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*CORRESPONDENCE Yinglao Zhang ⊠ zhangyl@ahau.edu.cn

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Pigments of aminophenoxazinones and viridomycins produced by termite-associated *Streptomyces tanashiensis* BYF-112

Shuxiang Zhang¹, Jun Wu¹, Zhou Jiang¹, Le Zhang¹, Tao Song¹, Xinhua Liu², Caiping Yin¹ and Yinglao Zhang^{1*}

¹School of Life Sciences, Anhui Agricultural University, Hefei, China, ²School of Pharmacy, Anhui Medical University, Hefei, China

Termite-associated Streptomyces tanashiensis BYF-112 was found as a potential source for yellow and green pigments, which were stable under the tested temperature, light and metal ions. Eight metabolites (1-8), including four new natural yellow pigments aminophenoxazinones (1-4), and two rarely iron dependent green pigments viridomycin A and F (9-10) were isolated from BYF-112 cultured in YMS and YMS treated with FeSO₄, respectively. The metabolites 2-4 displayed a significant safety performance on the normal liver cell line L-02, while the metabolite 1 showed weak cytotoxicity against the L-02 and several cancer cells. Especially, in the filter paper disc tests, the compound 1 possessed strong antibacterial activity against methicillin-resistant Staphylococcus aureus (MRSA) with the zone of inhibition (ZOI) of 15.3mm, which was equal to that of referenced levofloxacin (ZOI=15.2mm). And the metabolite 1 also showed moderate antibacterial activities against Micrococcus teragenus and S. aureus, with the ZOI values of 15.3 and 17.2mm. In addition, by the minimum inhibitory concentration (MIC) assay, the compound 1 displayed potential antibacterial activities against M. teragenus, S. aureus and MRSA, with the MIC values of 12.5, 12.5, and 25.0µg/ml, respectively. The present results indicate that BYF-112 may be a promising source for safe and bioactive pigments, which can be used for further development and industrial applications.

KEYWORDS

Streptomyces tanashiensis BYF-112, natural pigments, aminophenoxazinones, viridomycins, cytotoxic and antibacterial activities

1. Introduction

Edible pigments play a very important role in food qualities and consumer acceptance of products. With the increasing demand for pigments production in the world, the market of food colorants was estimated to reach USD 5.12 billion by 2023 with a five-year compound growth rate of 5.7% (Rana et al., 2021). Usually, food colors are categorized into

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natural and synthetic colors based on the origin (Teixeira et al., 2022). Synthetic colors have been widely used in the food industry for many years, but they pose serious health risks (Saini et al., 2018). Consumers are demanding natural pigments derived from animals, plants, insects and microbes for their safe, non-toxic, non-carcinogenic and biodegradable characteristic (Hosseini et al., 2021). Among these sources, natural edible colors by microbial fermentation have gained significant attention due to the abundance of raw materials without dependence on seasonal variations, cost effective production, and higher concentration with easier purification of products (Panesar et al., 2015; Pombeiro-Sponchiado et al., 2017). To satisfy these requirements, investigations on natural pigments, especially those derived from microorganisms, have increased (Chatragadda et al., 2019a; Chatragadda and Dufosse, 2021; Ramesh et al., 2022).

And several kinds of pigments derived from microorganisms have been developed as industry products, including prodigiosin (Ramesh et al., 2020), phycocyanin (Ashaolu et al., 2021), ankaflavin (Hsu et al., 2011; Dufossé, 2017), astaxanthin (Patel et al., 2022), β -carotene (Lopes et al., 2009; Dufossé, 2017), monascorubramin (Dufossé, 2018), riboflavin (De Faria et al., 2022), rubropunctatin (Zheng et al., 2010), anthraquinones (Venil and Lakshmanaperumalsamy, 2009). These food grade pigments obtained from microbial sources can also have health benefits as anti-cancer, anti-microbial and antioxidant, etc. (Rao et al., 2017).

Aminophenoxazinones are highly colored dyes, possessed a number of promising properties like anticarcinogenic, antifungal, antiparasitic, antibacterial or antimicrobial activities (Zorrilla et al., 2021) which consist of tricyclic structures with double bonds in aromatic systems containing oxygen and nitrogen atom (Pasceri et al., 2013). Thus, aminophenoxazinones are good candidate food grade pigments. Here we obtained four new natural aminophenoxazinones pigments 1-4, two rarely green pigments (9-10) from Streptomyces tanashiensis BYF-112 associated with Odontotermes formosanus for the first time. In this context, the objective of the work was to evaluate the potential of the production of natural pigments by insect associated actinomycete isolated from O. formosanus, that is (i) characterize the pigment by UV-Vis spectroscopy and evaluate its stability in relation to temperature, light and metal ions, (ii) separate and identify the pigments by column chromatography, electrospray ionization mass spectroscopy and NMR; and (iii) evaluate the cytotoxic and antimicrobial activities of the pigments.

