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Griseococcin (1) from *Bovistella radicata* (*Mont.*) *Pat* and antifungal activity



Yong Ye^{1,2}, Qinghua Zeng^{1,2} and Qingmei Zeng^{1,2*}

Abstract

Background: To evaluate the antimicrobial and microbicidel activity of *B. radicata* fermentation broth, the broth was purified by DEAE-cellulose and sephadex LC-20 column. The compounds were submitted to spectral analyses (HPLC, FT-IR, 1D and 2D NMR etc.).

Results: The purified compounds were identified as the Griseococcin(s) which were naphthoquinone derivatives, the Chemical formula and MW of Griseococcin (1) was determined as $C_{37}O_{10}H_{43}N$ and 661 Da. only Griseococcin (1) has good antimicrobial activity among the Griseococcin(s). The zone of inhibition (ZOI), minimum inhibitory concentration (MIC) and minimum bactericidal concentration (MBC) or minimum fungicidal concentration (MFC) of Griseococcin (1) were used to investigate the antimicrobial activity. Antifungal activity of Griseococcin (1) was significant, especially for main pathogenic fungus *Trichophyton rubrum* and *Trichophyton mentagrophytes*, MFC/MIC of Griseococcin (1) was 1, while MFC/MIC of postive control was greater than 4, the fungicidal effect of Griseococcin (1) was better than that of positive control.

Conclusions: In this paper, the secondary metabolite compound Griseococcin (1) from *B. radicata* was purified. The purified compound can restrain main pathogens (*T. rubrum* and *T. mentagrophytes*) leading to tinea pedis. The antifungal activity of Griseococcin (1) was similar to that of the positive control and the fungicidal effect of Griseococcin (1) was better than that of positive control, it might be suitable for pharmaceutical industries.

Keywords: Griseococcin, LH-20, DEAE, 1D NMR, 2D NMR, HPLC, FT-IR

Background

Tinea pedis is a chronic fungal infection of the feet [1]. Patients that have tinea pedis may be affected by several pathogens, including filamentous fungi named *Trichophyton rubrum* and *Trichophyton mentagrophytes* [2], as well as a yeast named *Candida albicans* [3]. *T. rubrum* is the main pathogenic fungi for tinea pedis, having a prevalence as high as 80% among all tinea-pedis associated pathogenic microbes [4]. Traditionally, to treat tinea pedis, synthetic fungicides such as fluconazole, itraconazole, echinocandins [5], and miconazole nitrate,

Puffballs are widely distributed in many provinces of China, and are various by more than 100 species [16].

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either by oral medication or external use [6], have been used to treat this disease. Vermes et al. (2000) found that flucytosine and AMB (amphotericin B) were moderately effective in fighting against invasive fungal infections [7–9]. Similar studies on itraconazole have demonstrated that it is effective against fungal infections [10]. However, due to side effects or the continuous drug resistance, some oral medications are unsafe for patients [11], and these chemicals also cause potential deleterious effects on the environment due to their residues [12, 13]. In general, plant natural products have been for decades one of the most successful sources of drugs to treat infectious diseases [14] and natural products extracted represent a rich resource for screening bioactive compounds [15].

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Ye et al. BMC Microbiology (2020) 20:276 Page 2 of 9

Calvatia gigantean (Batsch ex Pers) Lloyd, Calvatia lilacina (Mont.et Berk.) Lloyd, Lasiosphaera fenzlii Reich, Lycoperdon pyriforme Schaeff.:pers, Bovistella radicata (Mont.) Pat, Handkea utriformis (HU), H. excipuliformis (HE), and Vascellum pratense (VP) are all common medicinal puffballs. Although no longer edible in their mature state (because of their powdery consistency), these puffballs have been shown to be a source of active compounds of various biological activities. Puffballs are believed to have several therapeutic properties when used medicinally: hemostasis [17], cough relief [18], suppression of cell division, and antitumor [19] and antimicrobial [20] properties. Petrović P, et al. reported noticeable antimicrobial activity diversity for the methanol extracts obtained from Handkea utriformis (HU), H. excipuliformis (HE), and Vascellum pratense (VP) [21]. Specimen (Bovistella radicata (Mont.) Pat) was dried and deposited in the Institute of Biology, School fo Food and Biological Engineering, Hefei University of Technology (HFUT), China.

