STRUCTURE OF DUOCARMYCIN SA, A POTENT ANTITUMOR ANTIBIOTIC

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Recently, we reported the structures of the duocarmycins^{1,2)}, a unique antitumor antibiotics constituted with two structural moieties, a substituted pyrrologuinoline or its chemical equivalent cyclopropanopyrroloindole and a trimethoxyindolylcarbonyl group³⁾. In this communication we wish to report the structure of duocarmycin SA (1), a stable compound structurally related to the duocarmycin A (2). Duocarmycin SA (1) produced by an unidentified Streptomyces sp. is a novel and potent antitumor antibiotic, which is effective against murine lymphocytic leukemia P388 and murine sarcoma 180⁴⁾. The structure determination was accomplished by extensive 2D NMR experiments in addition to comparison of its spectra with those of duocarmycin A (2).

Physico-chemical properties were summarized in Table 1. ¹H and ¹³C chemical shifts for 1 and 2 were shown in Tables 2 and 3, respectively. A battery of 2D techniques including COSY, ¹³C-¹H COSY, and correlation spectroscopy *via* long range coupling⁵ (COLOC) experiments were used to determine all proton and carbon line assignments. ¹³C multiplicity data were derived from DEPT experiments. All the proton and carbon chemical shifts were related by ¹³C-¹H COSY experiment. The assignments of chemical shifts of 1 are based on analysis of 2D experiments (described below) and correlation with 2.

HR-MS of 1 (Table 1) exhibited the molecular formula $C_{25}H_{23}N_3O_7$ which was smaller by CH₂O than 2. Comparison of the NMR spectra of 1 and 2, revealed the existence of trimethoxyindolylcarbonyl group in both compounds. It was also confirmed by the mass fragment at m/z 234 (HR-MS, 234.0784; for $C_{12}H_{12}NO_4$, err+1.8 mmu), produced by

Table 1. Physico-chemical properties of duocarmycin SA (1).

Appearance	Yellow powder
Molecular formula	$C_{25}H_{23}N_{3}O_{7}$
EI-MS (m/z)	477 (M) ⁺ , 234
HR-MS (m/z) Calcd:	477.1535
Found:	477.1522
$[\alpha]_{\rm D}^{24}$ (c 0.1, MeOH)	$+180^{\circ}$
UV λ_{\max}^{MeOH} nm (ε)	235 (sh, 21,000), 316 (16,000),
	367 (27,000)
IR (CHCl ₃) ν cm ⁻¹	3460, 1714, 1642, 1619, 1258



Fig. 1. Structures of duocarmycin SA (1) and duocarmycin A (2).

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Notes

Proton	1	2
1-NH	10.25 (br s)	6.36 (s)
2-CH ₃		1.67 (s)
2-COOCH ₃	3.88 (s)	3.74 (s)
3-Н	6.56 (d, $J = 2.2$ Hz)	
4-H,	1.71 (dd, $J = 7.6, 4.7 \mathrm{Hz}$)	2.24 (dd, $J = 7.6$, 4.1 Hz)
4-H _h	1.54 (t, $J = 4.7$ Hz)	1.29 (dd, $J = 4.7$, 4.1 Hz)
4a-H	2.77 (d, t, $J = 7.6$, 4.7 Hz)	3.05 (m)
5-H,	4.46 (dd, $J = 10.6, 4.9$ Hz)	4.45 (dd, J = 10.3, 4.7 Hz)
5-H	4.37 (d, $J = 10.6$ Hz)	4.41 (br d, $J = 10.3$ Hz)
7-H	7.02 (s)	7.17 (s)
1'-NH	9.50 (br s)	9.49 (br s)
3'-H	6.92 (d, $J = 2.2 \mathrm{Hz}$)	6.94 (d, $J = 2.3$ Hz)
4'-H	6.76 (s)	6.78 (s)
5'-OCH3	3.87 (s)	3.88 (s)
6'-OCH3	3.92 (s)	3.93 (s)
7′-OCH3	4.03 (s)	4.06 (s)

Table 2. ¹H NMR data^a of duocarmycin SA (1) and duocarmycin A (2).

^a Measured on Bruker AM400 spectrometers with TMS (0 ppm) as an internal standard in CDCl₃.

