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STRUCTURE ELUCIDATION OF SCH 49088, A NOVEL EVERNINOMICIN ANTIBIOTIC CONTAINING AN UNUSUAL HYDROXYLAMINO-ETHER SUGAR, EVERHYDROXYLAMINOSE

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> Schering-Plough Research Institute, 2015 Galloping Hill Road, Kenilworth, N.J. 07033, U.S.A. [†]Paul M. Gross Chemical Laboratories, Duke University, Durham, N.C. 27706, U.S.A. (Dedicated to the memory of the late Professor Sir Derek Barton) Received 17 August 1998; accepted 8 September 1998

Abstract: Degradation of Sch 49088 (1) provided a novel, highly functionalized O-substituted hydroxylamine-containing sugar, everhydroxylaminose 11 isolated as a β -OMe glycoside. Notably the two acid sensitive orthoester moieties in the intact antibiotic 1 co-exist in the presence of three phenolic groups and a free carboxylic acid group. © 1998 Elsevier Science Ltd. All rights reserved.

Ziracin 2, a member of the everninomicin group of antibiotics produced by *Micromonospora Carbonacea*, is undergoing extensive clinical trials for its activity against methicillin resistant *Staphylococci* and vancomycin resistant *Enterococci.*¹ From the same fermentation, the nitroso 3, hydroxylamino 4, and amino 5 compounds have been isolated.² More recently we have isolated from this fermentation a polar, new antibiotic Sch 49088 (1),³ as a white, amorphous solid, C₈₈H₁₂₃NO₃₉Cl₂ (M+Na⁺ 1910.6904, calcd. 1910.6947),⁴ $[\alpha]_D^{25}$ -44.4 (methanol). As in ziracin 2,⁵ the ¹³C-nmr spectrum of 1⁶ indicated the presence of two orthoester carbons at δ 119.2 and δ 120.4. However, besides the absence of a nitro group in the infrared spectrum, there were other marked differences in the mass and nmr spectra of 1 compared to 2. These included a signal at δ 175.8 in 1 suggestive of a carboxylic acid or carboxamide carbon; also observed was an upfield shift of quarternary C65 from δ 89.9 in 2 to δ 59.8 in 1.⁷ Two sets of fragment ions of interest in the (HRFAB) mass spectrum of 1 at 954.3810 and 935.3396 suggested cleavage at the C16 orthoester as has been established for other everninomicins.¹ Fragmentation ions in the spectrum of 1 also showed identical sugar sequences from rings **A** to **J** as found in **2-5**. A new fragment peak at m/z 273.1860 (C₁₈H₂₅O₂) in 1 suggested its connectivity to ring **K** possibly through the quaternary carbon 65.

Treatment of 1 with excess diazomethane gave 1a (R = R' = Me) adding four methyl groups, three as methyl ethers and one as a methyl ester.⁸ The presence of three phenolic groups on rings A and I may account for the addition of three of the methyl groups as methyl ethers. In the (HRFAB) mass spectrum of 1a, a fragment peak at m/z 287.2012 (C₁₉H₂₇O₂) replaced the above m/z 273 fragment indicating addition of the remaining methyl functionality on the ring K side chain. Based on the well precedented Ganguly-Sarre degradation sequence,¹ two-phase acid hydrolysis of 1a in ethyl acetate followed by treatment of the intermediate ester 6 with diazomethane in methanol gave the lactone 7 and compound 8 as major products, the latter being identical in all respects to an authentic sample obtained from ziracin 2^{1b} in the same manner. The lactone 7, which contained the elements responsible for differences with ziracin 2, is an amorphous solid, C₄₉H₆₉NO₁₅Cl₂ (M+H⁺ 982.4113, calcd. 982.4123), [α]_D²⁵ -46.9 (CHCl₃). As in 1a, the mass spectrum of 7 had an important fragment peak at m/z 287.2012 (C₁₉H₂₇O₂) confirming the origin of the fragment ion m/z 273 derived from the parent antibiotic 1. Detailed nmr studies of 7^{6,9} revealed that the C₁₉H₂₇O₂ side chain consists of a trisubstituted phenyl ring with a C₇H₁₄ alkyl, a methyl and a *trans*-1,2-disubstituted double bond connected to a methylene carbonyl-methoxy unit.