The aim of the present study was to evaluate the antimicrobial activity of Griseococcin (1) extracted from *B. radicata* fermentation broth. The antimicrobial and microbicidel activities were evaluated in terms of their minimum inhibitory concentration (MIC), minimum

fungicidal concentration (MFC) or minimum bactericidal concentration (MBC) and zone of inhibition (ZOI) values [22], the physico-chemical characterization (HPLC, UV, FT-IR) of Griseococcin (1) and the chemical constituents responsible for this activity were also studied (1D and 2D NMR).

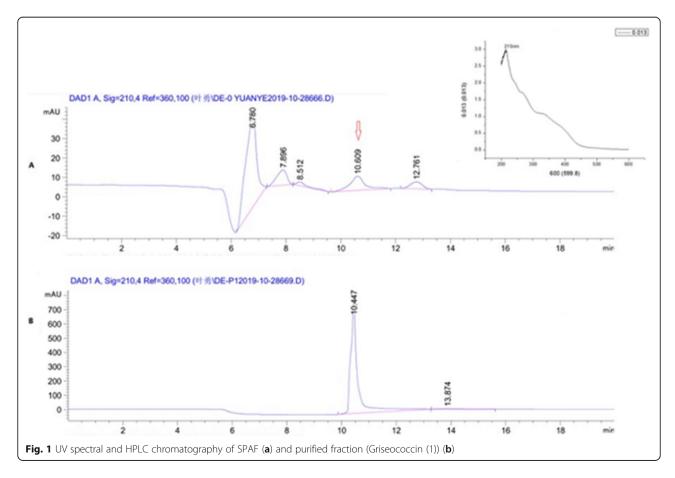
Results

Fermentation, extraction and purification of active compound from *B. radicata*

20% NaCl elution fraction from fermentation broth of *B. radicata* was named as SPAF by DEAE-cellulose column. The strongest antimicrobial activity fraction from SPAF was Griseococcin (1) by Sephadex LH-20 column. The UV $_{\rm max}$ of all the fraction was 215 nm, the HPLC chromatograms of SPAF and Griseococcin (1) were shown in Fig. 1(a ~ b). The chromatogram of B showed a single and symmetrical peak for Griseococcin (1) (Fig. 1b).

1D and 2D NMR of Griseococcin (1)

Griseococcin (1) was isolated as a white amorphous solid powder with the molecular formula of $C_{37}H_{43}NO_{10}$ derived from the high-resolution electrospray ionization mass spectrum (HR-ESI-MS). The complete assignments



Ye et al. BMC Microbiology (2020) 20:276 Page 3 of 9

for all protons and carbons were shown in Table 1. The ¹³C NMR spectra of Griseococcin (1) displayed signals of thirty seven carbons, including five carbonyl carbons $(\delta C215.7-175.1)$, five aromatic/olefinic methine carbons (δC 128.86, δC215.7-175.1), seven non-protonated aromatic/olefinic carbons (δC 161.06-109.99), four methyl carbons (δ C20.27)), and four olefin carbons (δC 166.01). The ¹H NMR spectrum of 1in D₂O exhibited signals of four methyls at δ H 2.14 (3H, s, H-14'), δ H 2.12 (3H, s, H-15'), δ H 1.06 (3H, s, H-16') and 1.07 (3H, s, H-17'), five aromatic protons δH 7.80 (1H, s, H-1), δH 7.93(1H, s, H-5), δH 7.72 (1H, s, H-6), δH 7.81 (1H, s, H-8) and 7.66 (1H, s, H-12)], four hydroxyl groups at δ H 8.37 (1H, br s, 4'-OH), δ H 7.81 (1H, br s, 9'OH) and δ H 7.80 (1H, br s, 11'-OH) and 9.63 (1H, br s, 13'-OH).