Carbon	1	2	Carbon	1	2
C-2	126.9	71.3	2-COOCH ₃	52.1	53.4
C-3	107.5	194.8	C-2'	128.5	128.2
C-3a	130.0	112.0	C-2'α	161.2	161.2
C-3b	31.3	30.6	C-3'	107.8	108.2
C-4	26.1	22.0	C-3'a	123.2	123.3
C-4a	23.6	22.3	C-4'	97.8	97.7
C-5	54.9	55.3	C-5'	150.5	150.6
C-6a	161.6	165.1	C-6'	141.1	141.3
C-7	112.5	113.2	C-7′	138.9	138.9
C-8	177.9	179.7	C-7'a	126.4	126.6
C-8a	131.6	164.4	5'-OCH3	56.3	56.3
2-CH ₃		21.1	6'-OCH3	61.4	61.5
2-COOCH3	161.0	168.0	7'-OCH3	61.1	61.2

Table 3. ¹³C NMR data^a for duocarmycin SA (1) and duocarmycin A (2).

^a Measured on Bruker AM400 spectrometers with TMS (0 ppm) as an internal standard in CDCl₃.

cleavage of amide bond.

In ¹H NMR spectrum of **1** (Table 2), one vinyl doublet (δ 6.56, J=2.2 Hz) coupled with the 1-NH proton (confirmed by ¹H homo decoupling experiment) was observed instead of the 2-methyl singlet observed in the spectrum of **2**. In the comparison of ¹³C NMR spectra of **1** and **2** (Table 3), the 2-methyl (δ 21.1) and 3-keto (δ 194.8) signals of the latter were lacking and trisubstituted double bond carbon signals (δ 126.9 and 107.5) were now present in **1**. These NMR data suggested the difference around the 2, 3 position.

Remaining protons were CH_2 -CH- CH_2 five spin system (position 4, 4a, 5), a methoxy group, and an isolated aromatic proton. ¹³C-¹H coupling constant

of a methylene (δ_c 26.0; $J_{CH} = 166$ Hz) in the five spin system suggests that it consists a part of cyclopropane ring system. Other connectivities of quaternary carbons were established by COLOC experiment (polarization delay is 50 mseconds). The diagnostic results were shown in Fig. 2.

Cross peaks between $\delta_{\rm H} 1.54$ (4-H_b), 7.02 (7-H) and $\delta_{\rm C} 161.6$ (C-6a); $\delta_{\rm H} 4.46$ (5-H_a), 7.02 (7-H) and $\delta_{\rm C} 31.3$ (C-3b); $\delta_{\rm H} 1.71$ (4-H_a) and $\delta_{\rm C} 130.0$ (C-3a); and $\delta_{\rm H} 7.02$ (7-H) and $\delta_{\rm C} 131.6$ (C-8a) indicate the cyclopropanoindole ring. No long range coupling was observed on $\delta_{\rm C} 177.9$ carbonyl carbon in gated decoupling experiment and no cross peak was observed on it in the COLOC experiment, but consideration of chemical shifts indicated the Fig. 2. COLOC experiment of 1.



location of the carbonyl group at 8-position. The methoxy group was shown to connect to carbonyl group, but location of methoxycarbonyl group was not indicated by this experiment. Cross peaks were observed from NH (1-H) proton and vinyl proton (3-H) to all the pyrrole carbons.

The orientation of pyrrole ring and the location of methoxycarbonyl group were determined by an NOE experiment on 1, where a 2.0% NOE from the vinyl proton to 4a-H, and 4.3% and 6.2% NOE from 4a-H and 4-H_a to the vinyl proton, respectively, were observed. Whereas no NOE was observed between methoxycarbonyl group and 4-H_a, 4a-H protons. Thus the position of the vinyl proton at δ 6.56 was established at 3 position. These NOE also indicated the configuration of 4a-H and 4-H_a to be β , supposed the cyclopropane ring to be α . Coupling constant between 4a-H and 5-H_a indicate that they are in *cis* configuration. Thus the relative structure of 1 was determined as shown in Fig. 1. 1 seems to be an intermediate in duocarmycin biosynthesis from its structural features.

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