

A Novel Testosterone 5 α -Reductase Inhibitor, 8',9'-Dehydroascochlorin Produced by *Verticillium* sp. FO-2787

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A new testosterone 5 α -reductase inhibitor, 8',9'-dehydroascochlorin, was isolated from the cultured mycelium of *Verticillium* sp. FO-2787. This compound (M.W.: 402, C₂₃H₂₇ClO₄) consists of 5-chloroorcyclic aldehyde and 2,3,4-trimethylcyclohexenone moieties, which are connected via a chain of (2*E*,4*E*)-3-methyl-2,4-pentadiene. The IC₅₀ value of the new microbial metabolite for testosterone 5 α -reductase activity prepared from rat prostate was 4×10^{-4} M.

Keywords microbial metabolite; testosterone 5 α -reductase inhibitor; 8',9'-dehydroascochlorin; *Verticillium*; cultured mycelium

Androgens, particularly 5 α -dihydrotestosterone (DHT), play a key role in the differentiation, growth and maintenance of the mammalian prostate. Benign prostate hyperplasia has been associated with an overproduction of DHT which is converted from testosterone by testosterone 5 α -reductase (T-5 α -reductase).¹⁾ In the course of our search for new nonsteroidal inhibitors of T-5 α -reductase, 8',9'-dehydroascochlorin was discovered from the cultural mycelium of *Verticillium* sp. FO-2787, together with five known compounds, LL-Z 1272*e* (**1**), LL-Z 1272*y*

(ascochlorin, **2**), LL-Z 1272*β* (**3**), LL-Z 1272*ζ* (**4**), LL-Z 1272*δ* (**6**) (Fig. 1). We report here the isolation of 8',9'-dehydroascochlorin and substances related to ascochlorin with details of their structural determination and T-5 α -reductase inhibitory effects.

Strain FO-2787 isolated from a soil sample collected in Shizuoka prefecture, Japan, was determined to be a *Verticillium* species from morphological studies. From the ethyl acetate extracts of cultured mycelium, six metabolites (**1**–**6**) were isolated by repeated silica gel column chromatography and by preparative HPLC with the guidance of T-5 α -reductase inhibitory activity. Among these six compounds, five compounds (**1**–**4**, **6**) were reported as antiviral and anti-*Tetrahymena pyriformis* by a number of investigators.^{2,3)}

A new inhibitor (**5**), white plates, mp 127–130 °C, [α]_D²⁴ –80° (*c* = 0.5, MeOH), showed characteristic UV absorption of an α,β -unsaturated carbonyl at 348.0 nm (log ϵ , 3.95) and 292.4 nm (log ϵ , 3.91). The molecular formula was determined to be C₂₃H₂₇ClO₄ (M⁺, *m/z* 402.1590, Calcd 402.1598) by the high resolution-electron impact (HR-EI) mass spectrum, which was a corresponding derivative less 2 mass units from ascochlorin (**2**) [C₂₃H₂₉ClO₄]. The fragmentation at *m/z* 199 (100%) in the EI mass spectrum of **5** can be assigned to the peak corresponding to the 5-chloroorcyclic aldehyde moiety.⁴⁾ In the ¹H-NMR spectrum of **5**, signals appearing at δ 12.71 (1H, s) and δ 10.15 (1H, s) were assigned to a chelated phenolic proton (C-2) and an aldehyde proton (C-1), respectively, which were in good agreement with those of **2** (Table I).⁵⁾ The signal observed at δ 3.54 (2H, d, *J* = 7.6 Hz) was assigned to benzylic protons adjacent to an olefinic proton measured at δ 5.55 (1H, t, *J* = 7.6 Hz). Furthermore, two doublet-methyl signals at δ 0.95 and 0.98 (3H, d each) were characteristic of the cyclohexanone moiety of ascochlorins.⁵⁾ In the ¹H–¹H correlation spectroscopy (¹H–¹H COSY) spectrum, the signal of an olefinic proton at δ 5.99 (2H, dd-like, *J* = 16.2, 3.3 Hz) was coupled with that of δ 6.56 (1H, dd, *J* = 9.9, 2.0 Hz) and that of δ 5.42 (1H, d, *J* = 16.2 Hz). The coupling system (1H d, *J* = 16.2 Hz, each) at δ 5.99 and 5.42 is a typical AB type doublet assignable to a *trans* conjugated double