The structure of Griseococcin (1) was deduced by comprehensive analysis of ¹H-¹HCOSY, HMBC, and HSQC spectra (Fig. 2a). In Griseococcin (1), the naphthoquinone substructure could be identified by the observation of HMBC correlations from H-8 (δH 7.80) to C-6 (δ C 137.21), C-4 (δC 138.60) and C-13 (δC 30.18), from H-1 (δ H 7.81) to C-3 (δ C 175.11), C-12 (δ C 166.07) and C-1' (δC 28.40), from H-5 (δH 7.93) to C-3 $(\delta C 175.11)$ and C-9 ($\delta C 138.56$), from H₂-13 ($\delta H 1.07$) to C-8 (δ C 135.45) and C-6 (δ C 137.21), from H₃-14' $(\delta H \ 1.85)$ to C-2' $(\delta C \ 215.7)$ and C-4' (OH) $(\delta C \ 73.60)$, from H₃-15' (δH 2.11) to C-6' (δC 215.70) and C-4' (OH) (δ C 73.60), from H₂-7' (δ H 1.08) to C-9' (δ C 71.25) and C-13' (δ C 71.18). The ¹H, ¹H three-bond couplings observed in the COSY spectrum from H-8' $(\delta H \ 1.94)$ to H-9' $(\delta H \ 3.62)$, from H-10' $(\delta H \ 1.29)$ to H-11' (δ H 3.49), from H-12' (δ H 1.73) to H-13' (δ H 3.51), together with the chemical shifts of the ¹³C resonances (C-8'-13') observed at alternating higher and lower fields, revealed the presence of cyclohexane with alternating hydroxyl and methyl groups. ¹H-¹H COSY correlations from H_2 -13 (δ H1.07, m) to H_2 -14 (δ H3.62, m), from H_2-14 ($\delta H_3.62$, m) to H_2-15 ($\delta H_3.49$, m) and from H_2 -16 (δH 3.55, m) to H_2 -17 (δH 3.51, m) and HMBC correlations from H_2 -13 (δH 1,07, m) to C-15 $(\delta C \ 166.02)$, from H₂-14 $(\delta H \ 3.62)$, m) to C-16 $(\delta C \ 166)$, from H_2 -15 (δ H3.49, m) to C-17 (δ C 166.01) and from H_2 -16 (δH 3.55, m) to C-18 (δC 23.15) identified coupled olefins. The key HMBC correlations from H_2 –1′ (δ H1.94, m) to C-3′ (δ C 23.4), from H – 3′ (δ H2.14, m) to C-5′ (δ C 29.05), from H_3 –14′ (δ H1.85, m) to C-2′ (δ C 215.7) and C-4′-OH (δ C73.6), from H3–15′ (δ H2.11, m) to C-6′ (δ C 215.7) and C-4′-OH identified two meta position carbonyl group and one ortho position hydroxyl group (Fig. 2b).

This connectivity was also secured by the observation of the HSQC correlations from H_3 –14′ to C-3′ and from H_3 –15′ to C-6′. Therefore, the complete structure of naphthoguinone was determined as shown in Fig. 2c.

Physico-chemical characterization of Griseococcin (1)

Griseococcin (1) was white powder and its solubility was 0.063 g/ml in water. It could be slight soluble in methanol and DMSO, but insoluble in n-hexane, dichloromethane, chloroform, ethyl acetate and acetone.

The FT-IR spectrum of Griseococcin (1) showed (Fig. 3) an intense and broad characteristic absorption peaks at 3417.2 cm⁻¹ represented the stretching vibration of O–H. The weak absorption peaks at 2356 and 2925.5 cm⁻¹ were resulted from the stretching vibration of C–H. The absorption bands at 1637.4 and 1618.1 cm⁻¹ are due to the vibration of C=C and C=O in the ester group. The absorptions peaks at 1456.1, 1414 and 624 cm⁻¹ were attributed to the presence of an internal C–H deformation. The strong absorption peak at 866 cm⁻¹ was resulted from aromatics. In conclusion, the typical absorption peak indicated that Griseococcin (1) was naphthoquinone with group O–H,C-H,C=C,C=O and so on [23].