2. Materials and methods

2.1. Influence of iron on the metabolite of BYF-112

Streptomyces tanashiensis BYF-112 was isolated from the workers of *O. formosanus* as described in our previous work and was preserved on YMS (1.0 g L^{-1} KNO3, 1.5 g L^{-1} yeast extract, 10 g L^{-1} soluble starch, and 20 g^{-1} agar) slants at 4°C until use

(Long et al., 2022). From a 7-day old YMS solid culture plate, a conidial spore suspension was prepared by adding 15 ml of distilled water containing 0.2% Tween 80 (w/v) under agitation by glass rod. To study the effect of iron, aliquots (2 ml) of spore suspension were added into 100 ml of YMS production media supplemented with deionized water dissolved FeSO₄•7H₂O at the concentrations of 50 mg/l. Equal amount of deionized water was added to the control group. The filter of each fermentation broth was extracted three times with equal amount of ethyl acetate (EtOAc) after fermenting at 28°C in a shaker rotating at 180 rpm for 7 days. The EtOAc extracts were fast analyzed by thin-layer chromatography (TLC).

2.2. Pigments stability

Solution ($250 \mu g/ml$) of yellow pigments (YP) and the green pigments (GP) were dissolved in DMSO and deionized water respectively, and stored in the dark. The pigments stability under heat, light and metal ions were performed by the methods described in detail previously with slight modification (Zhu et al., 2020; Wu et al., 2022). Heat stability: the tested tubes were placed into 101-3A electric thermostatic drying oven (Changzhou, Jiangsu, China) at 40, 60, 80, 100, and 120°C for 2 h. The samples were then removed and quickly cooled to room temperature in an ice bath for further analysis. Stability under condition with light and dark: the effect of light on pigment stability was determined by after exposure to indoor incandescent light and stored at dark place for 1–7 days. Stability under condition with metal ions: the effect of metal ions (Ca²⁺, Zn²⁺, Mg²⁺, Na⁺, and K⁺, 1 mmol/l) on the pigment was determined in the dark for 2 days.

The retention rate of YP and GY were calculated at the maximum absorbance 407 and 700 nm respectively, using UV–Visible spectrophotometer (UNICO UV-2800) according to following formula (Wu et al., 2022).

Retention rate of YP/GP = $A1/A0 \times 100\%$. A1/A0: the absorbance of YP/GP after/before treatment.

2.3. Fermentation

The fermentation procedure was described as the previously reference with slight modifications (Zhang et al., 2020). The fresh spores of *S. tanashiensis* BYF-112 cultivated on YMS agar plates at 28°C for 3–5 days were inoculated into 250 ml Erlenmeyer flask containing 100 ml YMS liquid culture medium (10g soluble starch, 1.5g yeast extract, 1.0g KNO₃, 1,000 ml deionized water). After 3 days of incubation at 28°C on rotary shakers at 180 rpm, 30 ml of cultured liquid was transferred as a seed into Erlenmeyer flasks (1,000 ml), each containing 400 ml YMS broth (or YMS broth supplemented with FeSO₄•7H₂O). The cultures were continuously shaken at 180 rpm for 7 days at 28°C to produce pigments.

2.4. Extraction, isolation, and identification of metabolites

The yellow culture broth (161) of BYF-112 grown on a YMS liquid medium was filtered by gauze to obtain the supernatant, which was then extracted with ethyl acetate (EtOAc, 3×161) at room temperature. The EtOAc phase was evaporated in vacuo to afford crude extract (6.6 g). The crude extract was subjected to a silica gel column eluting with a stepwise gradient of CH₂Cl₂/ MeOH (100, 0–100:16, ν/ν) to give five fractions (Fr1–Fr5). Compound 6 (15 mg) was crystallized from the CH₂Cl₂ solution from the Fr2 (CH₂Cl₂/MeOH, 100:1, ν/ν). Fr2 was repeatedly chromatographed over a Sephadex LH-20 column (MeOH) to yield metabolite 5 (6.0 mg). Fr3 (CH₂Cl₂/MeOH, 100:2, v/v) was further fractionated on a silica gel column, eluting with (CH₂Cl₂/MeOH, 100:0, 100:1 and 100:2, v/v) to give three subfractions (R1-R3). Subfractions R2 and R3 were loaded onto a Sephadex LH-20 column eluting with MeOH to give compounds 1 (40.2 mg) and 3 (5.2 mg), respectively. Compound 2 (30.1 mg) was crystallized from the methanoldichloromethane binary solvent solution of Fr4 (CH₂Cl₂/ MeOH, 100:4). The remaining part of Fr4 was further purified by a Sephadex LH-20 eluting with MeOH to yield compounds 7 (8.8 mg) and 8 (7.9 mg). Fr5 (CH₂Cl₂/MeOH, 100:16, v/v) was also loaded onto a Sephadex LH-20 column to yield compound 4 (8.6 mg).

