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# Streptocarbazoles A and B, Two Novel Indolocarbazoles from the Marine-Derived Actinomycete Strain *Streptomyces* sp. FMA

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#### **ABSTRACT**



Streptocarbazoles A (1) and B (2), two novel indolocarbazoles featuring unprecedented cyclic N-glycosidic linkages between 1,3-carbon atoms of the glycosyl moiety and two indole nitrogen atoms of the indolocarbazole core, were isolated from the marine-derived actinomycetes strain *Streptomyces* sp. FMA. Their structures were established by spectroscopic methods, CD spectra, and ECD quantum mechanical calculations. Compound 1 was cytotoxic on HL-60 and A-549 cell lines and could arrest the cell cycle of Hela cells at the  $G_2/M$  phase.

During the last 35 years, more than 130 indolocarbazoles (ICZs) have been isolated from different organisms including bacteria, fungi, and invertebrates. ICZs, kinds of alkaloids featuring a K252c<sup>2</sup> skeleton and represented by staurosporine, have succeeded in attracting great attention from chemists, biologists, physicians, and pharmaceutical companies for their unusual structures and important biological activities.<sup>4</sup> Cytotoxicity,<sup>5</sup> protein kinase C (PKC),<sup>6</sup> and topoisomerase (Topo)<sup>7</sup> inhibitions are by far the most widely searched and reported biological activities for ICZs. Several ICZs, such as UCN-01,<sup>8</sup> lestaurtinib,<sup>9</sup> and NSC655649,<sup>10</sup> are currently being tested in the cancer clinical trials. As part of our ongoing research on new antitumor compounds from marine-derived actinomycetes,<sup>11</sup> strain FMA, identified as *Streptomyces* 

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sp., was isolated from a mangrove soil collected in Sanya, Hainan Province of China. The EtOAc extracts of the fermentation broth of strain FMA exhibited significant cytotoxic effect on the P388 cell line. A series of metabolites contained in the extracts showed UV absorptions similar to those of ICZs at 227, 282, 325, and 355 nm<sup>3,12</sup> in an HPLC-UV profile (Figure S14, Supporting Information). Chemical investigation resulted in the isolation of two novel ICZs, which have been named the streptocarbazoles A (1) and B (2),<sup>13</sup> as well as five known analogues, K252c (indolocarbazole core),<sup>2</sup> K252a,<sup>14</sup> 3'-epi-K252a,<sup>14</sup> RK 286c, 15 and 4-bis(3-indolyl)-1*H*-pyrrole-2,5-dione. 16 Streptocarbazoles A (1) and B (2) possess exceptional cyclic N-glycosidic linkages between the 1,3-carbons of the glycosyl moiety and two indole nitrogen atoms of K252c. This connection was first discovered in the family of ICZs, and all 66 reported cyclic indolocarbazole glycosides were N-glycosidic linkages between the 1,5-carbons of the glycosyl moiety and the indolocarbazole core. In addition, there were only seven examples of cyclic O-glycosides in the literature, and the two glycosidic bonds were formed between the 1,2carbons of the glycosyl moiety and the aglycone. 11b,17

Streptocarbazole A (1) was assigned a molecular formula of C<sub>28</sub>H<sub>23</sub>N<sub>3</sub>O<sub>5</sub> on the basis of HRESIMS, <sup>18</sup> requiring 19 degrees of unsaturation. Its UV spectrum showed a characteristic peak of indolocarbazole chromophore at  $\lambda_{\rm max}$  227, 282, 325, and 355 nm.<sup>3,12</sup> The IR absorption bands at 3329, 1681, 1590, and 1454 cm<sup>-1</sup> suggested the presence of hydroxy, aromatic nucleus, and carbonyl groups. The <sup>1</sup>H and <sup>13</sup>C NMR spectra (Table 1) displayed signals similar to those of staurosporine, <sup>19</sup> further indicating an indolocarbazole unit that was confirmed by <sup>1</sup>H-<sup>1</sup>H COSY of H-1/H-2/H-3/H-4 and H-8/H-9/H-10/H-11 and by the key HMBC correlations of H-4 to C-13a and C-4b. H-6 to C-4c and C-7a, H-8 to C-7b and C-11a, and H-11 to C-7c (Figure 1). Moreover, careful comparison of its <sup>1</sup>H and <sup>13</sup>C NMR spectra (Table 1) with those of staurosporine<sup>19</sup> revealed that the glycosyl moiety is different, and two olefinic quaternary carbons ( $\delta_C$  134.2 and 145.2) are present in the sugar moiety of 1. This could be explained by the oxidation followed by enolization of the sugar moiety. <sup>1</sup>H-<sup>1</sup>H COSY from H-1' to H-2' and the key HMBC