In vitro antagonistic assay

Griseococcin (1) was assessed for antimicrobial and microbicidel activity against selected *Trichophyton rubrum* (ATCC 28188), *Trichophyton mentagrophytes* (ATCC 9533), *Epidermophyton floccosum* (ATCC 52066), *Candida albicans* (ATCC 10231), *Staphylococcus aureus* (ATCC 6538), *Bacillus subtilis* (ATCC 6051), *Escherichia coli* (ATCC 8739) and *Pseudomonas aeruginosa* (ATCC 27582) The results were shown in Table 1, it displayed strong antifungal activity against *T. rubrum*, *T. mentagrophytes* with ZOI values of 18.06 ± 0.85, 15.01 ± 1.02 mm, as compared to the positive control

Table 1 Zone of inhibition (ZOI) of Griseococcin (1) on microbial strains

Pathogenic	Zone of inhibition (mm)					
	T. rubrum	T. mentagrophytes	E. floccosum	C. albicans		
Griseococcin (1)	18.1 ± 0.9	15 .0 ± 1.0	5.0 ± 2.3	1.2 ± 2.2		
Terbinafine	20.7 ± 1.6	28.3 ± 2.2	10.02 ± 1.02	6.3 ± 0.6		
	S. aureus	E. coli	B. subtilis	P. aeruginosa		
Griseococcin (1)	10.0 ± 1.9	8.0 ± 1.1	3.0 ± 1.2	3.7 ± 1.4		
Gentamicin sulfate	31.7 ± 1.5	28.5 ± 1.4	10.0 ± 1.0	33.6 ± 1.4		

Ye et al. BMC Microbiology (2020) 20:276 Page 4 of 9

with $ZOI = 20.67 \pm 1.58$, 28.33 ± 2.15 mm, respectively. While antibacterial activity was weak.

J.Meletiadis et al. reported that compounds were considered bactericidal or fungicidal when the MBC/MIC or MFC/MIC ratio is ≤ 4 [24]. In this study, it was inportant to discern whether the Griseococcin (1) possesses bactericidal and fungicidal activities. The fungicidal activities of Griseococcin (1) were assessed as MIC, MFC and MFC/MIC. The results were shown in Table 2. Griseococcin (1) showed the high fungicidal activities by means of lowest values of MIC and MFC against the four fungi, especially for main pathogenic fungi (T. rubrum), the MIC, MFC and MFC/MIC values were 31.2 ± 2.7 , $31.2 \pm$ 3.1 µg/ml and 1, while MIC, MFC and MFC/MIC values of Terbinafine were 15.6 ± 1.2 , $31.2 \pm 1.6 \,\mu\text{g/ml}$ and 6. Fungicidal activities of Griseococcin (1) are revealed more effective than that of commercial reagents (Terbinafine).

Griseococcin (1) also showed high bactericidal activities with MIC, MBC and MBC/MIC values ranged between $62.5 \sim 125 \,\mu\text{g/ml}$, $125-500 \,\mu\text{g/ml}$ and 2-4 against

examined bacteria (*S. aureus*, *E. coli* and *P. aeruginosa*). The results were shown in Table 3. Griseococcin (1) showed the highest bactericidal activity for *S. aureus* and *E. coli*, MBC/MIC value of Griseococcin (1) was 2.0, while the MBC/MIC ratio was 3.0 and 4.0 for positive control (Gentamicin sulfate).