The green culture broth (81) of BYF112 grown on a YMS liquid media supplemented with $FeSO_4 \cdot 7H_2O$ was centrifuged to afford the liquid phase at 6,000 rpm for 10 min. The liquid phase was passed through a macroporous resin column D-101 (Shangxi Lebobiochem Co., Ltd., China), rinsed with deionized water until the color of the effluent turned colorless, and then eluted with methanol. The organic fraction was evaporated *in vacuo* to afford crude extract (2.6 g). The crude extract was subjected to a silica gel column eluting with a stepwise gradient of $CH_2Cl_2/MeOH$ (100:4, 100:8, and 100:16, v/v) to give two fractions (Fr1–Fr2). Fractions Fr1 and Fr2 were further purified by a Sephadex LH-20 eluting with MeOH to yield compounds **9** (20.1 mg) and **10** (10.9 mg), respectively.

Structural identifications of the metabolites were made on the basis of the spectroscopic analysis. NMR spectra were recorded on an Agilent II DD2 instrument operating at 600 MHz for ¹H and 150 MHz for ¹³C, while 2D spectra (COSY, HMQC, HMBC, and DEPT) were obtained using standard Agilent software. Chemical shifts were given in parts per million (δ) downfield from the TMS internal standard. A Mariner Mass 5,304 instrument was used to measure the HR-ESI-MS spectra.

2.5. Cytotoxicity assay

The metabolites **1–4** were evaluated for their cytotoxic activities against human malignant melanoma cell line (A375), human ovarian cancer cell line (SKOV-3), human breast cancer

cell line (MDA-MB231), human gastric cancer cell line (MGC-803) and human normal cell line (L-02) with the MTT method (Wang et al., 2022). The cell lines were maintained in Dulbecco's modified eagle medium (DMEM) supplemented with 10% fetal bovine serum (FBS) along with 1% penicillin and streptomycin. Cells were grown at 37°C in an atmosphere with 5% CO₂ and then seeded in 96-well plates and permitted to grow for 20 h. Then tested compounds at preset concentrations were added. The 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT; $20\,\mu l,\,5\,mg/ml)$ was added into each well after a $48\,h$ exposure period. After an additional 4h further incubation, the medium was replaced by DMSO (150 µl), then shaking the plates for 15 min. The absorbance at 490 nm were recorded. The percentage of cell viability was calculated using the following formula: Cell viability (%) = the OD of tested group/the OD of CK \times 100%. Finally, IC₅₀ values were determined by EXCEL. The azithromycin was used as positive control.

2.6. Antimicrobial activity

The Staphylococcus aureus (ATCC6538), methicillin-resistant S. aureus (MRSA, SA2), Escherichia coli (ATCC8739) and Micrococcus tetragenus (ATCC35098) donated by Professor Ting Xue from Anhui Agricultural university were used as a target pathogen for the examination of antibacterial activity of the metabolites. The disc diffusion (Özogul et al., 2022) and minimum inhibitory concentrations (MICs; Cui et al., 2022) methods were performed to evaluate the antibacterial activities of the new natural metabolites 1, 2, and 4. Filter paper disks (6mm) with metabolite dissolved in DMSO in a concentration (30 µg/filter paper) were added to the culture medium, and the plates were incubated at 37°C for 12h. Filter papers with DMSO and levofloxacin were set as negative and positive controls, respectively. All experiments were performed in triplicate, and data were shown as mean values standard deviation. And the MICs of the compounds were determined by the method as we previously described (Cui et al., 2022).

3. Results and discussion

3.1. Effect of iron on the metabolic profile of *Streptomyces tanashiensis* BYF-112

Effect of Iron on the metabolic profile of *S. tanashiensis* BYF-112 was displayed in Supplementary Figure S1A. The result showed that the color of the fermentation broth exhibited a great difference between YMS and YMS culture broth treated with iron, which presented yellow and green, respectively. Furthermore, TLC profiles indicated that two green spots were observed in iron treated extract, and only yellow spots were seen in the extracts of iron free cultures (Supplementary Figure S1B). These results

showed that iron had a great influence on the metabolite profiles of the strain BYF-112.