Further degradation of **7** in refluxing methanol in the presence of p-toluene sulfonic acid provided a lipophilic monosugar α -methyl glycoside **9** C₂₈H₄₅NO₆ (M+H⁺ 492.3311, calcd. 492.3325), [α]²⁵_D +11.1 (CHCl₃) and its β -

0040-4039/98/\$ - see front matter © 1998 Elsevier Science Ltd. All rights reserved. *PII*: S0040-4039(98)01923-6 anomer 10 (M+H⁺ 492.3311), $[\alpha]_D^{25}$ -45.8 (CHCl₃) as thick syrups in a 1:2 ratio.¹⁰ Acid hydrolysis of 1 under identical conditions provided the same α - and β - OMe glycosides 9 (63-C<u>H</u>, dd, δ 4.4, J = 5.1, 1.7 Hz) and 10 (63-C<u>H</u>, d, δ 4.64, J = 4.5 Hz) respectively.¹⁰ Their detailed mass and nmr spectral analysis confirmed that the C₁₈H₂₅O₂ fragment was attached to ring K through the nitrogen at the quaternary C-65, with all of the structural features mentioned above remaining intact during the degradation.

The connections between all subunits in 9 and 10, especially the unusual C-N-O-C linkage, were elucidated by a series of selective INEPT experiments, which showed that the O-methyl at C-88, C-87 CH, and the C-86 CH are all long-range coupled to C-88. These protons are either two or three bonds from C-88. The C-71 CH was found to be two or three bonds from quaternary C-72 and C-73, one methine carbon (C-77) in the aromatic region, and two methylene carbons (C-79 and C-80) in the aliphatic region. In addition C-71 CH is not long-ranged coupled to C-65 at 59.7 ppm, which is four bonds away in 9 or 10. Similar connectivities were found in the HMBC⁵ spectrum of 1.

Methylation of **10** with NaH/CH₃I in tetrahydrofuran yielded gem-dimethylated **12** C₃₀H₄₉NO₆ (M+H⁺ 520.3648, calcd. 520.3639). A doublet of doublets at 3.24 ppm (2H) in the pmr spectrum of **10** was replaced by a singlet at 1.46 ppm (6H) in **12** indicating a methylene spacer between the C-86 and C-88 in the former. Hydrogenation of **10** (Pd/C) gave the dihydro sugar **13** C₂₈H₄₇NO₆ (M+H⁺ 494.3458, calcd. 494.3481).

Basic hydrolysis of **10** (KOH/dioxane/H₂O) at room temperature provided everhydroxylaminose, the novel sugar carboxylic acid **11** C₂₇H₄₃NO₆ (M+H⁺ 478.3159, calcd. 478.3169), $[\alpha]_D^{25}$ -53.3 (CHCl₃), as its β-OMe glycoside.¹¹ Treatment of **11** with diazomethane regenerated **10** confirming the presence of a methyl ester at C-88 and therefore a carboxy group in **11** and **1**.

As expected, reductive N-O bond cleavage of both **9** and **10** in methanolic alkaline solutions with nickel/aluminium alloy^{12,13} conveniently provided the corresponding amino sugars **15** and **16** and the lipophilic aromatic acid **14**, as waxy crystals, m.p. $50-51^{\circ}$ C, C₁₈H₂₁NO₄ (M+Na⁺ 315.1933, calcd. 315.1936), $[\alpha]_D^{25}$ -16.9 (CHCl₃).¹⁴ Acetylation of the α -anomer **15** with acetic anhydride yielded the crystalline **17** m.p. 158-159°C, C₁₁H₂₁NO₄ (M+Na⁺ 254.1370, calcd. 254.1368), $[\alpha]_D^{25}$ +60.5 (CHCl₃), which was identical in all respects (including single crystal X-ray data)¹⁵ to the same compound isolated from evernitrose.^{1b} Thus the stereochemistry of the new sugar, everhydroxylaminose **11** (isolated as β -OMe glycoside) at C65, C66 and C67 could be assigned in a relative as well as an absolute sense.¹¹

We have reported on the determination of the absolute stereochemistry at the C16 orthoester of the everninomicin antibiotics.^{1a} As can be seen from a comparision of ¹H and ¹³C chemical shifts of 1 and 2 (rings **B**, **C** and **D**), there can be little doubt that the same connectivity and stereochemical relationship exists at C16 in the structure of 1. (Table)

