

ORIGINAL ARTICLE

Discovery of new hazimycin congeners from *Kitasatospora* sp. P07101



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Abstract In an analytical study of microbial broths, the actinomycete strain *Kitasatospora* sp. P07101 was found to produce three new congeners, which were designated hazimycins B (**1**), C (**2**), and D (**3**), together with the previously reported hazimycin (renamed hazimycin A (**4**)). The structures of these hazimycins were examined using various spectroscopic methods including nuclear magnetic resonance (NMR), and the results revealed that **1–3** were analogues of hazimycin with the replacement of one of the two isonitrile groups in **4** by an NH-formyl group in **1**, the two isonitrile groups and an amide group by two NH-formyl groups and a nitrile group in **2**, and the two isonitrile groups and two amide groups by two NH-formyl groups and two nitrile groups in **3**. Only hazimycin A exhibited moderate antimicrobial activities against Gram-positive bacteria and *Candida albicans*. These results indicated that the presence of two isonitrile groups in the hazimycin structure is essential for antimicrobial activity.

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1. Introduction

Our research group has focused on discovering novel compounds from microbial metabolites^{1–4}. Compounds were screened from our original culture collection using LC–UV and LC–MS/MS instruments. During this chemical screening program, the actinomycete strain *Kitasatospora* sp. P07101 was found to produce unidentified compounds. Novel hazimycins, hazimycins B (1), C (2), and D (3), were recently isolated from the fermentation broth along with the known antibiotic hazimycin⁵ (renamed hazimycin A (4), Fig. 1). These new congeners possessed a diaryl skeleton that contained isonitrile and nitrile groups, which are rare among microbial metabolites. The isolation, structure elucidation, and biological activities of 1–3 have been described in the present study.

2. Results and discussion

2.1. Structure elucidation of 1–3

The physicochemical properties of compounds 1–3 are summarized in Table 1. Compounds 1–3 showed UV absorption between approximately 212 nm and 289 nm, which was identical to that of 4. The IR absorption at 2150–2300 cm⁻¹ suggested the presence of isonitrile and/or nitrile groups in their structures. These results indicated that the basic skeleton of 1–3 was similar to that of 4.

The structure of 1 was elucidated from various spectral data including NMR experiments. The molecular formula of 1 was

determined to be C₂₀H₂₀N₄O₅ based on HR-ESI-MS measurements, which indicated that the molecular formula of 1 has one oxygen atom and two hydrogen atoms more than that of 4. The ¹³C-NMR spectrum showed 20 resolved signals, which were classified into two *sp*³ methylene carbons, two *sp*³ methine carbons, six *sp*² methine carbons, four *sp*² quaternary carbons, two *sp*² quaternary oxycarbons, one *sp* carbon, two *sp*² carbonyl carbons, and one *sp*² formyl carbon by ¹H–¹³C heteronuclear single-quantum correlation (HSQC) analysis. The ¹H NMR spectrum (in DMSO-*d*₆) displayed 18 proton signals. The connectivity of the proton and carbon atoms was established from the ¹H–¹³C HSQC spectrum (Table 2). A comparison of the NMR spectra of 1 and 4 indicated that they both possessed a dihydroxydiaryl skeleton. However, most double signals were observed in the ¹H and ¹³C NMR spectra of 1, suggesting that 1 was a heterodimer, while 4 was a homodimer. A formyl proton signal (δ 7.92) and amide proton signal (δ 8.17) were observed in 1, but were absent in 4, which indicated that one of two isonitrile groups was converted to an NH-formyl group in 1. Cross peaks were observed from H-2'' (δ 4.43) to C-4'' (δ 160.9) as well as from NH-2'' (δ 8.17) to C-4'' in the ¹³C–¹H heteronuclear multiple-bond correlation (HMBC) experiments (Fig. 2A). The structure satisfied the unsaturation number, UV spectra, and molecular formula. These results indicated that compound 1 was a 2''-NH-formyl hazimycin, as shown in Fig. 1.