3.2. Stability of yellow and green pigments

Previous reports showed that the color stability of natural pigment was affected by external factors such as heat, light, and metal ions (Zhang et al., 2006; Zhu et al., 2020). Therefore, the stability of YP and GP under different environmental conditions (heat, light, and metal ions) were evaluated in this study.

As shown in Figures 1A,B, both YP and GP presented strong heat stability with the residual rate of 89.44-98.22% under exposure to $40-100^{\circ}$ C for 2 h at the concentration of $250 \,\mu$ g/ml. The result indicated that microbial pigments might be applied in the food industry due to their superior thermal stability, compared to other natural pigments (Vendruscolo et al., 2013; Teran Hilares et al., 2018).

The color stability of YP and GP stored in light and dark was conducted under being treated for 1-7 days at 25°C (Figures 1C,D). The results showed that the YP retention rate

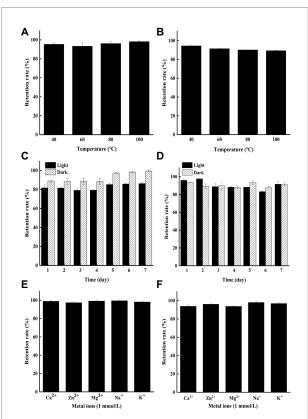


FIGURE 1

The stability of the yellow pigments (YP) and green pigments (GP) in different tested conditions. (A) Effects of heat on the stability of the YP. (B) Effects of heat on the stability of the GP. (C) Effects of light on the stability of the YP. (D) Effects of light on the stability of the GP. (E) Effects of metal ions on the stability of the YP. (F) Effects of metal ions on the stability of the GP.

(86.23%) in the indoor incandescent light decreased slightly, compared to that of in the dark for 7 days (99.44%), while the GP remained stable under light and dark for 7d with the retention rate of 91.57 and 91.42%, respectively. Hence, both pigments can be preserved in light to some extent.

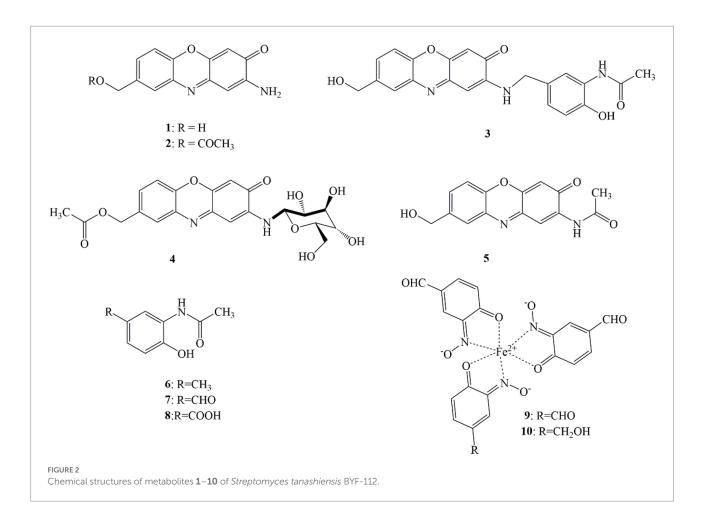
The effect of metal ions $(Ca^{2+}, Zn^{2+}, Mg^{2+}, Na^+, K^+, 1 \text{ mmol/l})$ on the pigment was determined in the dark for 2 days. The results showed that both YP (Figure 1E) and GP (Figure 1F) were stable with the retention rate of more than 92% under the treatments of different metal ions.

3.3. Identification of the secondary metabolites from *Streptomyces tanashiensis* BYF-112

Four new natural compounds **1–4**, along with six known metabolites **5–10** were isolated from the culture of *S. tanashiensis* BYF-112. Chemical structures of compounds **1–10** are shown in Figure 2.

Compound 1 was obtained as a yellow brown powder, and its molecular formula $C_{13}H_{10}N_2O_3$ was deduced from HR-ESI-MS data (m/z [M+H]⁺ 243.0757, which was quite in accordance with ¹H and ¹³C NMR data (Supplementary Figures S2–S4). ¹H NMR and ¹³C NMR data were as following: ¹H NMR (DMSO- d_6 , 600 MHz) δ_H 4.55 (2H, d, J=5.6 Hz), 5.30 (1H, m), 6.33 (H, s), 6.33 (H, s), 6.74 (2H, brs), 7.38 (1H, d, J=8.3 Hz), 7.43 (1H, d, J=8.3 Hz), 7.60 (1H, brs); ¹³C NMR (DMSO- d_6 , 150 MHz) δ_C 62.5 (CH₂), 98.8 (CH), 103.6 (CH), 115.9 (CH), 125.7 (CH), 127.5 (CH), 133.8 (C), 140.2 (C), 141.0 (C), 147.6 (C), 148.5 (C), 149.2 (C), 180.6 (C). The data were almost no different from 2-amino-8-hydroxymethyl-3H-phenoxazin-3-one, a semisynthetic derivative of the pigment aminophenoxazinone described in the literature (Pasceri et al., 2013). However, to our knowledge, the compound was first discovered as a new natural product.

