86 also induced tumor cell apoptosis by activating both intrinsic and extrinsic pathways and they also exhibited significant in vivo antitumor activity in mice. These results suggest that 85 and 86 are effective antitumor agents in vitro and in vivo, and should be further evaluated for their potential in clinical



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Fig. (1). Proposed biosynthesis hypothesis of ETPs.

usage [75]. 11,11'-dideoxyverticillin A (**75**) was found to act as a potent angiogenesis inhibitor, which inhibited angiogenesis, required for tumor growth, *via* various steps and through interfering with VEGF signal transduction [70] and to possess growth factor receptor tyrosine kinaseinhibitory effect with anti-tumor activity against four human breast tumor cell lines with an average IC₅₀ value of 0.2 μ M [110]. Yanagihara *et al.* found leptosins C (**98**) and F (**111**) inhibited DNA topoisomerases (topo) I and/or II, targets of anticancer agents, and induced apoptosis by inactivation of Akt/protein kinase B [111].

5. CONCLUSIONS

To date, about 120 ETPs with broad-spectrum bioactivities have been identified from the fungi, and the members of ETPs family are still expanding. The wide range of bioactivities of ETPs are still explored extensively, and SAR studies of ETPs suggest that disulfide or polysulfide moieties play the key role for their bioactivities, which is particularly important for synthesis and design of novel anticancer drug candidates. Although it is challenging to understand the biosynthetic pathways for ETPs, the development in identification of the putative ETP biosynthetic gene cluster will make biosynthetic pathway to be understood. Meanwhile, the plausible biosynthetic hypotheses, that ETPs are derived from amino acid and cyclic dipeptides, can provide access to synthesize natural ETPs and their analogues with high bioactivity. Demand for ETPs in medicinal lead identification and in understanding biosynthesis and mechanism of activities will lead to continue the research in this field.

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ABBREVIATIONS

CCR2 =	Chemokine	C-C motif	receptor 2

- CD = Circular dichroism
- DNA = Deoxyribonucleic acid
- $EC_{50} = 50\%$ effective concentration
- $ED_{50} = 50\%$ effective dose
- EGF = Epidermal growth factor
- HIF-1 = Hypoxia-inducible factor-1
- HIV-1 = Human immunodeficiency virus type 1
- hMCP-1 = Human monocyte chemoattractant protein-1

HMTs	=	Histone methyltransferases
IC ₅₀	=	50% inhibitory concentration
ID ₅₀	=	50% infective dose
IDO	=	Indoleamine 2,3-dioxygenase
МСР	=	Monocyte chemoattractant protein
MIC	=	Minimum inhibitory concentration
MRSA	=	Methicillin-resistant Staphylococcus aureus
MTT	=	3-(4,5-Dimethylthiazol-2-yl)-2,5- diphenyltetrazolium bromide
NMR	=	Nuclear magnetic resonace
ORD	=	Optical rotatory dispersion
QRSA	=	Quinolone-resistant Staphylococcus aureus
RNA	=	Ribonucleic acid
SAR	=	Structure-activity relationship
SCID	=	Severe combined immunodeficiency
0/T/C	_	Detion of the suminal time of the treate

- %T/C = Ratio of the survival time of the treated animal group (T) and that of the control group (C)
- VEGF = Vascular endothelial growth factor

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