In summary, Sch 49088 (1) is a novel everninomicin antibiotic containing a unique O-alkylated hydroxylamine substituted sugar subunit. In *in vitro* antibacterial assays against *Staphylococci* as well as *Enterococci*, **1** was found to be only weakly active compared to ziracin (2), 3, 4 and 5.^{1,2}

References and Notes:

- (a) A. K. Ganguly, J. L. McCormick, T-M. Chan, A. K. Saksena and P. R. Das, *Tetrahedron Lett.*, 1997, <u>38</u>, 7989; (b) A. K. Ganguly, B. Pramanik, T-M. Chan, Y.T. Liu, J. Morton and V. Girijavallabhan, *Heterocycles*, 1989, <u>28</u>, 83; (c) V. M. Girijavallabhan and A. K. Ganguly, *Kirk-Othmer Encyclopedia of Chemical Technology*, <u>1992</u>, 4th ed., Vol. 3, 259.
- 2. A. K. Ganguly. V. M. Girijavallabhan, G. H. Miller and O. Z. Sarre, J. Antibiotics, 1982, 35, 561.
- 3. Isolation of Sch 49088: The fermentation broth² was adjusted to pH 10.5 and centrifuged to remove mycelia; the aqueous phase was adjusted to pH 7 and the resulting precipitate collected by filtration and vacuum dried at room temperature. A 40 g portion of this crude powder was chromatographed over silica gel (3.2 Kg). The column was eluted with 20% acetone/CH₂Cl₂ (28 L), 30% acetone/CH₂Cl₂ (40 L), acetone (28 L) and 5% H₂O/acetone (20 L). Fractions containing Sch 49088 (100% acetone and 5% H₂O/acetone) were combined and evaporated to dryness *in vacuo* to provide almost pure (TLC)material (18.3 g). A 7 g portion of this product was rechromatographed over silica gel (300 g) using 4-5% methanol/methylene chloride as



78 Me

75

<u>14</u> R"=H

3.82

1.35

1.34

68.80

18.80

18.46

ÕMe <u>**15**</u> $R^1 = OMe; R^2 = H; Z = H$ <u>**16**</u> $R^1 = H; R^2 = OMe; Z = H$ **<u>17</u>** $R^1 = OMe; R^2 = H; Z = Ac$

ö

3.91 78.45

*δ (± 0.1 ppm); **δ (± 0.02 ppm)

3.82

1.35

1.34

68.76

18.76

18.55

28

eluent. Fractions (100 mL each) were collected and monitored by TLC (SiO2, 10% methanol/CH2Cl2). Fractions 15-18 were combined and evaporated in vacuo to provide a white amorphous solid which was dissolved in a minimum of ethyl acetate and precipitated with efficient stirring into n-heptane. Pure Sch 49088 was dried in vacuum and obtained as a white powder (3.8 g). To prevent decomposition this material was stored as a dry powder at -40°C. FAB mass spectra were obtained on the JEOL JMS-HX110A instrument equipped with a FAB gun using xenon gas. The use of