The molecular formula of 2 was identical to that of 1. However, two proton signals of an NH-formyl group (δ 8.06 and 8.86) were newly observed, and one of the amide proton signals of the two carboxamide

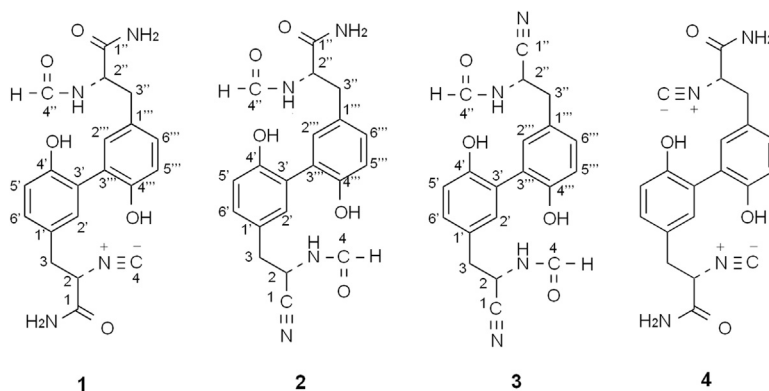


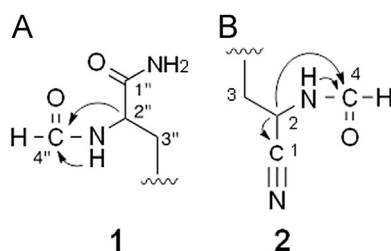
Figure 1 Structures of 1–4.

Table 1 Physicochemical properties of 1–3.

Parameter	1	2	3
Appearance	Pale yellow powder	Colorless oil	Colorless oil
Molecular formula	C ₂₀ H ₂₀ N ₄ O ₅	C ₂₀ H ₂₀ N ₄ O ₅	C ₂₀ H ₁₈ N ₄ O ₄
Molecular weight	396	396	378
HR-ESI-MS (<i>m/z</i>)			
Found	419.1339 [M+Na] ⁺	419.1338 [M+Na] ⁺	401.1227 [M+Na] ⁺
Calcd.	419.1331	419.1331	401.1226
UV (MeOH) λ _{max} (nm)/ε	212/30,254 289/4514	212/38,945 289/4672	214/36,931 289/5065
[α] _D ²⁸ (c=0.1, MeOH)	–1.4°	–1.6°	–10.6°
IR (KBr) ν _{max} (cm ⁻¹)	3400, 3205 3086, 2150 1671, 1621	3315, 3027 2252, 1674 1614, 1504	3379, 3021 2248, 1673 1612, 1500

Table 2 ^1H and ^{13}C NMR chemical shifts of **1–3**.

Position 1		2		Position	3	
δ_{C}	δ_{H}	δ_{C}	δ_{H}		δ_{C}	δ_{H}
1	167.1s –	119.0s –	–	1, 1''	119.0s –	–
1-NH ₂	– 7.48 (1H, s), 7.71 (1H, s)	–	–	1-NH ₂ , 1''-NH ₂	–	–
2	58.9d 4.49 (1H, dd, $J=4.8, 4.4$)	40.1d 4.98 (1H, dd, $J=8.0, 7.6$)	–	2, 2''	40.4d 4.90 (1H, dd, $J=8.0, 7.6$)	–
2-NH	–	– 8.86 (1H, d, $J=7.6$)	–	2-NH, 2''-NH	–	8.86 (1H, d, $J=7.6$)
3	37.8t 2.86 (1H, m), 3.03 (1H, dd, $J=8.8, 4.8$)	36.5t 2.98 (1H, m)	–	3, 3''	36.5t 2.98 (1H, m)	–
4	158.0s –	161.1d 8.06 (1H, s)	–	4, 4''	161.1s 8.06 (1H, s)	–
1'	126.0s –	125.3s –	–	1', 1'''	125.2s –	–
2'	132.2d 7.07 (1H, s)	132.5d 7.07 (1H, s)	–	2', 2'''	132.5d 7.08 (1H, s)	–
3'	125.4s –	126.0s –	–	3', 3'''	125.7s –	–
4'	153.7s –	153.7s –	–	4', 4'''	153.9s –	–
5'	115.7d 6.80 (1H, d, $J=8.0$)	115.8d 6.81 (1H, d, $J=8.8$)	–	5', 5'''	115.8d 6.80 (1H, d, $J=8.4$)	–
6'	128.9d 7.03 (1H, d, $J=8.0$)	128.9d 7.05 (1H, d, $J=8.8$)	–	6', 6'''	129.0d 7.05 (1H, d, $J=8.4$)	–
1''	172.8s –	172.8s –	–			
1''-NH ₂	– 7.02 (1H, s), 7.48 (1H, s)	– 7.04 (1H, s), 7.48 (1H, s)	–			
2''	52.7d 4.43 (1H, ddd, $J=8.4, 4.8, 4.0$)	52.7d 4.44 (1H, ddd, $J=8.4, 4.8, 4.0$)	–			
2''-NH	– 8.17 (1H, d, $J=8.4$)	– 8.16 (1H, d, $J=8.4$)	–			
3''	37.0t 2.65 (1H, m), 2.90 (1H, m)	36.7t 2.60 (1H, m), 2.91 (1H, m)	–			
4''	160.9d 7.92 (1H, s)	160.8d 7.92 (1H, s)	–			
1'''	127.8s –	127.8s –	–			
2'''	132.2d 7.02 (1H, s)	132.2d 7.02 (1H, s)	–			
3'''	126.0s –	125.3s –	–			
4'''	153.1s –	153.0s –	–			
5'''	115.5d 6.74 (1H, d, $J=8.0$)	115.4d 6.75 (1H, d, $J=8.0$)	–			
6'''	128.8d 6.98 (1H, d, $J=8.0$)	128.8d 6.98 (1H, d, $J=8.0$)	–			