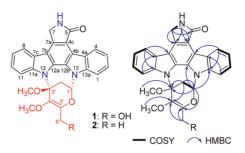


Figure 1. Structures and key  ${}^{1}H-{}^{1}H$  COSY and HMBC correlations of 1 and 2.

correlations of H-2' to C-1', C-3' and C-4', 3'-OCH<sub>3</sub> to C-3', 4'-OCH<sub>3</sub> to C-4', H-6' to C-4' and C-5', and H-1' to C-3', C-5' and C-12b suggested that the glycosyl moiety is 1,3-disubstituted 3,4-di-O-methyl-2-deoxy-4,5-dehydropyranoglucose and the glycosides are formed via the linkages of  $C_{1'}$ - $N_{13}$  and  $C_{3'}$ - $N_{12}$ . Cis- and trans- could be two possible candidate relative configurations of 1.

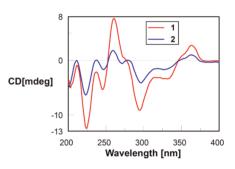


Figure 2. CD spectra of streptocarbazoles A (1) and B (2).

Model building of each via HyperChem 7.5 soft MM+ energy minimization calculations suggested that *trans-*1 possessed significant intraring strain and higher molecular energy (Figure S1, Supporting Information). Furthermore, chemical shifts of <sup>13</sup>C NMR were computed at the B3LYP/6-311++G(2d,p)//B3LYP/6-31G(d) level.<sup>20</sup> The magnetic shielding values were converted into chemical shifts after the corrections using slope and intercept of the linear-square functions,<sup>20</sup> and the relative errors of chemical shifts were computed by subtracting the calculated <sup>13</sup>C NMR from the experimental shifts. The relative errors between the measured <sup>13</sup>C shifts and the calculated <sup>13</sup>C shifts are less than 7.2 ppm in *cis-*1, while the maximum error at C-5′ reached 14.9 ppm in *trans-*1 (Table S2, Supporting Information). Thus, the relative configuration of 1 was deduced as *cis.* 

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<sup>(18)</sup> Streptocarbazole A (1): yellow, amorphous powder,  $[\alpha]^{21}_{D}$  – 54 (c 0.1, MeOH); UV (MeOH)  $\lambda_{\rm max}$  ( $\log \varepsilon$ ) 227 (3.32), 282 (3.65), 325 (3.15), 355 (3.09) nm; CD (MeOH)  $\lambda_{\rm max}$  ( $\Delta \varepsilon$ ) 364 (+1.5), 292 (-4.3), 259 (+4.1), 246 (-2.7), 237 (+0.6), 224 (-6.1), 212 (+1.5), 202 (-3.5) nm; IR (KBr)  $\nu_{\rm max}$  3329, 2936, 1590, 1454, 1398, 1354, 1330, 1139, 1099, 1057, 1021, 746 cm<sup>-1</sup>; <sup>1</sup>H and <sup>13</sup>C NMR, see Table 1; HR ESIMS m/z 482.1715 [M + H]<sup>+</sup> (calcd for  $C_{28}H_{24}N_{3}O_{5}$ , 482.1716).

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**Table 1.** <sup>1</sup>H and <sup>13</sup>C NMR Data for Streptocarbazoles A (1) and B (2) (600, 150 MHz, DMSO- $d_6$ , TMS,  $\delta$  ppm)