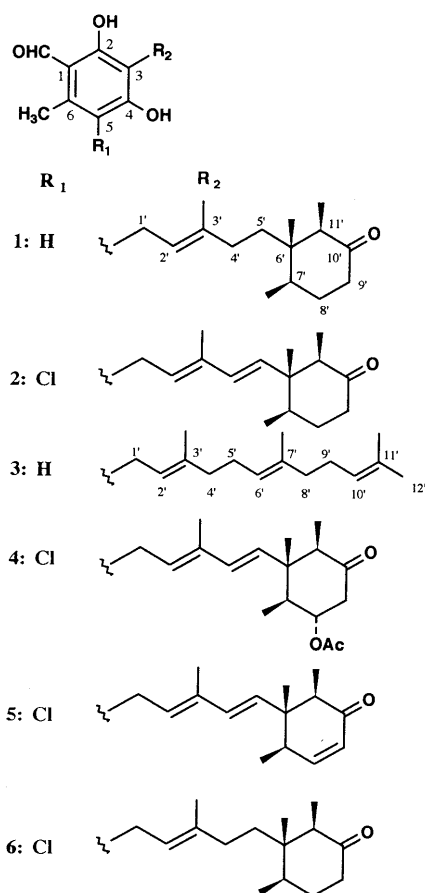


Fig. 1. Structures of Ascochlorins

TABLE I. ^1H -NMR Chemical Shifts of **1**–**6** in CDCl_3

H	1	2	3	4	5	6
1-CHO	10.06 (1H, s)	10.14 (1H, s)	10.06 (1H, s)	10.14 (1H, s)	10.15 (1H, s)	10.14 (1H, s)
2-OH	12.70 (1H, s)	12.71 (1H, s)	12.73 (1H, s)	12.71 (1H, s)	12.71 (1H, s)	12.69 (1H, s)
4-OH	6.92 (1H, br s)	—	6.44 (1H, br s)	—	—	—
5	6.24 (1H, s)	—	6.21 (1H, s)	—	—	—
6-Me	2.48 (3H, s)	2.60 (3H, s)	2.49 (3H, s)	2.61 (3H, s)	2.61 (4H, s)	2.60 (3H, s)
1'	3.37 (2H, d, 6.6)	3.53 (2H, d, 7.6)	3.40 (2H, d, 7.3)	3.54 (2H, d, 7.6)	3.54 (2H, d, 7.6)	3.39 (2H, d, 7.3)
2'	5.28 (1H, t, 6.6)	5.52 (1H, t, 7.6)	5.27 (1H, t, 7.3)	5.54 (1H, t, 7.6)	5.55 (1H, t, 7.6)	5.25 (1H, dd) ^d
4'a	2.10–1.83 (1H, m)	5.90	2.03 (1H, m)	5.92 (1H, d, 16.0)	5.99 (2H, dd) ^b	1.98 (1H, m)
4'b	1.63 (1H, m)	—	2.03 (1H, m)	—	—	1.89–1.77 (1H, m)
5'a	1.40 (2H, m)	5.38 (1H, d, 16.0)	2.03 (1H, m)	5.32 (1H, d, 16.0)	5.42 (1H, d, 16.2)	1.38 (1H, m)
5'b	1.40 (2H, m)	—	2.03 (1H, m)	—	—	1.38 (1H, m)
6'	—	—	5.07 (1H, br s)	—	—	—
7'	2.10–1.83 (1H, m)	1.99–1.87 (1H, m)	—	2.01 (1H, m)	2.61 (1H, s)	1.98 (2H, m)
8'a	2.10–1.83 (1H, m)	1.99–1.87	2.03 (1H, m)	4.89 (1H, m)	6.56 (1H, dd) ^c	1.89–1.77 (1H, m)
8'b	2.10–1.83 (1H, m)	1.63 (1H, ddd) ^a	2.03 (1H, m)	—	—	1.63 (1H, m)
9'a	2.34 (1H, m)	2.39 (3H, m)	2.03 (1H, m)	2.42–2.38 (1H, m)	5.99 (2H, dd) ^b	2.31 (1H, m)
9'b	—	—	2.03 (1H, m)	—	—	—
10'	—	—	5.07 (1H, br s)	—	—	—
11'	2.56–2.44	2.39 (1H, m)	—	2.42–2.38 (1H, m)	2.46 (1H, q, 6.8)	2.45 (1H, q, 6.6)
12'	—	—	1.67 (1H, s)	—	—	—
3'-Me	1.83 (3H, s)	1.92 (3H, s)	1.81 (3H, s)	1.92 (3H, s)	1.93 (3H, s)	1.81 (3H, s)
6'-Me	0.57 (3H, s)	0.69 (3H, s)	—	0.73 (3H, s)	0.80 (3H, s)	0.61 (3H, s)
7'-Me	0.92 (3H, d, 6.6)	0.83 (3H, d, 6.6)	1.59 (3H, br s)	0.86 (3H, d, 6.6)	0.95 (3H, d, 6.9)	0.90 (3H, d, 6.9)
11'-Me	0.88 (3H, d, 6.9)	0.81 (3H, d, 6.9)	1.59 (3H, br s)	0.86 (3H, d, 6.6)	0.98 (3H, d, 7.3)	0.88 (3H, d, 5.3)
Ac-Me	—	—	—	2.06 (3H, s)	—	—