Compound 2 was purified as a yellow brown powder, and its molecular formula C15H12N2O4 was deduced from HR-ESI-MS ion peak at m/z $[M + H]^+$ 285.0860, which was consistent with ¹H and ¹³C NMR data (Table 1; Supplementary Figures S5–S11). The ¹H NMR and ¹³C NMR spectra of 2 were resembled to those of 1, except an acetyl unit signal ($\delta_{\rm C}$ 170.1, 20.6; $\delta_{\rm H}$ 2.09, s, 3H) at position 12 in 1 appeared to be COCH₃, which corresponded to the increase in molecular of compound 2 by 42 amu compared to 1. Further structure elucidation of 2 was established by the 1D and 2D NMR spectra. The ¹H NMR data exhibited the protons resonances of a trisubstituted aromatic ring proton signals at $\delta_{\rm H}$ 7.49, (1H, d, J=8.4Hz, H-6), $\delta_{\rm H}$ 7.44, (1H, d, J=9.5Hz, H-7) and $\delta_{\rm H}$ 7.69, (1H, s H-9), another two aromatic protons at $\delta_{\rm H}$ 6.34 (1H, s, H-1), 6.36 (1H, s, H-4), one oxygenated methylene proton peak at $\delta_{\rm H}$ 5.16 (2H, s, H-11), one methyl protons of acetyl group resonances line at $\delta_{\rm H}$ (2.09, 3H, s, H-14) and one α -amino proton signal at $\delta_{\rm H}$ 6.81 (2H, d, J = 6.7 Hz, NH₂-2). The observation agreed with its ¹³C NMR spectra, which displayed as anticipated one methyl carbon ($\delta_{\rm C}$ 20.6, CH₃, C-14), one methylene carbon ($\delta_{\rm C}$



64.6, CH₂, C-11), one carbonyl carbon ($\delta_{\rm C}$ 180.1, CO, C-2), and 11 aromatic carbon resonance lines ($\delta_{\rm C}$ 98.4, 142.4, 103.4, 148.8, 141.4, 115.9, 128.3, 133.3, 127.2, 133.4, and 148.4). These substructures were pieced together by the 2D NMR experiments including its ¹H–¹H COSY spectrum highlighting the two coupling sequences (11-H to 7-H and 9-H), and its HMBC spectrum (Supplementary Figure S12) indicating the key correlations of H-1 to C-3 and C-4a, and of H-4 to C-2 and C-10a, and of H-6 to C-5a, C-8 and C-9a, and of H-7 to C-11 and C-5a, and of H-9 to C-5a and C-11, and of H-8 to C-8 and C-9, and of 2-NH₂ to C-1 and C-3. Thus, the structure of metabolite **2** was determined as a new aminophenoxazinone derivative and named exfoliazone B.

Compound **3** was purified as a red yellow powder, and its molecular formula was evidenced to be $C_{22}H_{19}N_3O_5$ from the quasimolecular ion peak at m/z [M+Na]⁺ 428.1217 in its HR-ESI-MS spectrum, which was consistent with ¹H and ¹³C NMR data (Table 1; Supplementary Figures S13–S19). The ¹H NMR (DMSO- d_6 , 600 MHz) spectrum of **3** exhibited five aromatic protons signals (δ_H 6.04, 6.36, 7.45, 7.38 and 7.58) and one imino proton resonance (δ_H 7.60) and one oxygenated methylene signal (δ_H 4.54 and 5.33), and the ¹³C NMR (DMSO- d_6 , 125 MHz) spectrum displayed 11 aromatic carbon resonance lines (δ_C 96.3, 145.7, 103.3, 149.3, 140.7, 115.6, 127.2, 140.0, 125.39, 133.5, 149.9),