	1		2	
position	$\delta_{ m C}$	$\delta_{\rm H}(J~{\rm in~Hz})$	$\delta_{ m C}$	$\delta_{\mathrm{H}}\left(J\ \mathrm{in}\ \mathrm{Hz} ight)$
1	110.5, CH	7.82, d(8.3)	110.3, CH	7.82, d(8.2)
2	126.5, CH	7.55, t(8.3)	126.5, CH	7.55, t (8.2)
3	121.1, CH	7.35, t(7.8)	121.1, CH	7.36, t(7.8)
4	$126.2, \mathrm{CH}$	9.37, d (7.8)	126.3, CH	9.37, d(8.2)
4a	123.4, C		123.4, C	
4b	116.3, C		116.3, C	
4c	120.1, C		120.1, C	
5	172.3, C		172.3, C	
6-NH		8.65, s		8.67, s
7	$45.8, \mathrm{CH}_2$	4.98, s	$45.8, \mathrm{CH}_2$	4.99, s
7a	134.2, C		133.3, C	
7b	116.9, C		116.8, C	
7c	123.5, C		123.5, C	
8	$121.6, \mathrm{CH}$	8.03, d(7.8)	121.6, CH	8.03, d (7.8)
9	$121.2, \mathrm{CH}$	7.35, t (7.8)	121.1, CH	7.36, t(7.8)
10	$126.5, \mathrm{CH}$	7.53, t(8.2)	126.5, CH	7.52, t(8.2)
11	$115.4, \mathrm{CH}$	8.23, d(8.7)	115.4, CH	8.24, d (8.2)
11a	141.1, C		141.1, C	
12a	133.9, C		133.9, C	
12b	125.5, C		125.5, C	
13a	139.7, C		139.6, C	
1'	77.5, CH	7.04, d(5.1)	77.6, CH	7.04, d(6.0)
2'	$37.9, \mathrm{CH}_2$	3.41, dd	$37.8, CH_2$	3.42, dd
		(14.5, 6.0);		(15.6, 5.9);
		3.57, d (14.5)		3.55, d (15.6)
3'	87.8, C		87.8, C	
4'	134.2, C		134.0, C	
5'	145.3, C		142.9, C	
6′	$56.3, \mathrm{CH}_2$	3.68, d(12.8); 3.82, d(12.8)	$14.8$ , $CH_3$	1.59, s
3'-OCH <sub>3</sub>	$51.6, CH_3$	3.56, s	$51.6, CH_3$	3.57, s
4'-OCH <sub>3</sub>	, ,	3.28, s	$62.1,\mathrm{CH}_3$	3.14, s

An extension of the lactone sector rule<sup>21</sup> could be used to determine the absolute configuration of 1. The molecule was viewed from the line on the plane of the amide group along the bisectrix of the N-C=O angle, i.e., the line from C-5 to C-1' as shown in Figure 3 for (1'R, 3'S)-1. The functional group at C-3' lying in the back upper left sector was responsible for the negative Cotton effect  $(\pi - \pi^*)$ transition of conjugated lactam), which was good in accordance with the measured negative CD Cotton effect at  $\lambda_{\text{max}}$  292 nm of 1 (Figure 2). Moreover, the quantum chemical ECD calculation method<sup>22</sup> was used to confirm the absolute configuration of 1. The preliminary conformational distribution search was performed by HyperChem 7.5 software. The corresponding minimum geometries were further fully optimized by using DFT at the B3LYP/6-31G(d) level as implemented in the Gaussian 03 program package. The stable conformers obtained were submitted to ECD calculation by the TDDFT [B3LYP/6-31G(d)] method. The overall predicted ECD spectrum of 1 was subsequently compared with the measured one. The measured

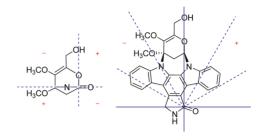


Figure 3. Application of an amide sector rule to 1.

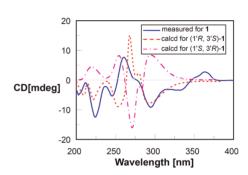


Figure 4. Measured and calculated ECD spectra for 1.

CD curve of **1** showed Cotton effect at  $\lambda_{\text{max}}(\Delta \varepsilon)$  292 (-4.3), 259 (+4.1) and 224 (-6.1) nm, matching with the calculated ECD curve of (1'*R*, 3'*S*)-**1** and opposite to that of (1'*S*, 3'*R*)-**1** (Figure 4), and the B3LYP/6-31G(d)-optimized conformations were also used in optical rotation ( $[\alpha]_D$ ) computations at the B3LYP/6-311G(d,p) level.<sup>23</sup> The calculated  $[\alpha]_D$  value for (1'*R*, 3'*S*)-**1** was -56.6, close to the measured value of -54. Thus, the absolute configuration of **1** was unambiguously established to be (1'*R*,3'*S*).