- 4 DMSO as the solvent and 3NBA as the matrix with NaCl addition gave the best results producing abundant sodiated molecular ions. Negative ion FAB-MS analysis was performed in the DMSO-SNBA matrix without the addition of NaCI. High resolution fragment ions spectra (positive and negative FAB) were utilized to identify sugar sequences. The peak matching experiments were performed at a mass resolving power of 5000. Detailed mass and nmr spectral discussion will be presented elsewhere. T-M. Chan, R. M. Osterman, J. B. Morton and A. K. Ganguly, *Magnetic Resonance in Chemistry*, **1997**, <u>35</u>, 529.
- 5
- NMR spectra were obtained on a Varian XL-400 instrument, operating at a frequency of 400 MHz for ¹H and 100.6 MHz for 6 13C. Complete NMR assignments are based on COSY, NOESY, HETCOR, HMBC and SINEPT experiments according to reference 5. Selective spectral data is given here.
- Numbering system follows according to reference 1c and has been used consistently including the degradation products except 7. the amino sugars 15, 16 and the acetamido derivative 17.
- All degradation experiments were conducted with 1a due to its improved solubility and stability to storage compared to 1. 8
- Lactone 7: ¹H NMR [CDCk] & 0.80 (m, 70-CH3, 83-CH3, 84-CH3), 1.00 and 1.42 (m, 80-CH2A, 80-CH2B), 1.05 (s, 68-CH3), 9 1.10 and 1.25 (m, 82-CH2a, 82-CH2B), 1.29 (m,81-CH), 1.41 (d, 21-CH3), 1.42 (d, 15-CH3), 1.50 and 2.10 (d and dd 64-CH2a, 64-CH2B), 1.64 and 1.72 (m, 79-CH2A, 79-CH2B), 1.76 and 2.39 (m, 11-CH2A, 11-CH2B), 2.31 (s, 78CH3), 2.36 (s, 8-CH3), 2.56 and 3.09 (dd 17-CH2A, 17-CH2B), 3.02 (d, 66-CH), 3.22 (dd, 87-CH2), 3.32 (dd, 19-CH), 3.43 (s, 69-OCH3), 3.48 (m, 67-CHO), 3.60 (m, 14-CHO), 3.70 (s, 88-COOCH3), 3.87 (s, 7-OCH3) 3.91 (s, 1-C-OCH3), 3.92 (m, 12-CHO), 3.99 (m, 18-CHO), 4.18 (m, 20-CHO), 4.21 (s, exchangeable), 4.59 (dd, 10-CHO), 4.71 (t, 71-CHO), 4.86 (d, 63-CHO), 4.94 (t, 13-CHO), 5.14 (broad, exchangeable), 6.08 (dt, 86-CH), 6.80 (d, 85-CH), 7.01 (d, 75-CH), 7.03 (s, 77-CH), 7.32 (d, 74-CH). ¹³C NMR [CDCl₃] CH₃ δ 11.4 (C-83), 17.9, 18.1, 18.2, 18.3, 18.8, 19.2 (C-84), 21.4; CH₂ 29.3 (C-82), 32.7, 33.4, 36.1, 36.5, 38.0, 38.6 (C-87); CH 34.4 (C-81; OCH3) 51.9, 60.7, 61.3, 62.0; tert-C 59.6 (C-65); CHO 66.0 (C-67), 67.3 (C-18), 71.5, 71.7, 74.4, 75.7 (C-13), 82.3 (C-66), 83.3, 86.0 (C-71, C-14), 94.2 (C-63), 101.0 (C-10); tert-C 121.3, 122.7 (C-86), 125.7, 126.2, 126.6 (C-74), 127.3 (C-77), 128.1 (C-75), 131.1 (C-85), 132.6, 135.3, 139.5, 153.3 (C-1/C-3), 154.7 (C-1/C-3), 165.6 (C-9/C-16), 172.0 (C-88).
- Isolation of 9 and 10: A solution of the lactone 7 (0.34 g) in dry methanol (30 ml) containing p-toluene sulfonic acid (0.018 g) was refluxed for 18 hrs. After cooling NaHCO3 (0.025 g) was added, the reaction mixture stirred for -30 min and evaporated to dryness in vacuo. The residue was dissolved in minimum CH2Cl2 and chromatographed over silica gel (50 g). The column was successively eluted with n-hexane and increasing concentration of ethyl acetate (5%, 8%, 10%, 12%) in n-hexane. Appropriate fractions were combined (TLC) to provide the less polar 10 (β-anomer, 0.099 g) and the more polar 9 (α-anomer, 0.056 g). β-anomer 10: ¹H NMR [CDCl3] δ 3.71 (s, 88-OCH3), 3.24 (dd, J = 7.2, 1.5 Hz; 87-CH2), 6.08 (dt, 15.5, 7.2, 7.2 Hz; 86-CH), 6.84 (broad d, J = 15.5 Hz; 85-CH), 7.34 (d, J = 7.8 Hz; 74-CH), 7.08 (broad s, 77-CH), 7.03 (dd, J = 7.8, 1.8 Hz; 75-CH2), 2.34 (s, 78-CH3), 4.75 (dd, J = 7.4, 5.9 Hz; 71-CH), 1.65, 1.78 (m, 79-CH2), 1.05, 1.45 (m, 80-CH2), 1.3 (m, 81-CH), 1.1, 1.3 (m 82-CH2), 0.81 (t, J = 7.2; 83-CH3), 0.82 (t, J = 6.8 Hz; 84-CH3), 4.64 (d, J = 4.5 Hz; 63-CH), 2.15 (dd, J = 4.5, 13.8 Hz; 64-CH2A), 1.56 (d, J =13.8 Hz; 64-CH2B), 3.16 (d, J = 9.8 Hz; 66-CH), 3.67 (m, 67-CH), 1.28 (d, J = 6.1 Hz; 70-CH3), 1.1 (s, 68-CH3), 3.54 (s, 69-OCH3), 3.28 (s, 63-OCH3).