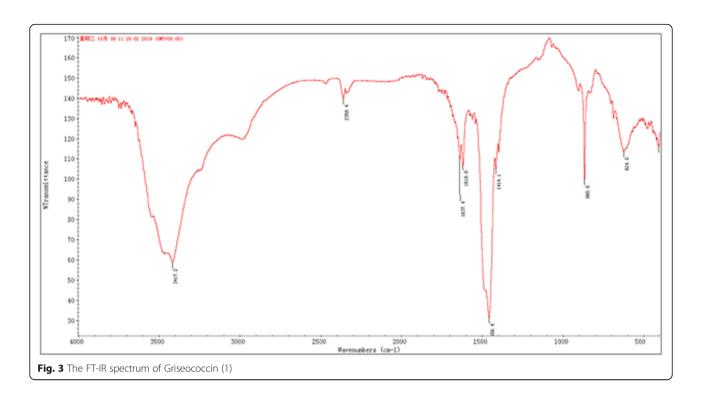
Due to side effects and the continuous drug resistance, commercial reagents might be unsafe for patients [11], Therefore, the development of fungicidal therapies is crucial, above results (MIC, MFC or MBC and MFC/MIC) add more value to Griseococcin (1).

Discussion

In the present study, Griseococcin (1) purified from selected puffball (*Bovistella radicata (Mont.) Pat*) had remarkable antifungal activities. These data are consistent with previous findings on the minimum inhibitory concentrations (MICs) and zone of inhibition (ZOI) of *B. radicata* fermentation [20].

According to the Chinese Pharmacopeia, the puffball can restrain *S. aureus* and *P. aeruginosa*. The antifungal

Ye et al. BMC Microbiology (2020) 20:276 Page 5 of 9



function of puffball has not been reported previously, hence, the present study is interesting and original. The novel application of *B. radicata* might be due to different geographic sources of the material used and different strains used [25].

In this study, the purification extraction Griseococcin(s) from fermentation broth of *B. radicata* obtained through celluous DE-52 and sephadex LH-20 column. In Vitro study on antifungal effects of Griseococcin (1) on fungi showed that the most sensitive fungi strains were the main pathogenic fungi (*T. rubrum* and *T. mentagrophytes*) causing tinea pedis, ZOIs were 18.06 ± 0.85 and 15.01 ± 1.02 mm, MICs were 31.2 ± 2.7 and 31.2 ± 1.8 µg/ml, MFCs were 31.2 ± 3.1 µg/ml and 31.2 ± 2.1 µg/ml, MFC/MICs were 1 and 1 against *T. rubrum* and *T. mentagrophytes*. ZOI values of positive control (Terbinafine)

were $20.67 \pm 1.58 \,\text{mm}$ $28.33 \pm 2.15 \,\text{mm}$, MICs were 15.6 ± 1.6 and $7.8 \pm 1.2 \,\mu\text{g/ml}$, MFCs were 93.6 ± 2.1 and $39.0 \pm 2.2 \,\mu\text{g/ml}$, MFC/MICs were 6 and 5 respectively. The antifungal effect of Griseococcin (1) was similar with that of positive control, the fungicidal effect of Griseococcin (1) was better than that of positive control. The most sensitive bacterial species for Griseococcin (1) was S. aureus and E. coli, MICs and MBCs were $62.5 \pm$ 1.5, 125 ± 2.3 and 125 ± 3.1 , $250 \pm 2.1 \,\mu\text{g/ml}$ respectively, P. aeruginosa and B. subtilis were more resistant. MFC/ MICs and MBC/MICs of Griseococcin (1) were less than positive control which meant that antibacterial activity Griseococcin (1) was better than that of the commercial drugs. This study is important for the development of new drugs with low toxicity, overcoming drug resistance and recurrence.