a) $J=13.2, 12.9, 5.6\text{ Hz}$; b) $J=16.2, 3.3\text{ Hz}$, overlapped; c) $J=9.9, 2.0\text{ Hz}$; d) $J=7.3, 6.2\text{ Hz}$.

TABLE II. ^{13}C -NMR Chemical Shifts of **1**–**6** in CDCl_3

C	1	2	3	4	5	6
1	113.1 (s)	113.7 ^a (s)	113.2 (s)	114.6 ^a (s)	113.7 ^b (s)	114.3 ^a (s)
2	163.7 (s)	162.2 (s)	163.6 ^b (s)	162.2 (s)	162.2 (s)	162.2 (s)
3	112.1 (s)	113.6 ^a (s)	116.6 (s)	113.6 ^a (s)	113.6 ^b (s)	113.6 ^a (s)
4	162.3 (s)	156.2 (s)	162.6 ^b (s)	156.1 (s)	156.2 (s)	156.3 (s)
5	110.6 (d)	113.1 ^a (s)	110.8 (d)	113.1 ^a (s)	113.2 ^b (s)	113.2 ^a (s)
6	141.9 (s)	137.8 (s)	142.0 (s)	137.8 (s)	137.8 (s)	137.6 (s)
1'	21.2 (t)	22.2 (t)	21.2 (t)	22.2 (t)	22.2 (t)	22.1 (t)
2'	121.4 (d)	127.5 (d)	120.9 (d)	128.2 (d)	128.0 ^c (d)	121.0 (d)
3'	138.0 (s)	134.1 (s)	139.5 (s)	134.1 (s)	134.1 (s)	136.6 (s)
4'	32.6 (t)	133.2 (d)	39.7 ^c (t)	133.8 (d)	134.0 (d)	42.6 (t)
5'	35.6 (t)	135.6 (d)	26.6 ^d (t)	135.4 (d)	134.2 (d)	35.6 (t)
6'	43.6 (s)	48.4 (s)	124.3 ^e (d)	47.2 (s)	47.9 (s)	43.5 (s)
7'	36.0 (d)	40.8 (d)	135.6 (s)	45.6 (d)	41.9 (d)	36.1 (d)
8'	30.9 (t)	31.1 (t)	39.6 ^c (t)	73.7 (d)	152.1 (d)	31.0 (t)
9'	41.5 (t)	41.5 (t)	26.2 ^d (t)	45.3 (t)	127.9 ^c (d)	41.6 (t)
10'	215.0 (s)	212.8 (t)	123.5 ^e (d)	207.6 (s)	201.4 (s)	214.1 (s)
11'	50.5 (d)	53.6 (d)	131.3 (s)	53.8 (d)	51.8 (d)	50.4 (d)
12'	—	—	25.7 (q)	—	—	—
1-CHO	192.9 (d)	193.2 (d)	193.0 (d)	193.2 (d)	193.2 (d)	193.3 (d)
Methyl group						
6	18.0 (q)	14.4 (q)	18.0 (q)	12.7 (q)	14.4 (q)	14.5 (q)
3'	16.4 (q)	12.6 (q)	16.2 (q)	12.6 (q)	12.7 (q)	16.4 (q)
6'	15.3 (q)	10.3 (q)	—	11.6 (q)	10.0 (q)	15.3 (q)
7'	7.5 (q)	16.3 (q)	16.0 (q)	14.5 (q)	15.1 (q)	7.5 (q)
11'	15.0 (q)	8.9 (q)	17.6 (q)	8.9 (q)	9.0 (q)	15.1 (q)
Acetyl group						
Me	—	—	—	21.0 (q)	—	—
C=O	—	—	—	170.1 (q)	—	—

a–e) Assignments may be interchangeable.

