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LEUHISTIN, A NEW INHIBITOR OF AMINOPEPTIDASE M, PRODUCED BY *Bacillus laterosporus* BMI156-14F1

II. STRUCTURE DETERMINATION OF LEUHISTIN

SHIGEMI YOSHIDA, HIROSHI NAGANAWA, TAKAAKI AOYAGI and Tomio Takeuchi

Institute of Microbial Chemistry, 3-14-23 Kamiosaki, Shinagawa-ku, Tokyo 141, Japan

YASUO TAKEUCHI and YOSHIO KODAMA

Pharmaceutical Research Center, Meiji Seika Kaisha, Ltd., Morooka-cho, Kouhoku-ku, Yokohama 222, Japan

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Leuhistin, a new inhibitor of aminopeptidase M, has been isolated from the culture broth of *Bacillus laterosporus* BMI156-14F1. The structure of leuhistin was determined by NMR studies. X-Ray and chemical analysis confirmed the absolute structure to be (2R,3S)-3-amino-2-hydroxy-2-(1H-imidazol-4-ylmethyl)-5-methylhexanoic acid.

In the preceding paper¹, we have described the taxonomy, isolation, physico-chemical properties and biological properties of leuhistin, a novel inhibitor of aminopeptidase M (AP-M). In this paper, we describe the structure determination of leuhistin. The molecular formula of leuhistin was elucidated as $C_{11}H_{19}N_3O_3$ from the mass spectrum and the elemental analysis. The IR spectrum (KBr) of leuhistin showed absorption bands at 3400, 3130, 2970, 1625, 1470, 1395, 1270, 1220, 1150, 1090, 835 cm⁻¹.

Table 1 shows ¹H and ¹³C NMR data for leuhistin monohydrochloride. A positive color reaction with ninhydrin reagent suggested the presence of an amino group in the molecule. The chemical shifts of 3-H (δ 3.57) and C-3 (δ 55.8) indicated the amino group was attached to C-3. ¹H-¹H COSY spectrum of

Table 1. ¹H and ¹³C NMR spectral data for leuhistin monohydrochloride in D_2O .

Assignment	$^{1}\mathrm{H}^{\mathrm{a}}$	¹³ C ^b
1-CO		176.9 (s)
2-C		78.2 (s)
3-CH	3.57 dd (J=2.5, 10.8)	55.8 (d)
4-CH ₂	1.58 ddd $(J=3.2, 10.8, 13.2),$	37.1 (t)
2	1.70 ddd $(J=2.5, 10.2, 13.2)$	
5-CH	1.75 m	24.7 (d)
6-CH ₃	0.96 d (J = 5.6)	20.8 (q)
7-CH3	1.02 d (J=6.4)	23.5 (q)
8-CH	3.05 d (J=15.0), 3.25 d (J=15.0)	31.2 (t)
2'-CH	8.61 d $(J=1.6)$	133.9 (d)
4'-C	` ´ ´	128.7 (s)
5'-CH	7.28 br s	118.0 (d)

^a 400 MHz, δ in ppm, J in Hz.

^b 100 MHz, δ in ppm.

leuhistin revealed the partial structure (Fig. 1A). A positive color reaction with Pauly reagent and NMR data suggested the presence of an imidazole moiety (Fig. 1B). The other three carbons are shown as C,



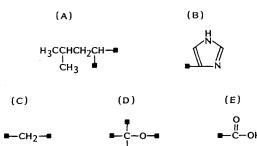
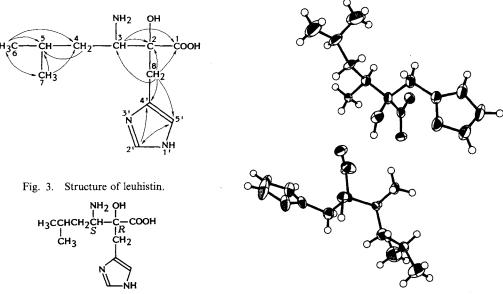


Fig. 4. An ORTEP drawing of leuhistin.

Fig. 2. Long range ¹H-¹³C coupling pattern of leuhistin.