Table 2 MIC, MFC and MIC/MFC of Griseococcin (1) on fungal strains

Pathogenic fungi	MIC (μg/ml)	MIC (μg/ml)				
	T. rubrum	T. mentagrophytes	E. floccosum	C. albicans		
Griseococcin (1)						
MIC (μg/ml)	31.2 ± 2.7	31.2 ± 1.8	250.0 ± 5.2	250 ± 7.6		
MFC (µg/ml)	31.2 ± 3.1	31.2 ± 2.1	500 ± 3.1	500 ± 6.7		
MFC / MIC	1	1	2	2		
Terbinafine						
MIC (μg/ml)	15.6 ± 1.6	7.8 ± 1.2	62.5 ± 2.1	62.5 ± 2.6		
MFC (µg/ml)	93.6 ± 2.1	39.0 ± 2.2	250.0 ± 2.1	125 ± 2.6		
MFC/ MIC	6	5	4	2		

Ye et al. BMC Microbiology (2020) 20:276 Page 6 of 9

Table 3 MIC, MBC and MIC/MFC of Griseococcin (1) on bacterial strains

Pathogenic bacteria	MIC (μg/ml)				
	S. aureus	E. coli	B. subtilis	P. aeruginosa	
Griseococcin (1	1)				
MIC (µg/ml)	62.5 ± 1.5	125 ± 2.3	125 ± 1.5	62.5 ± 3.6	
MBC (µg/ml)	125 ± 3.1	250 ± 2.1	> 500	250 ± 6.7	
MBC / MIC	2.0	2.0	> 4.0	4.0	
Gentamicin sul	lfate				
MIC (μg/ml)	15.6 ± 1.3	31.2 ± 2.4	31.2 ± 3.0	62.5 ± 2.6	
MBC (µg/ml)	46.8 ± 1.6	124.8 ± 2.2	202.8 ± 2.1	350 ± 2.2	
MBC/MIC	3.0	4 .0	6.5	5.6	

The FT-IR spectrum of Griseococcin (1) showed the strong absorption band, stretching vibration and bending vibration of O-H, C=O, C=C and C-H which belong to a unsaturated coupled bond and aromatic form of naphthoquinone. According to HR-ESI-MS analysis, MW of Griseococcin (1) was 661 Da. Based on the results of different spectral (HPLC, FT-IR, DSC, 1D and 2D NMR etc.) studies and physicochemical properties, the molecular formula of Griseococcin (1) was $C_{37}H_{43}NO_{10}$ and the molecular structure of Griseococcin (1) was shown in Fig. 1. MBC/MICs of were 6 and 5 respectively.

Previously, many authors reported the various biological activity of fermentation broth from puffball like anticancer activity [26, 27], antioxidant activity [28], antifatigue effect [29], etc. In the present study, the antifungal activity of *B. radicata* was another important biological function. The biological activities of organic compounds are related to their molecular weight, functional groups, the length of chain, the composition of group and the number of branches, hydrophilic and hydrophobic group. It means that the structure-activity relationship should be disclosed.

Conclusions

Future work concentrating on determining the antifungal mechanisms of Griseococcin (1) will be performed, which will be helpful in laying a foundation for overcoming the drug resistance that pathogens quickly develop against tinea pedis.

In this paper, the antifungal secondary metabolite compound Griseococcin (1) from *B. radicata* were studied. The compound from *Bovistella radicata* (*Mont.*) *Pat* was purified. Molecular weight and molecular formula of the purified compound (Griseococcin (1)) were 661 Da and $C_{37}H_{43}NO_{10}$ respectively, it can restrain main pathogens (*T. rubrum* and *T. mentagrophytes*) leading to tinea pedis. The antifungal activity of Griseococcin (1) was similar to that of the positive control.

Methods

Sample Bovistella radicata (Mont.) Pat collection and tested microorganisms

Bovistella radicata (Mont.) Pat was obtained from Shuinan town, Jishui county, Jiangxi province, China. After strain identification and authenticated by Professor Qingmei Zeng, it belongs to Agaricalesorder, Lycoperdaceae family, Bovistella genus, Bovistella radicata (Mont.) Pat species. The tested pathogenic fungi included Trichophyton rubrum (ATCC 28188), Trichophyton mentagrophytes (ATCC 9533), Epidermophyton floccosum (ATCC 52066), and Candida albicans (ATCC 10231). Four strains of test pathogenic bacteria included Staphylococcus aureus (ATCC 6538), Bacillus subtilis (ATCC 6051), Escherichia coli (ATCC 8739) and Pseudomonas aeruginosa (ATCC 27582). All standard bacterial and fungal strains were obtained from rom the Microbiology Laboratory at Department of Biology, Anhui Medical University, Anhui.