bond. The residual coupling systems (1H, d, $J=9.9\text{ Hz}$) at δ 6.56 and 5.99 were assigned to α,β -unsaturated protons (C-8', C-9') in the cyclohexenone moiety of the molecule. The catalytic hydrogenation on 5% Pd/C of **5** gave

2. From these results, **5** was determined to be 8',9'-dehydroascochlorin. The ^{13}C -NMR chemical shifts of **5** were also assigned by ^1H - ^{13}C COSY spectrum (Table II). The relative structure of **5** was anticipated by nuclear

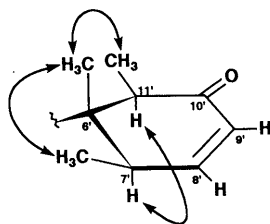
Fig. 2. NOESY Experiments for the Cyclohexenone Moiety of **5**

TABLE III. CD Cotton Effects of Ascochlorins

Compound	$\Delta\epsilon$ (nm)	Stereochemistry
1	-1.7 (290)	6'S, 7'R, 11'R
2	-7.3 (290)	6'R, 7'R, 11'R
4	-4.9 (290)	6'S, 7'S, 8'S, 11'R
5	-2.2 (327) -8.3 (244)	6'R, 7'R, 11'R
6	-1.3 (290)	6'S, 7'R, 11'R

TABLE IV. Testosterone 5 α -Reductase Inhibitory Activities

Compound	IC ₅₀ (M)
1	3.7×10^{-4}
2	3.4×10^{-4}
3	3.6×10^{-4}
4	3.4×10^{-4}
5	14.0×10^{-4}
6	3.7×10^{-4}
Riboflavin	1.3×10^{-3}

Overhauser effect spectroscopy (NOESY) experiments (Fig. 2). Several correlation peaks between 6'-methyl and 11'-methyl, 6'-methyl and 7'-methyl, and 7'-proton and 11'-proton were observed. Therefore, of the three methyl groups attached to the cyclohexenone ring, two were in equatorial (C-7', C-11') and the other in axial position (C-6'). The final confirmation of the absolute configuration of **5** was made by chemical transformation of **5** to ascochlorin (**2**) by catalytic hydrogenation on 5% Pd/C. Ascochlorin (**2**), whose absolute configuration was clarified by X-ray structure analysis,^{6,7)} showed a negative Cotton effect at 290 nm ($\Delta\epsilon$, -7.3) on the circular dichroism (CD) spectrum, whereas, **5** showed negative Cotton effects at 327 nm ($\Delta\epsilon$, -2.2) and at 244 nm ($\Delta\epsilon$, -8.3). Furthermore, ascochlorin (**2**) derived from **5** gave a negative Cotton effect at 290 nm ($\Delta\epsilon$, -7.2). These results clearly indicate that **5** has the same absolute configuration series as that of **2** (6'R, 7'R, 11'R). Consequently, the stereochemistry of **5** was confirmed to be 6'R, 7'R, 11'R. Other compounds (**1**, **4**, **6**) also showed similar Cotton effects to that of **2** (Table III). Therefore, a series of cyclohexanone rings of ascochlorin-analogs possesses the same absolute configuration as that of **2**. Compound **5** has already been prepared synthetically by Ellestad *et al.*,³⁾ but the present study is the first isolation and full characterization of **5** of microbial origin.

The inhibitory effect of compounds (**1**–**6**) on rat prostate testosterone 5 α -reductase was tested, and the IC₅₀ values of the compounds were 3.4 – 14.0×10^{-4} M as shown in Table IV. These compounds showed no

antimicrobial activities against gram-positive and gram-negative bacteria, fungi or yeast at a concentration of 500 μ g/ml.

As inhibitors of testosterone 5 α -reductase activity, several nonsteroidal compounds, including riboflavin, were isolated from the fermentation broth of microorganisms.^{8–11)} According to our present experiment, the inhibitory activities of **5** and related compounds were more than 10-fold greater than that of riboflavin. The structure of **5** was completely different from that of riboflavin and related compounds. Therefore, it would be of interest to examine **5** in *in vivo* experiments.