**Figure 2** Key HMBCs of **1** and **2**.

groups (δ 7.48 and 7.71) disappeared in the ^1H NMR spectrum of **2**. Furthermore, a new *sp* carbon signal (δ 119.0) was observed in place of one of the two carboxamide carbon signals (δ 167.1) in the ^{13}C NMR spectrum of **2**. These results indicated the formylation of another isonitrile group of **1** and the conversion of one of the two carboxamide groups of **1** to a nitrile group in **2**. The position of the nitrile group was confirmed by ^{13}C - ^1H HMBC experiments (Fig. 2B): cross peaks were observed from H-2 (δ 4.98) to C-1 (δ 119.0) and C-4 (δ 161.1). Thus, compound **2** was elucidated to be 2,2''-NH-formyl and 2-nitrile hazimycin (Fig. 1).

As listed in Table 1, the molecular formula of **3** has one oxygen atom and two hydrogen atoms fewer than that of **2**. Its ^1H -NMR spectrum revealed homodimer-type proton signals, and was almost identical to that of **2** except for the disappearance of the amide proton signals of the carboxamide groups (δ 7.04 and 7.48) in **3**. Furthermore, the presence of a nitrile carbon signal (δ 119.0) was confirmed as well as **2** in the ^{13}C -NMR spectrum, which indicated that another carboxamide group of **2** was converted to a nitrile group in **3**. Finally, cross peaks were observed from H-2'' (δ 4.90) to C1'' (δ 119.0) and C4'' (δ 161.1) as well as from NH-2'' (δ 8.86) to C4'' in

the ^{13}C - ^1H HMBC experiments. Thus, compound **3** was elucidated to be a 2,2''-NH-formyl and 2,2''-nitrile hazimycin (Fig. 1)

Regarding the absolute stereochemistry of the novel hazimycin analogs, dityrosine was prepared by hydrolyzing **4** under acidic conditions because its optical rotation has already been accurately described in previous studies⁶. The results obtained in this study were consistent with those of L,L-dityrosine. Thus, the absolute stereochemistry of **2** and **2''** of **4** was defined as *S*. Similarly, the hazimycin congeners **1–3** are considered to be derived from L,L-dityrosine because they are generated *via* common biosynthetic pathway. Thus, compounds **1–3** should have the same absolute stereochemistry as **4**.

Waltz et al.⁷ previously reported that hazimycin A was biosynthesized from tyrosine and methionine based on ^{14}C -labeling experiments. They postulated that an isonitrile group was formed through the NH-formyl residue generated due to the methylation, reduction, and oxidation of an amino group of tyrosine. In the present study, we demonstrated the presence and structure of not only the NH-formyl product, but also the new family of nitrile-type hazimycins in the culture broth of the actinomycete strain.

2.2. Antimicrobial activities of **1–3**

We examined antimicrobial activity against 7 test microorganisms using the paper disk method⁸. As shown in Table 3, compounds **1–3** and dityrosine did not inhibit the growth of these microorganisms; only compound **4** exhibited antimicrobial activity against *Mycobacterium smegmatis*, *Bacillus subtilis*, *Staphylococcus aureus*, *Micrococcus luteus*, and *Candida albicans* with inhibition zones of 19, 14, 23, 26 and 20 mm, respectively. These results indicated that the

Table 3 Antimicrobial activities of **1–4**^a.

Test organism	Inhibition zone (diameter, mm)				Dityrosine
	1	2	3	4	
Gram-positive bacteria					
<i>Mycobacterium smegmatis</i>	– ^b	–	–	19	–
<i>Bacillus subtilis</i>	–	–	–	14	–
<i>Staphylococcus aureus</i>	–	–	–	23	–
<i>Micrococcus luteus</i>	–	–	–	26	–
Gram-negative bacteria					
<i>Escherichia coli</i>	–	–	–	–	–
<i>Pseudomonas aeruginosa</i>	–	–	–	–	–
Yeast					
<i>Candida albicans</i>	–	–	–	20	–

^a10 µg/6 mm disk.^bCan not detect antimicrobial activity.

attachment of both isonitrile groups in the side chain of the dihydroxydiaryl skeleton was crucial for antimicrobial activity.

