## **NOTES**

# EI-2346, a Novel Interleukin-1β Converting Enzyme Inhibitor Produced by Streptomyces sp. E-2346

### II. Structure Elucidation

Fumito Koizumi\*, Atsuhiro Hasegawa<sup>a</sup>, Mayumi Yoshida<sup>b</sup>, Yuzuru Matsuda<sup>c</sup> and Satoshi Nakanishi<sup>d</sup>

Tokyo Research Laboratories, Kyowa Hakko Kogyo Co., Ltd., 3-6-6 Asahimachi, Machida-shi, Tokyo 194-8533, Japan

(Received for publication January 9, 2004)

The IL-1 $\beta$  converting enzyme (ICE) is a cysteine protease which cleaves biologically-inactive 31 kDa precursor to biologically-active IL-1 $\beta^{1,2}$ , a key mediator of inflammation<sup>3,4</sup>). Thus, ICE inhibitors would be useful as anti-inflammatory agents. As described in previous paper<sup>5</sup>, we isolated novel ICE inhibitory compound, EI-2346, from culture broth of *Streptomyces* sp. E-2346. In this paper, we describe the structure elucidation of EI-2346.

The physico-chemical properties of EI-2346 (1) were summarized in previous paper<sup>5)</sup>. The molecular formula of

Fig. 1. Structure of EI-2346.

$$H_3C$$
 $H_3C$ 
 $H_3C$ 

1 was determined by high resolution FAB-MS to be  $C_{22}H_{26}O_9$ . The <sup>1</sup>H and <sup>13</sup>C NMR data of 1 were summarized in Table 1, and COSY, HMBC, and NOESY correlations were shown in Fig. 2. The <sup>13</sup>C NMR spectrum (Table 1) showed 22 carbon signals which supported the molecular formula of 1. UV and IR spectra of 1 were consistent with

Table 1. <sup>13</sup>C and <sup>1</sup>H NMR data for EI-2346.

No.	$\delta C^a (ppm)$	δH <sup>b</sup> (ppm, multi, J in Hz)
1	70.6	5.11, m
3	95.1	
4	41.5	2.95, d, <i>J</i> =16.5
		2.86, d, <i>J</i> =16.5
4a	144.3	
5	120.6	7.33, s
5a	131.1	
6	185.9	
7	136.8	7.10, s
8	146.1	
9	189.9	•
9a	114.4	
10	159.0	
10-OH		
10a	134.8	
11	37.1	1.98, m (2H)
12	19.1	1.43, m
		1.27, m
13	14.4	0.89, t, <i>J</i> =7.4
14	29.2	1.54, s
15	95.2	5.72, s
17	84.4	3.64, m
18	62.3	3.64, m
19	72.4	4.19, dd, <i>J</i> =4.5, 10.3
		3.57, dd, <i>J</i> =10.3, 10.3
21	62.4	3.89, dd, <i>J</i> =2.0, 12.2
		3.74, dd, <i>J</i> =5.1, 12.2

a 13C NMR (100 MHz in CD<sub>3</sub>OD)

b 1H NMR (400 MHz in CD<sub>3</sub>OD)

<sup>\*</sup> Corresponding author: fumito.koizumi@kyowa.co.jp

<sup>&</sup>lt;sup>a</sup> Present address: Pharmaceutical Research Institute, Kyowa Hakko Kogyo Co., Ltd., 1188 Shimotogari, Nagaizumi-cho, Sunto-gun, Shizuoka 411-8731, Japan.

b Present address: Genomic Sciences Center, RIKEN, 1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama-shi, Kanagawa, 230-0045, Japan.

<sup>&</sup>lt;sup>c</sup> Present address: President, Kyowa Hakko Kogyo Co., Ltd., 1-6-1 Ohtemachi, Chiyoda-ku, Tokyo 100-8185, Japan.

d Present address: Strategic Planning Department, Pharmaceuticals Company, Kyowa Hakko Kogyo Co., Ltd., 1-6-1 Ohtemachi, Chiyoda-ku, Tokyo 100-8185, Japan.