Experimental

¹H- and ¹³C-NMR spectra were recorded on a JEOL JNM-EX 270 spectrometer (¹H 270.05 MHz, ¹³C 67.8 MHz). The ¹H- and ¹³C-NMR signals were assigned with the aid of the ¹H–¹H, and ¹H–¹³C COSY, heteronuclear multiple bond connectivity (HMBC) and NOESY experiments. Melting points were measured using a Yanaco micromelting point apparatus and are uncorrected. Kieselgel 60 (70–230 mesh, 230–400 mesh, Merck) was used for column chromatography, and DC-Alufolien Kieselgel 60 F₂₅₄ (Merck) was used for TLC analysis. Analytical and preparative HPLC was carried out with MeOH–H₂O (8:2) and CH₃CN–H₂O (8:2) using Cosmosil 5C18-AR packed columns (Nacalai Tesque, 5 mm, i.d. 4.6 $\times$ 250 mm, i.d. 20 $\times$ 250 mm) employing a UV monitoring system (210 or 254 nm). CD spectra were recorded on a JASCO J-720 spectropolarimeter in MeOH.

Testosterone 5 α -Reductase Assay The assay mixture consisting of 30 μ l of 0.25 M sucrose–0.1 M HEPES buffer (pH 7.4), 10 μ l of 50 mM NADPH, 10 μ l of 0.75 nmol [¹⁴C]testosterone, 30 μ l of the test sample, and 20 μ l of the enzyme solution prepared from the prostate of Donryu male rats⁸⁾ was incubated for 2 h at 37 °C. *n*-Hexane (400 μ l) was added, and the mixture was shaken. The *n*-hexane layer (20 μ l) was applied to 0.25 mm-precoated silica gel plates (Merck), and the plates were developed with a solvent system of cyclohexane–ethyl acetate (1:1). The radioactive areas of [¹⁴C]testosterone and [¹⁴C]-5 α -dihydrotestosterone were determined using an AMBIS image analyzer. Riboflavin was purchased from Kanto Chemical Co., Inc.

Fermentation of Strain FO-2787 and Isolation of Active Compounds A stock culture of the producing organism was inoculated into a test tube (i.d. 2 $\times$ 20 cm) containing 10 ml of seed medium consisting of 2% glucose, 0.2% yeast extract, 0.05% MgSO₄·7H₂O, 0.5% polypeptone, 0.1% KH₂PO₄ and 0.1% agar (pH 6.0 before sterilization). The tube was incubated at 27 °C for 72 h on a reciprocal shaker. Then, 2 ml portions of the growth were transferred to a 500 ml Erlenmeyer flask containing 100 ml of the seed medium. The flask was incubated at 27 °C for 72 h on a rotary shaker (210 rpm), and 600 ml of the resulting culture was transferred into a 50-liter fermentor containing 30 l of the production medium consisting of 3% soluble starch, 1% glycerol, 2% soybean meal, 0.3% yeast extract, 0.3% KCl, 0.2% CaCO₃, 0.05% MgSO₄·7H₂O and 0.05% KH₂PO₄ (pH 6.5). The fermentation was carried out at 27 °C for 96 h using an agitation rate of 160 rpm and an aeration rate of 60 l per minute.

The ethyl acetate extracts of cultured mycelium were chromatographed on a silica gel (70–230 mesh) column using CHCl₃–MeOH (9:1). Furthermore, each active fractions was submitted to a silica gel (230–400 mesh) column using CHCl₃–MeOH (20:1→9:1) as the developing solvent. Finally, isolation of each compound (**1**–**6**) by use of preparative HPLC gave **1** (80 mg), **2** (60 mg), **3** (30 mg), **4** (6.5 mg), **5** (8 mg) and **6** (8 mg), respectively.

Catalytic Hydrogenation of 5 Compound **5** (1.0 mg) was dissolved in 3 ml of 0.01 N KOH–MeOH and hydrogenated at 1 atm over 5% palladium carbon at room temperature for 5 min. The resulting solution was neutralized by 0.01 N HCl and filtered through a Cosmonice filter (W-type, 0.45 μ m, Nacalai Tesque). The product was purified by preparative HPLC (MeOH–H₂O (8:2)) to give 0.65 mg of pure **2** (yield, 64.7%). The reaction product was identified as ascochlorin (**2**) by EI-MS, ¹H-NMR and co-HPLC with the authentic sample.

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