one carbonyl carbon signal ($\delta_{\rm C}$ 179.9) and one methylene carbon ($\delta_{\rm C}$ 63.2). These data showed almost no difference with exfoliazone described in the literature (Zhang et al., 2015). Apart from these signals of exfoliazone skeleton, there were six aromatic carbon ($\delta_{\rm C}$ 128.5, 121.1, 126.5, 147.9, 115.9, and 123.5), two aromatic proton ($\delta_{\rm H}$ 7.68 and 6.80), one carbonyl carbon ($\delta_{\rm C}$ 169.1), one methyl carbon ($\delta_{\rm C}$ 23.6), one methyl proton ($\delta_{\rm H}$ 2.05), one methylene carbon ($\delta_{\rm C}$ 45.2) one methylene proton ($\delta_{\rm H}$ 4.33), one imino proton ($\delta_{\rm H}$ 7.68) and one hydroxyl proton ($\delta_{\rm H}$ 9.73) as shown in the Table 1. These unassigned data indicated that a subgroup N-(2-hydroxy-5methyl-phenyl)-acetamide were existed in **3**, which corresponded to be increase in molecular weight of **3** by 164 amu compared to that of exfoliazone. These hypotheses were verified by the key HMBC (Fig. S12) correlations in compound **3**. Thus, the metabolite **3** was identified as a new natural product, named exfoliazone C.

Compound 4 was obtained as a yellow red powder, and its molecular formula $C_{21}H_{22}N_2O_9$ was deduced from HR-ESI-MS ($m/z [M+H]^+ 447.1378, [M+Na]^+469.1205$ calcd for $C_{21}H_{23}N_2O_9$, 447.1404 and $C_{21}H_{22}N_2O_9Na$, 469.1223), which was consistent with the ¹H and ¹³C NMR data (Table 2; Supplementary Figures S20–S26). The ¹H NMR (DMSO- d_6 , 600 MHz) spectrum of 4 exhibited five aromatic protons signals $\delta_H 6.45$ (1H, s, H-1), 6.44 (1H, s, H-4), 7.54 (1H, d, J=8.4Hz, H-6), 7.52 (1H, d, J=9.5Hz, H-7) and 7.76 (1H, s, H-9; one methylene proton δ_H 5.18 (2H, s, H-11); one

No.	2		3		
	$\delta_{ extsf{H}}$, mult (J in Hz)	δ_{C}	$\delta_{ extsf{H}}$, mult (J in Hz)	δ_{C}	
1	6.36, s	98.4, CH	6.04, s	96.3, CH	
2		142.4, C		145.7, C	
3		180.1, C		179.9, C	
4	6.36, s	103.4, CH	6.36, s	103.3, CH	
4a		148.8, C		149.3, C	
5a		141.4, C		140.7, C	
6	7.49, d, 8.0	115.9, CH	7.45, d, 8.0	115.6, CH	
7	7.44, d, 9.5	128.3, CH	7.38, d, 8.0	127.2, CH	
8		133.3, C		140.0, C	
9	7.69, brs	127.2, CH	7.58, s	125.3, CH	
9a		133.4, C		133.5, C	
10a		148.4, C		149.9, C	
11	5.16, s	64.6, CH ₂	4.54, s	62.2, CH ₂	
13		170.1, C			
14	2.09, s	20.6, CH ₃			
1'				128.5, C	
2'			7.68, s	121.1, CH	
3'				126.5, C	
4'				147.9, C	
5'			6.80, d, 7.9	115.9, CH	
6'			6.95, d 7.9	123.7, CH	
8'				169.1, C	
9'			2.05, s	23.6, CH ₃	
10'			4.33, d, 6.18	45.2, CH2	
2-NH ₂ /NH	6.81, brs, NH ₂		7.60, d, 6.1, NH		
11-OH			5.33, br, s		
4'-OH			9.73, br, s		
7'-NH			9.31, s		

TABLE 1 ¹H and ¹³C NMR data for compounds 2 and 3 in DMSO-d₆.

methyl proton $\delta_{\rm H}$ 2.08 (3H, s, H-14) and one imino proton $\delta_{\rm H}$ 6.96 (1H, d, J = 6.7 Hz, NH-8'); and the ¹³C NMR (DMSO- d_6 , 125 MHz) spectrum displayed 11 aromatic carbon resonance lines δ_C 99.6 (CH, C-1), 145.1 (C, C-2), 103.4 (CH, C-4), 148.9 (C, C-4a), 141.7 (C, C-5a), 116.0 (CH, C-6), 129.3 (CH, C-7), 133.5 (C, C-8), 127.5 (CH, C-9), 133.3 (C, C-9a), 148.4(C, C-10a), one carbonyl carbon signal 179.6 (CO, C-2), one methylene carbon signal 64.6(CH₂, C-11) and one methyl carbon signal 20.6 (CH₃, C-14). These proton and carbon signals of substructures were very similar to that of metabolite 1 suggesting that 4 might be a derivative of 1.The residual proton signals $\delta_{\rm H}$ 4.49 (1H, m, H-1'), 3.37 (1H, m, H-3'), 3.15 (1H, m, H-4'), 4.98 (1H, d, J=5.2Hz, OH-4'), 3.30 (1H, m, H-5'), 5.06 (1H, d, J=4.3 Hz, OH-5'), 3.34 (1H, m, H-6'), 5.23 (1H, d, J=5.3 Hz, OH-6'), 3.70 (1H, m, H-7), 3.46 (1H, m, H-7'), 4.50

