## Structure Revision of FD-891, a 16-Membered Macrolide Antibiotic

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FD-891 was isolated from the fermentation broth of *Streptmyces graminofaciens* A-8890 in 1994, and was shown to have a cytotoxic activity *in vitro* against several tumor cell lines<sup>1</sup>). The structure of FD-891 was first proposed to be an 18-membered macrolactone by spectroscopic means<sup>2</sup>), and recently, the stereochemistry of

Fig. 1. Proposed structures of FD-891.

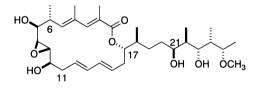
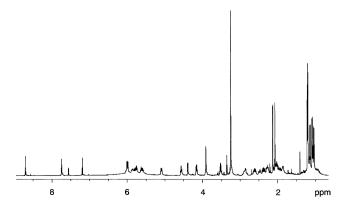


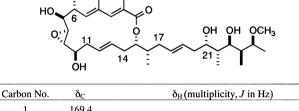
Fig. 2. <sup>1</sup>H-NMR spectrum of FD-891 in pyridine- $d_5$  (400 MHz).



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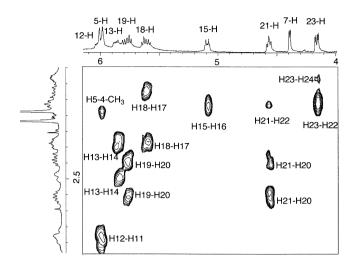
each chiral center of FD-891 was determined through synthetic studies of relevant fragments as well as X-ray diffraction of degradative derivatives as shown in Fig. 1<sup>3)</sup>. During our synthetic studies of FD-891 and its related compound, FD-892, we noticed the synthesized fragments including the C12-C15 conjugated double bond moiety showed a quite different signal pattern in <sup>1</sup>H NMR, especially the chemical shifts of the double bonds compared to those reported in natural FD-891 and FD-892.

Table 1. <sup>1</sup>H and <sup>13</sup>C NMR data of FD-891 in pyridine- $d_5$ .



Carbon No.	δ <sub>C</sub>	$\delta_{\rm H}$ (multiplicity, J in Hz)
1	169.4	
2	124.0	
3	145.5	7.62 (s)
4	135.2	
5	144.4	5.88 (d, 5.2)
6	38.0	3.12 (m)
7	71.3	4.27 (d, 3.6)
8	56.1	3.23 (s)
9	57.8	3.79 (s)
10	72.8	3.78 (m)
11	39.0	2.76 (m)
12	131.0	5.90 (m)
13	127.5	5.73 (ddd, 4.8, 9.6, 14.8)
14	35.1	2.36 (brd, 13.6 ), 2.15 (m)
15	77.0	4.96 (dt, 6.4, 3.2)
16	35.1	1.90 (m)
17	36.5	2.15 (m), 1.90 (m)
18	129.1	5.48 (dt, 15.2, 7.2)
19	130.9	5.64 (dt, 15.2, 6.8)
20	39.0	2.49 (dt, 13.6, 7.2) 2.25 (dt, 13.6, 6.8)
21	71.4	4.46 (t, 6.0)
22	40.4	1.90 (m)
23	74.8	4.04 (dd, 3.2, 8.0)
24	41.5	1.74 (m)
25	80.8	3.39 (quintet, 6.4)
26	8.9	1.08 (d, 6.0)
2-CH3	14.0	2.01 (s)
4-CH <sub>3</sub>	15.8	1.95 (s)
6-CH <sub>3</sub>	17.1	1.02 (d, 6.8)
16-CH <sub>3</sub>	10.8	0.91 (d, 6.4)
22-CH <sub>3</sub>	16.7	0.96 (d, 6.8)
24-CH <sub>3</sub>	16.6	1.09 (d, 6.8)
25-OCH <sub>3</sub>	56.2	3.11 (s)

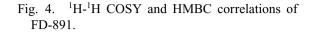
Fig. 3.  ${}^{1}\text{H}{}^{-1}\text{H}$  COSY specrum of FD-891 in pyridine- $d_{5}$ .

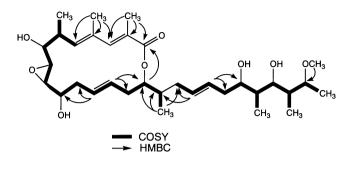


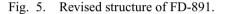
The numbering is according to Table 1.

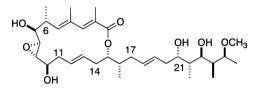
Therefore, we undertook re-investigation of the structure of FD-891.

In order to clarify the position of the double bonds, we attempted several different solvents for <sup>1</sup>H NMR studies, since the original description suggested complex signal overlaps in CDCl<sub>3</sub><sup>2)</sup>. As a result, moderate spectral resolution was observed in pyridine- $d_5$  as shown in Fig. 2. Under these conditions, we examined several NMR spectra including COSY and HMBC spectra. The resulting <sup>1</sup>H and <sup>13</sup>C NMR data are summarized in Table 1. As shown in <sup>1</sup>H-<sup>1</sup>H COSY spectrum (Fig. 3), the conectivities of each double bond to the neighboring protons were clearly observed. These results clearly indicated that two disubstituted E-double bonds must be in isolated environments, but not in conjugated to each other. Based on these results, the plain structure of FD-891 has now been finalized to be a 16-membered macrolactone as shown in Fig. 4. This plain structure turned out to be the same as BE-45653<sup>4)</sup> reported in 1997. Previously, we encountered certain difficulty in explaining the results of ozonolysis studies, however, the present revised structure of FD-891 is quite consistent with our degradation studies<sup>3)</sup>. The absolute structure of FD-891 is now depicted as shown in Fig. 5.









Although we cannot confirm the structure of FD-892, analog of FD-891, because the organism does not produce FD-892 any more, it seems likely that FD-892 also has the same carbon skeleton.

## References

- SEKI-ASANO, M.; T. OKAZAKI, M. YAMAGISHI, N. SAKAI, K. HANADA & K. MIZOUE: Isolation and characterization of new 18-membered macrolides FD-891 and FD-892. J. Antibiotics 47: 1226~1233, 1994
- SEKI-ASANO, M.; Y. TSUCHIDA, K. HANADA & K. MIZOUE: Structures of new 18-membered macrolides FD-891 and FD-892. J. Antibiotics 47: 1234~1241, 1994
- EGUCHI, T.; K. KOBAYASHI, H. UEKUSA, Y. OHASHI, K. MIZOUE, Y. MATSUSHIMA & K. KAKINUMA: Stereostructure of a novel cytotoxic 18-membered macrolactone antibiotic FD-891. Org. Lett. 4: 3383~3386, 2002
- OGAWA, H.; S. NAKAJIMA, H. SUZUKI, K. OJIRI & H. SUDA: Antitumor agent BE-45653 manufacture with streptomyces. Jpn. Kokai Tokkyo Koho, 09087285, 1997