naphthoquinone chromophore, and the UV spectrum of 1 was similar to that of exfoliamycin<sup>6,7)</sup> which also has a naphthoquinone chromophore. The <sup>1</sup>H and <sup>13</sup>C NMR spectra of 1 were similar to those reported for K1115B<sub>1</sub><sup>8</sup>, another secondary metabolite isolated from Streptomyces sp., which has a naphthoquinone structure and a 1,3-dioxan ring. Upon comparison of NMR data (chemical shifts of <sup>1</sup>H and <sup>13</sup>C, and <sup>1</sup>H-<sup>1</sup>H coupling constants) of 1 with those of K1115B<sub>1</sub><sup>8)</sup>, naphthoquinone and 1,3-dioxan moieties were found as common partial structure. The molecular formula of 1 (C<sub>22</sub>H<sub>26</sub>O<sub>9</sub>) which was determined by high resolution FAB-MS corresponds to H<sub>2</sub>O more than that of K1115B<sub>1</sub>  $(C_{22}H_{24}O_8)$ . The NMR analysis of 1 indicated that this additional unit was located in the 3,4-double bond of pyran ring of K1115B<sub>1</sub>. In the <sup>1</sup>H and <sup>13</sup>C NMR spectra of 1, signals which were attributed to 3,4-double bond of pyran ring ( $\delta$  C-3 158.7,  $\delta$  C-4 100.0, and  $\delta$  H-4 5.6) in K1115B<sub>1</sub> were replaced by those of hemiacetal carbon ( $\delta$  C-3 95.1) and methylene ( $\delta$  C-4 41.5,  $\delta$  H-4 2.95 and 2.86). Therefore, the structure of 1 should be the corresponding 3,4-hydrate derivative of K1115B<sub>1</sub>. This was also confirmed by the results that <sup>1</sup>H and <sup>13</sup>C NMR spectra of tricyclic naphthoquinone and tetrahydropyran moieties of 1 were similar to those reported for the known exfoliamycin which also has the same tricyclic structure. Moreover, this structure was supported by the 2D NMR study for 1 (Fig. 2).

Relative configuration of the 1,3-dioxan ring in 1 was determined by coupling constants and NOE experiment as shown in Fig. 2. NOEs were observed between H-15 ( $\delta_{H}$ 

Fig. 2. Summary of COSY, HMBC, and NOESY correlations for EI-2346.

5.72) and H-17 ( $\delta_{\rm H}$  3.64), H-15 and H-19<sub>ax</sub> ( $\delta_{\rm H}$  3.57). These results revealed that 1,3-dioxane moiety was chair conformation, and that these protons were of the 1,3-diaxial orientation. A large coupling constant between H-18 and H-19<sub>ax</sub> (J=10.3 Hz) indicated that these protons were of the 1,2-diaxial orientation and 18-OH was of the equatorial orientation. These results revealed the relative configuration of 1,3-dioxan ring shown in Fig. 2. Stereochemistries at C-1 and C-3 of EI-2346 were not determined yet.

### Acknowledgment

We would like to express thanks to Dr. SHINGO KAKITA for helpful discussion.

#### References

- THORNBERRY, N. A.; H. G. BULL, J. R. CALAYCAY, K. T. CHAPMAN, A. D. HOWARD, M. J. KOSTURA, D. K. MILLER, S. M. MOLINEAUX, J. R. WEIDNER, J. AUNINS, K. O. ELLISTON, J. M. AYALA, F. J. CASANO, J. CHIN, G. J.-F. DING, L. A. EGGER, E. P. GAFFNEY, G. LIMJUCO, O. C. PALYHA, S. M. RAJU, A. M. ROLANDO, J. P. SALLEY, T.-T. YAMIN, T. D. LEE, J. E. SHIVELY, M. MACCROSS, R. A. MUMFORD, J. A. SCHMIDT & M. J. TOCCI: A novel heterodimeric cysteine protease is required for interleukin-1β processing in monocytes. Nature 356: 768~774, 1992
- 2) Gerretti, D. P.; C. J. Kozlosky, B. Mosley, N. Nelson, K. V. Ness, T. A. Greenstreet, C. J. March, S. R. Kronheim, T. Druck, L. A. Cannizzaro, K. Huebner & R. A. Black: Molecular cloning of the interleukin-1 $\beta$  converting enzyme. Science 256: 97~100, 1992
- 3) DINARELLO, C. A.: Interleukin-1 and interleukin-1 antagonism. Blood 77: 1627~1652, 1991
- DINARELLO, C. A. & S. M. WOLFF: The role of interleukin-1 in disease. N. Engl. J. Med. 328: 106~113, 1993
- 5) KOIZUMI, F.; A. HASEGAWA, K. OCHIAI, K. ANDO, H. KONDO, M. YOSHIDA, Y. MATSUDA & S. NAKANISHI: EI-2346, a novel interleukin-1β converting enzyme inhibitor produced by *Streptomyces* sp. E-2346. 1. Taxonomy of producing strain, fermentation, isolation, physicochemical properties, and biological properties. J. Antibiotics 56: 985~992, 2003
- 6) POTTERAT, O.; H. ZAHNER, C. VOLKMANN & A. ZEECK: Exofoliamycin and related metabolites, new naphtoquinone antibiotics from *Streptomyces exfoliatus*. J. Antibiotics 46: 346~349, 1993
- VOLKMANN, C.; A. ZEECK, O. POTTERAT, H. ZAHNER, F. M. BOHNEN & R. H. IRMER: Metabolic products of microorganismen. 270<sup>†</sup>. The structures of the exofoliamycin. J. Antibiotics 48: 431~432, 1995
- 8) NARUSE, N.; M. GOTO, Y. WATANABE, T. TERASAWA & K. DOBASHI: K1115A, a new anthraquinone that inhibits the binding of activator-1 (AP-1) to its recognition sites. J. Antibiotics 51: 545~552, 1998