TABLE 2 ¹ H and ¹³ C NMR	data for compound	4 in	DMSO-d ₆
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No.	$\delta_{ extsf{H}}$, mult (J in Hz)	δ_{C} ,
1	6.45, s	99.6, CH
2		145.1, C
3		179.6, C
4	6.44, s	103.4, CH
4a		148.9, C
5a		141.7, C
6	7.54, d, 8.4	116.1, CH
7	7.52, d, 9.5	129.3, CH
8		133.5, C
9	7.76, s	127.5, CH
9a		133.3, C
10a		148.4, C
11	5.18, s	64.6, CH ₂
13		170.2, C
14	2.09, s	20.6, CH3
1'	4.49, m	83.3, CH
3'	3.37, m	77.0, CH
4'	3.15, m	70.0, CH
5'	3.30, m	78.1, CH
6'	3.34, m	72.2, CH
7'	3.70, m	60.8, CH
	3.46, m	
8'	6.96, d, 6.7	
4'-OH	4.98, d, 5.2	
5'-OH	5.06, d, 4.3	
6'-ОН	5.23, d, 5.3	
7′-OH	4.50, d, 5.1	

(1H, d, J=5.1 Hz, OH-7') and carbon signals $\delta_{\rm C}$ 83.3 (C-1'), 77.0 (C-3'), 70.0 (C-4'), 78.1 (C-5'), 72.2 (C-6'), 60.8 (C-7') indicated that a hexose were existed in **4**. These substructures were pieced together by the 2D NMR experiments including its ¹H-¹H COSY spectrum highlighting the two coupling sequences (11-H to 7-H and 9-H), and its HMBC spectrum (Supplementary Figure S12) indicating the key correlations of H-1 to C-2 and C-3, and H-4 to C-3, C-2 and C-10a, and H-6 to C-5a and C-8, and H-7 to C-11 and C-5a, and H-9 to C-11 and C-5a, and H-11 to C-7 and C-9, and H-14 to C-1, and H-8' to C-1, C-3, C-1' and C-6'. Thus, the structure of metabolite **3** was determined as a new glycosylated derivative of **1** and named exfoliazoneglycoside.

The known compounds **5–10** were identified as exfoliazone (5) (Zhang et al., 2015), N-(2-Hydroxy-5-methylphenyl acetamide (6) (Suzuki et al., 2006a), 3-acetylamino-4-hydroxybenzaldehyde (7) (Suzuki et al., 2006b), 3-acetylamino-4-hydroxybenzonic acid

(8) (He et al., 2018), viridomycin A (9) and viridomycin F (10) (Omura et al., 1999), by comparing the NMR and MS data with those reported in the literature. It was reported that the known compounds N-(2-Hydroxy-5-methylphenyl) acetamide (6), 3s-acetylamino-4-hydroxybenzaldehyde (7) and 3-acetylamino-4-hydroxybenzoic acid (3,4-AHBA; Gould et al., 1996). The remaining aminophenoxazinones were presumably synthesized from the precursor 3,4-AHBA in the gene cluster, which contained a key tyrosinase-like copper-containing monooxygenase responsible for the C-nitrosation (Noguchi et al., 2010). *Streptomyces* has been reported as pigments producer (Chatragadda et al., 2019b; Zhu et al., 2020). Our results enriched the pigment molecules diversity produced by the *Streptomyces*.

3.4. Cytotoxicity of the new metabolites 1–4

In vitro toxicity of the compounds 1–4 was evaluated against the cell lines using the MTT method (Table 3). The results showed that the metabolites 2–4 showed no cytotoxic activities against the human normal liver cell line L-02 with the IC₅₀ values of more than 100 μ M. These safety performance of 2–4 might be an advantage for applying in the food industry. Furthermore, the metabolite 2 showed selective cytotoxic activities toward human ovarian cancer cell line SKOV-3 and gastric cancer cell line MGC-803 with the IC₅₀ values of 75.78 and 69.88 μ M, respectively, which were lower than those of positive azithromycin with the IC₅₀ values of 0.41 and 1.57. The compound 1 exhibited inhibitory activity against the tested human cancer cell lines (human malignant melanoma cell

TABLE 3 Cytotoxic activity (IC $_{50}$ in μM) of the secondary metabolites 1–4.

line A375, SKOV-3, breast cancer cell line MDA-231 and MGC-803) with the IC_{50} values of 22.33–67.70 μ M. However, it is low toxicity against the normal cell L-02 with the IC_{50} value of 72.01 μ M would be a limitation in using as a food pigment.