3. Material and methods

3.1. General procedures

The actinomycete strain *Kitasatospora* sp. P07101 was originally isolated from a soil sample collected at Minato-ku, Tokyo, Japan. The genus was determined based on taxonomic studies and genetic analysis of 16S rRNA by the identification services of TechnoSuruga Laboratory Co., Ltd., Shizuoka, Japan. This strain was used to produce **1–3**. UV spectra were recorded on a spectrophotometer (8453 model, Agilent). IR spectra were recorded on a Fourier transform infrared spectrometer (FT-710, Horiba). Optical rotations were measured with a digital polarimeter (DIP-370, JASCO). FAB-MS spectra and HR-FAB-MS spectra were recorded on a mass spectrometer (JMS-AX505HA, JEOL). The various NMR spectra were collected with a spectrometer (XL-400, Varian).

3.2. Isolation of **1–3**

The three-day-old fermentation broth (6.5 L) of *Kitasatospora* sp. P07101 was centrifuged to separate the mycelia and supernatant. A part of the supernatant (4.0 L) was then added to a Diaion HP-20 column (volume: 0.2 L, Mitsubishi Chemical Co.) After washing with distilled water (1.0 L), the desired substances were eluted with MeOH (1.0 L) and concentrated *in vacuo* to dryness to produce red brown materials (2.3 g), which were then dissolved in MeOH and purified using high performance liquid chromatography (HPLC, column: DevelosilC30, 250 mm × 20 mm, Nomura Scientific Co.; solvent: a gradient system of 40 min from 5% CH₃CN containing 0.05% trifluoroacetic acid (TFA) to 55% CH₃CN containing 0.05% TFA; detection: UV at 210 nm; flow: 8.0 mL/min). Under these conditions, the peaks eluted at retention

time of 18.8 and 27.8 min were repeatedly collected and concentrated to dryness to give **1** (1.0 mg) and **4** (110 mg) as pale yellow powders, respectively.

A residual supernatant (2.5 L) was extracted twice with an equal volume of ethylacetate, and concentrated *in vacuo* to dryness to produce red brown materials (1.2 g). This sample was dissolved in MeOH, and then purified using HPLC (column: PEGASIL ODS, 250 mm × 20 mm, Senshu Scientific Co.; solvent: a gradient system of 60 min from 10% CH₃CN containing 0.05% TFA to 45% CH₃CN containing 0.05% TFA; detection: UV at 210 nm; flow: 6.0 mL/min). Under these conditions, the peaks eluted at retention time of 27 and 40 min were repeatedly collected and concentrated to dryness to give **2** (44.4 mg) and **3** (55.2 mg) as colorless oils, respectively.

3.3. Preparation of dityrosine

Dityrosine was prepared according to a previous method⁶ by hydrolyzing hazimycin A under acidic conditions. Hazimycin A (20.0 mg) was dissolved in 12 mol/L HCl (1.0 mL), and then hydrolyzed at 60°C for 12 h. The reaction solution was neutralized with 10 mol/L NaOH, and then centrifuged to precipitate the resulting salts. Finally, the collected supernatant was purified using HPLC (column: Develosil C30, 250 mm × 20 mm; solvent: 3.0% CH₃CN; detection: UV at 210 nm; flow: 6.0 mL/min). The peak eluted at 16.7 min was repeatedly collected and concentrated to dryness to give dityrosine (10.2 mg) as a white powder. The spectroscopic data listed below were consistent with the findings of a previous study⁶.

Dityrosine: FAB-MS-positive; [M+H]⁺ = 361, ¹H NMR in DMSO-*d*₆ (600 MHz) δ_H: 2.90 (2H), 3.03 (2H), 3.93 (2H), 6.80 (2H), 7.01 (2H), 7.16 (1H), 7.19 (1H), ¹³C NMR in DMSO-*d*₆ (150 MHz) δ_C: 34.8, 35.4, 54.0, 54.5, 115.8, 116.0, 125.6, 126.0, 126.3, 129.5, 130.1, 133.2, 133.4, 153.2, 153.3, 170.6, 170.8, [α]_D²⁸: –3.54 (c = 0.1, 1 mol/L HCl).

3.4. Assay for antimicrobial activity

Antimicrobial activity against 7 test microorganisms as listed in Table 3 was measured using paper disks (6 mm, ADVANTEC) containing a sample, according to our established method⁸.

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