Fermentation, extraction and purification of *Bovistella* radicata (Mont.) Pat

The mature *B.radicata* should be dried at 40 °C for at least 2 days, and taken out when its weight is no longer changed, the sporophore and spore powder were ground together and filtered through a 100 mesh sieve. The mixed powder of B.radicata was inoculated into 100 mL of potato dextrose broth (PDB) in 250 ml flask. The flask was kept in rotary shaker at 25 °C with 115 rpm for 72 h. The pH and moisture content of PDB was also determined according Maguireboyle (2014) and Mcauliffe (2016) [30, 31]. For every 12 h, the fermentation was taken to perform antimicrobial activity against main pathogens T. rubrum and T. mentagrophytes by zone of inhibition (ZOI) method. Then the fermentation were centrifuged at 7000 rpm for 20 min and filtered over Whatman No.4 paper to get the final clear supernatant and preserve at 4 °C. 50 ml clear supernatant was purified firstly using 100 ml DEAEcellulose column and eluted by different concentration NaCl (10-30%) to get different fractions. 20% NaCl elution fraction showed best antifungal activity against pathogens and was named as SPAF. Furthermore, SPAF (20% NaCl elution fraction) was purified by sephadex LH-20 column. Different purified fractions (named Griseococcin(s)) were obtained from SPAF, only Griseococcin(1) (500 µg/ml) has antifungal activity and it's biochemical characteristics and spectral (HPLC, FT-IR, 1D and 2D NMR etc.) studies were assessed.

Antimicrobial activity

The examined methods were the minimum inhibitory concentrations (MICs) [22], minimum bactericidal concentration (MBCs) or minimum fungicidal concentration (MFCs) [32, 33] and zone of inhibitions (ZOIs) [34].

Ye et al. BMC Microbiology (2020) 20:276 Page 7 of 9

ZOI is qualitative analysis and MIC is quantitative analysis of antimicrobial activity [35]. The MICs, MBCs and MFCs of Griseococcin (1) were determined in the 96well plates by the double micro dilution method against pathogens. 100 µL dilutions (approximately 10⁶ CFU/ mL) of T. rubrum, T. mentagrophytes, E. floccosum, and C. albicans in potato dextrose broth and S. aureus, B. subtilis, E. coli and P. aeruginosa in Nutrient Broth [36] were inoculated into 96-well plates. Then, 100 µL Griseococcin (1) solutions were added after a double dilution with the corresponding medium broth (from $500 \,\mu g/mL$ to $7.8 \,\mu g/mL$). 0.9% (v/v) NaCl was used as the negative control. Gentamicin sulfate or Terbinafine were dissolved in normal saline (NS) to a concentration of 1 mg/mL for the subsequent tests as positive control against bacteria or fungi. The Petri dishes were incubated at 37 °C for 24 h with S. aureus, E. coli, B. subtilis and P. aeruginosa, for 48 h with T. rubrum, T. mentagrophytes, E. floccosum, and C. albicans. Griseococcin (1) was also dissolved in NS at 1 mg/mL. The MIC was recorded as the lowest concentration of sample showing no detectable growth. MFC or MBC was determined as the concentration causing no visible growth and killing 99.5% of the original inoculum. Ten microliters of subinhibitory concentrations of Griseococcin (1) was placed in the corresponding solid medium for 48 h to determine the MBC or MFC values according to the growth of the microbial colonies. Each sample was performed twice. The zones of inhibition (ZOI) of Griseococcin (1) (100 µg/ml) was also evaluated. The prepared Griseococcin (1) was filled into the wells. After incubating for 24 h at 37 °C, the measurements were done basically from the edge of the zone to the edge of the well [34].