3.5. Antibacterial activities of the metabolites

The disc diameters of zone of inhibition (ZOI) values of metabolites (1, 2 and 4) against bacterial pathogens were presented in Table 4; Supplementary Figure S27. The results showed that metabolite 1 possessed strong antibacterial activity against methicillin-resistant S. aureus (MRSA) with the ZOI of 15.3 mm, which was equal to that of referenced levofloxacin (ZOI = 15.2 mm). In addition, the metabolite 1 showed moderate antibacterial activities against Gram-positive S. aureus (ZOI=17.2 mm) and *M. tetragenus* (ZOI=15.mm), which were weaker than those of referenced levofloxacin with the ZOI values of ZOI = 28.9 and 24.8 mm, respectively. And in the MICs test (Table 4), the compound 1 exhibited potential antibacterial activities against *M. tetragenus*, *S. aureus* and MRSA in the MIC tests with the MIC values of 12.5, 12.50, and 25.0 µg/ml, which were comparable to those of positive gentamycin sulfate with the MIC values of 3.13, 3.13, and 6.25 µg/ml, respectively. However, both compounds 2 and 4 did not showed antibacterial activities against the test strains. By comparing the structure of compounds 1 and 2, the oxhydryl substituent at C-11 in 2 could reduce antibacterial activities against tested bacteria. It indicated that the 11-position hydroxyl group might play a very important role in inhibiting the growth of the Gram-positive bacteria.

Metabolites	A375	SKOV-3	MDA-231	MGC-803	L-02
1	67.70 ± 4.31	22.33 ± 3.78	55.85 ± 7.47	50.18 ± 7.01	72.01 ± 7.22
2	>100	75.78 ± 5.77	>100	69.88 ± 1.94	>100
3	>100	>100	>100	>100	>100
4	>100	>100	>100	>100	>100
Azithromycin ^a	0.79 ± 0.13	0.41 ± 0.12	4.03 ± 0.43	1.57 ± 0.03	7.34 ± 1.15

^aPositive control.

TABLE 4 Zone of inhibition (ZOI, mm) and minimum inhibitory concentrations (MICs; µg/ml) of the second metabolites 1, 2, and 4 against the tested bacteria.

Metabolites	Escherichia coli		Micrococcus tetragenus		Staphylococcus aureus		MRSA	
	ZOI	MIC	ZOI	MIC	ZOI	MIC	ZOI	MIC
1	_ ^b	>100	15.3 ± 0.2	12.5	17.2 ± 0.2	12.5	15.3 ± 0.1	25.0
2	_ ^b	>100	_ ^b	50.0	_ ^b	>100	_ ^b	>100
4	_ ^b	>100	_ ^b	>100	_ ^b	>100	_ ^b	>100
Levofloxacinª	33.0±1.0	3.13	24.8 ± 0.6	3.13	28.9 ± 0.8	3.13	15.2 ± 0.1	6.25

^aPositive control 30 µg/disc.

^bNo inhibited

4. Conclusion

Here, the metabolites of the strain S. tanashiensis BYF-112 associated with the termite (O. formosanus) were investigated. The results showed that iron elements had a great influence on the color of fermentation broth and metabolites profile of BYF-112. Total of four new natural aminophenoxazinones pigments, along with two rarely green pigments were purified and identified from the fermentation broth of BYF-112 for the first time. Both the yellow and green metabolites showed good stability under heat, light and metal conditions. And the new compounds showed weak or almost no cytotoxicity to the normal cell line L-02. Especially, the metabolite 1 presented strong antibacterial activity against the MRSA, and moderate activities against Gram-positive M. tetragenus and S. aureus. Therefore, these new natural compounds showed stability and safety as food pigments to a certain extent. In summary, the termite-associated S. tanashiensis BYF-112 can be considered a promising new source of bioactive natural pigments that can be used for industrial applications in the future.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary material, further inquiries can be directed to the corresponding author.

Author contributions

YZ designed and supervised the study. SZ, JW, ZJ, LZ, TS, and XL performed the experiments and analyzed the data. SZ wrote

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Conflict of interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Supplementary material

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