The molecular formula of streptocarbazole B (2) was assigned to be  $C_{28}H_{23}N_3O_4$  based on the HRESIMS,<sup>24</sup> which was only one oxygen less than that of 1. Careful comparison of its <sup>1</sup>H and <sup>13</sup>C NMR spectra (Table 1) with those of 1 showed that a methyl signals at  $\delta_{C/H}$  14.8/1.59 in 2 replaced the corresponding hydroxymethyl signals at  $\delta_{C/H}$  56.2/3.68 and 3.82 in 1, suggesting that the glycosyl moiety in 2 is 6-deoxygenated. The HMBC correlations from H<sub>3</sub>-6' to C-5' and C-4' also supported this change in 2 (Figure 1). The CD Cotton effects at  $\lambda_{max}(\Delta\varepsilon)$  292 nm ( $\Delta\varepsilon$  -1.4), 259 nm ( $\Delta\varepsilon$  +1.2) and 224 nm ( $\Delta\varepsilon$  -2.4) (Figure 2) and the specific rotation ([ $\alpha$ ] -23) were almost

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<sup>(24)</sup> Streptocarbazole B (2): yellow, amorphous powder,  $[\alpha]^{21}_{\rm D}-23$  (c 0.1, MeOH); UV (MeOH)  $\lambda_{\rm max}$  (log  $\varepsilon$ ) 227 (3.30), 282 (3.58), 325 (3.13), 355 (3.05) nm; CD (MeOH)  $\lambda_{\rm max}$  ( $\Delta\varepsilon$ ) 364 (+0.4), 292 (-1.4), 259 (+1.2), 246 (-0.8), 236 (+0.3), 224 (-2.4), 211 (+0.7), 200 (-2.2) nm; IR (KBr)  $\nu_{\rm max}$  2924, 1632, 1454, 1387, 1354, 1330, 1295, 1144, 1062, 1020, 747 cm $^{-1}$ ; 'H and  $^{13}$ C NMR, see Table 1; HRESIMS m/z 466.1758 [M + H]+ (calcd for  $C_{28}H_{24}N_3O_4$ , 466.1767).

Scheme 1. Plausible Biosynthetic Pathway of 1 and 2

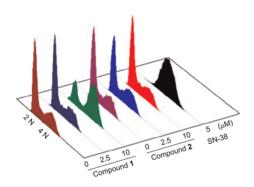


Figure 5. DNA histograms of Hela cells treated with different concentrations of streptocarbazoles A (1) and B (2).

the same to those of 1, indicating the same absolute configuration. Thus, the structure of streptocarbazole B (2) was clearly elucidated as 6'-deoxystreptocarbazole A.

A plausible biogenetic pathway for streptocarbazoles A (1) and B (2) was postulated (Scheme 1). The indolocarbazole unit (K252c) was derived from tryptophan, <sup>25</sup> while the glycosyl moiety was probably developed from 2-deoxy-pyranoglucose. The glycosidation of 12-NH of K252c with 2-deoxypyranoglucose produced glycoside a that serves the key biosynthetic precursor. Glycoside a underwent an

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oxidation and an enolization to yield another glycoside **b** which further underwent aldol condensation followed by methylation to yield **1**. Streptocarbazole B (**2**) was probably formed from K252c and 2,6-dideoxy-D-pyranoglucose by the same biogenetic procedures.

Compounds 1 and 2 were tested for cytotoxic effects on the HL-60, A-549, and P388 cell lines using the SRB method<sup>26</sup> and on the Hela cell line using the MTT method.<sup>27</sup> The results showed that compound 1 is effective on HL-60, A-549, P388 and Hela cells, with  $IC_{50}$  values of 1.4, <sup>13</sup> 5.0, <sup>13</sup> 18.9, and 34.5  $\mu$ M, respectively. And compound 2 was active against P388 and Hela cells with IC<sub>50</sub> values of 12.8 and 22.5 µM, respectively, but no effects on HL-60 and A-549 cells were observed (IC<sub>50</sub> > 50  $\mu$ M). The inhibitory effects of 1 and 2 on the cell cycle<sup>28</sup> and the kinase<sup>29</sup> were also evaluated. The results showed that compound 1 arrested the cell cycle of Hela cells in the G<sub>2</sub>/M phase at a concentration of 10 µM (Figure 5) and both new compounds did not show obvious inhibitory effects on the kinase GSK3-beta, Aurora-A, PKC-beta and PKC-epsilon (Table S1, Supporting Information).

Compounds 1 and 2 had been applied for the invention patents in China for the novel structures and the cytotoxicity against HL-60 and A-549 cell lines. <sup>13</sup> In this paper, the absolute configurations were revised and the bioactivities such as inhibition of cell cycle and the kinases and the cytotoxicity against Hela and P388 cell lines were complemented.

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**Supporting Information Available.** Experimental details, the NMR spectra of 1 and 2, and bioassay protocols used. This material is available free of charge via the Internet at http://pubs.acs.org.

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The authors declare no competing financial interest.