13C NMR [CDCl3] & 51.8 (C-88-OCH3), 172.0 (C-88), 38.5 (C-87), 122.6 (C-86), 131.1 (C-85), 132.5 (C-73), 139.4 (C-72), 126.5 (C-74), 127.2 (C-77), 128.0 (C-75), 137.2 (C-76), 21.3 (C-78), 83.1 (C-71), 33.3 (C-79), 32.7 (C-80), 34.4 (C-81), 29.2 (C-82), 11.3 (C-83), 19.1 (C-84), 98.7 (C-63), 38.1 (C-64), 59.7 (C-65), 82.8 (C-66), 65.2 (C-67), 18.5 (C-70), 18.1 (C-68), 61.4 (C-69), 54.6 (C-63-OCH3).

- 11. Everhydroxylaminose β-methyl glycoside 11: Except for the absence of C-88-OCH3, the ¹H and ¹³C NMR spectra of 10 and 11 were virtually identical. At this stage we do not know the absolute stereochemistry at C-71 and C-81. Early attempts to provide a crystalline derivative of either 11 or 14 failed to provide crystals suitable for X-ray analysis.
- 12. G. Lunn, E. B. Sansone, L. K. Keefer, Synthesis, 1985, 1104.
- 13. Ni-Al alloy reduction of 9: Ni-Al alloy (3.5 g) was added in portions to a solution of 9 (0.35 g) in methanol (20 ml) and IN KOH (14 ml). The reaction mixture was stirred at room temperature for ~15 hrs. Additional Ni-Al alloy (1 g) was added and the mixture stirred for another 8 hrs. At the end of this time the reaction mixture was filtered through celite to remove inorganic salts and washed with methanol/water (1:1, ~10 mL). The combined filtrates were evaporated to dryness in vacuo and residue taken up in water (~15 mL, pH ~14). The aqueous solution was extracted with CH₂Cl₂ (2 x 10 mL), the organic phase dried (Na₂SO₄) and evaporated to provide the amino sugar 15 (α -anomer; 0.106 g) as a thick syrup. The remaining aqueous phase was adjusted to pH 2 by addition of citric acid and extracted with ethyl acetate (2 x 10 mL). The ethyl acetate phase was washed with brine (~5 mL), dried (Na2SO4) and evaporated to dryness in vacuo to provide the aromatic acid 14 (0.249 g)as an oil. Similar treatment of 10 with Ni-AI alloy provided the amino sugar 16 (β-anomer) and the same aromatic acid 14.
- 14. Aromatic acid 14: ¹H NMR [CDCl3] δ 2.42 (t, J = 7.2 Hz; 87-CH2), 1.92 (quintet, J = 7.6 Hz; 86-CH2), 2.67 (m, 85-CH2), 7.29 (broad s, 74-CH), 7.04 (d, J = 7.8 Hz; 77-CH), 7.01 (dd, J = 1.5, 7.8 Hz; 75-CH), 2.33 (s, 78-CH3), 4.89 (dd, J = 7.6, 5.5 Hz; 71-CH), 1.72 (m, 79-CH2), 1.1, 1.55 (m, 80-CH2), 1.45 (m, 81-CH), 1.15, 1.45 (m, 82-CH2), 0.85 (t, J = 7 Hz; 83-CH3), 0.86 (d, J = 6.7 Hz; 84-CH3)

13C NMR [CDCl3] δ 178.8 (C-88), 33.4 (C-87), 26.4 (C-86), 31.2 (C-85), 134.8 (C-73), 142.4 (C-72), 129.5 (C-74), 126.3 (C-77), 128.1 (C-75), 136.2 (C-76), 21.1 (C-78), 70.7 (C-71), 36.3 (C-79), 33.1 (C-80), 34.4 (C-81), 29.3 (C-82), 11.3 (C-83), 19.2 (C-82). Note: Concomitant reduction of C85, C86 double bond could not be avoided under the N-O bond cleavage conditions.

Atomic parameters bond lengths, bond angles and torsion angles for 17 have been deposited at the Cambridge Crystallographic Data Center, Cambridge, CB2 1EZ, U.K. Any request should be accompanied by full literature citation for this communication.