D and E in Fig. 1. The heteronuclear multiple-bond correlation (HMBC) experiment (Fig. 2) indicated the linkage of these partial structures. Cross peaks between 8-H (δ 3.05 and 3.25) and imidazole carbons (C-4' δ 128.7 and C-5' δ 118.0) indicated that the methylene group is linked to the imidazole moiety. Long range coupling between 3-H and C-2 (δ 78.2), 8-H and C-3, and 8-H and C-2 suggested the quaternary carbon (C-2) is located between C-3 and C-8. Long range coupling between 3-H and C-1 (δ 176.9), and between 8-H and C-1 indicated that C-2 is connected to C-1. The C-1 carbon was identified as the carbon of carboxyl group from its chemical shift (δ 176.9). This was supported by the pKa values (2.3, 7.5 and 9.6) of leuhistin. Another oxygen atom, which is linked to the quaternary oxycarbon (C-2), was identified as a hydroxyl group from consideration of the molecular formula. Thus the structure of leuhistin was determined to be 3-amino-2-hydroxy-2-(1*H*-imidazol-4-ylmethyl)-5-methylhexanoic acid (Fig. 3). The absolute stereochemistry at C-3 was determined by oxidation of leuhistin. Treatment with potassium permanganate gave leucine that was identified as the L-enantiomer by TLC on a chiral precoated HPTLC plate. Thus the configuration at C-3 was found to be *S*.

The relative stereochemistry of C-2 and C-3 of leuhistin was determined by a crystal X-ray diffraction analysis using leuhistin free base. An $ORTEP^{2}$ drawing of leuhistin is shown in Fig. 4. The X-ray analysis showed that the relative configuration of C-2 and C-3 is *threo*.

Therefore, the structure of leuhistin was determined to be (2R,3S)-3-amino-2-hydroxy-2-(1H - imidazol-4-ylmethyl)-5-methylhexanoic acid (Fig. 3).

Experimental

The mass spectrum was obtained on a Hitachi M-80H mass spectrometer, the IR spectrum on a Hitachi 260-10 spectrophotometer and the NMR spectra on a Jeol JNM-GX400 NMR spectrometer with ¹H NMR at 400 MHz and ¹³C NMR at 100 MHz.

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Chemicals

Chemicals employed were as follows: PK208 (cation exchange resin) from Nippon Rensui Co., Tokyo, Japan; HPTLC pre-coated CHIR plates from E. Merck, Darmstadt, FRG. All other chemicals were of analytical grade.

Oxidation of Leuhistin

Potassium permanganate (450 mg) was added to a solution of 100 mg of leuhistin in 5 ml of water. After 5 hours at room temperature, the reaction mixture was filtered, and the filtrate was applied to a column of PK208 (free acid form). Elution with 0.5 N ammonium hydroxide and concentration of the ninhydrin positive fractions gave a brownish powder. The powder was subjected to a silica gel TLC with BuOH - AcOH - H₂O (2:1:1). The extract from the Rf 0.72 fraction was concentrated to give leucine as a colorless powder.

Determination of Configuration of Leucine

The solution of leucine obtained from the oxidation of leuhistin was examined by HPTLC on a Merck CHIR pre-coated plate eluting with MeOH - H_2O - MeCN (1:1:4). The configuration of the leucine, Rf 0.69, was determined by comparison with authentic D- and L-leucine which gave Rf values of 0.59 and 0.69, respectively.

X-Ray Diffraction Analysis

A colorless prism crystal of leuhistin having approximate dimensions $0.2 \times 0.1 \times 0.02$ mm was mounted on a glass fiber. All measurements were made on a Rigaku AFC5R diffractometer with graphite monochromated CuK α radiation and a 3KW rotating anode generator. The lattice constants were derived from a least-squares refinement using the setting angles of 25 carefully centered reflections in the range $31.0^{\circ} < 2\theta < 51.6^{\circ}$. Crystal data: C₁₁H₁₉N₃O₃, monoclinic, P2₁, a=5.6163(4), b=23.640(2), c=9.639(1)Å, β =96.020(8)°, U=6473(2)Å³, Z=4, D_{cale}=1.259 g/cm³, μ for CuK α radiation=7.74 cm⁻¹.

Intensities were measured by a $2\theta - \omega$ scan method with a scan speed 8°/minute in ω . Backgrounds were measured at each end of the scan for half the total scan time. The weak reflections (I<10.0 σ (I)) were rescaned (maximum of 2 rescans) and the counts were accumulated to assure good counting statistics. A total of 2,087 reflections in the 2θ range $6^{\circ} - 120^{\circ}$ was measured.

The crystal structure was determined by direct method using MAGEX³⁾ program in MITHRIL package⁴⁾. In the final refinement, the non-hydrogen atoms were refined anisotropically by full-matrix least-squares. The hydrogen atoms were eventually included in calculated positions but were not refined. The final R value was 0.045 for 1,022 observed reflections (I > 3σ (I)). The atomic parameters, bond lengths and angles have been sent to the Cambridge Crystallographic Data Centre.

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