General experimental procedures

The UV_{max} absorption spectrum of SPAF was analyzed at full-wave spectra (200-900 nm) by UV/vis 2802 spectrophotometer. The FT-IR spectrum of Griseococcin(s) were recorded on a Thermo Nicolet Spectrum FT-IR in a range of $4000-400\,\mathrm{cm}^{-1}$ with KBr pellets. HR-ESI-MS data were obtained on an Agilent 1260 Infinity LC coupled to a 6230 TOF. 20 mg of the dried sample was dissolved in 0.55 mL of deuteroxide (99.99% D) in a NMR tube. 1D and 2D NMR spectra were acquired on an AVANCE-600 NMR spectrometer (Bruker Inc., Rheinstetten, Germany) at 50 °C. The chemical shifts were given in δ (ppm) and referenced to the solvent signal (D₂O-d₆, δ H 2.50, δ C 39.5). Column chromatography (CC) was conducted on DEAE-cellulose and Sephadex LH-20. The fractions Griseococcin(s) were also monitored by HPLC (Agilent 1260 chromatography system, USA) which was equipped with a diode array detector (DAD). The DAD detector was set at 215 nm to acquire chromatograms. The separation of

Table 4 ¹H (700 MHz) and ¹³CNMR (175 MHz) spectroscopic data for Griseococcin (1): in DMSO

Position	δ H (mult, J inHz)	δC
1	7.81	166
2		128.56
3		175.11
4		138.60
5	7.93	135.7
6	7.72	137.21
7		136.28
8	7.80	135.45
9		138.56
10		175.07
11		128.57
12	7.66	166.07
13	1.07	30.18
14	3.62	166.01
15	3.49	166.02
16	3.55	166.03
17	3.51	166.01
18	1.25	23.15
19	1.94	23.16
20	2.09	184.15
1'	1.94	28.4
2 '		215.7
3 '	2.14	23.4
4 '	4.47	73.6
4 ' -OH	8.37	
5 '	2.12	29.05
6 '		215.7
7 '	1.08	25.52
8 '	1.94	26.81
9 '	3.62	71.25
9 ' -OH	7.81	
10 '	1.29	23.45
11 '	3.49	71.15
11 ' -OH	7.80	
12 '	1.73	23.47
13 '	3.51	71.18
13 ' -OH	7.93	
14 '	1.85	20.27 CH ₃
15 '	2.11	20.27 CH ₃
16 '	1.06	20.27 CH ₃
17 '	1.07	20.27 CH ₃

Ye et al. BMC Microbiology (2020) 20:276 Page 8 of 9

compound was performed on a Hypersil RP-C18 column (5 μ m, 250×10.0 mm, Thermo Fisher Scientific, USA) at a temperature of 25 °C. Injection volume: 20 μ L.

Griseococcin (1)

Griseococcin (1): IR (neat) v max 3417, 2926, 2356, 1637, 1618, 1456, 1414, 866, 624 cm $^{-1}$; UV (D₂O) λ max 215 nm; 1 H and 13 CNMR data see Table 4; HR-ESI-MS m/z 661.1970 [M + H] + (calcd for C₃₇ H₄₃NO₁₀, 661.1968).

Supplementary information

Supplementary information accompanies this paper at https://doi.org/10. 1186/s12866-020-01961-x.

Additional file 1.

Abbreviations

NMR: Nuclear Magnetic Resonance; HPLC: High Performance Liquid Chromatography; FT-IR: Fourier transform infrared spectroscopy; DMSO: Dimethyl Sulfoxide; ZOI: Zone of inhibition; MIC: Minimum inhibitory concentration; HR-ESI-MS: High-resolution electrospray ionization mass spectroscopy; DAD: Diode array detector; MW: Molecular weight

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Not applicable.

Authors' contributions

YY conceived and designed the experiments, performed the experiments and analyzed the data, wrote and edited the manuscript. QZ1 analyzed the data. The excellent technical support by QZ2. All authors read and approved the final manuscript

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Availability of data and materials

The datasets used and analysed during the current study are available from the corresponding author on resonable request.

Ethics approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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Ye et al. BMC Microbiology (2020) 20:276 Page 9 of 9

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