

Synthetic Studies on Tetrazomine: Stereochemical Assignment of the β -Hydroxypipelicolic Acid

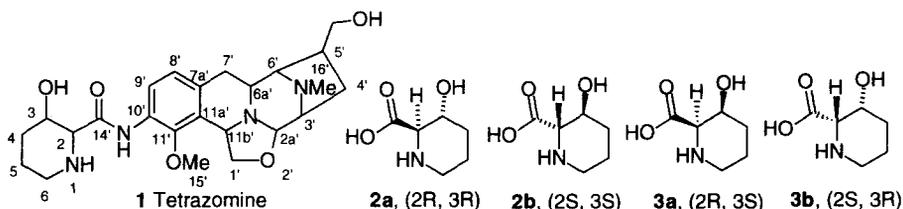
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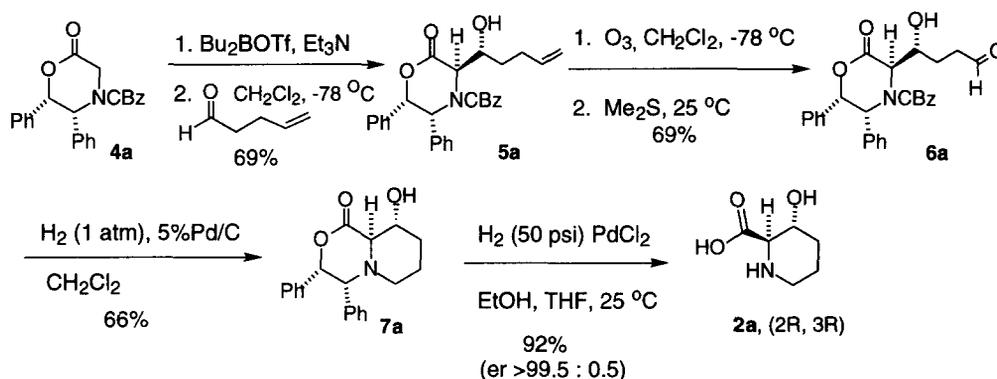
Abstract. The asymmetric syntheses of 2(R), 3(R)- and 2(S), 3(S)- β -hydroxypipelicolic acids are described, including the determination of the absolute stereochemistry of the β -hydroxypipelicolic acid moiety of tetrazomine. © 1998 Elsevier Science Ltd. All rights reserved.

Tetrazomine (**1**), is an antitumor antibiotic that was isolated from *Saccharothrix mutabilis*¹ by the Yamanouchi Pharmaceutical Co. in Japan. Preliminary antitumor/antimicrobial assays of this substance indicate that tetrazomine displays potent antitumor activity against P388 leukemia *in vivo* and displays good antimicrobial activity against both Gram-negative and Gram-positive organisms. Tetrazomine is structurally similar to the quinocarcin and bioxalomycin class of antitumor antibiotics and the mechanism of oxidative DNA cleavage mediated by tetrazomine has been reported from these laboratories.² Tetrazomine contains the unusual amino acid β -hydroxypipelicolic acid. The relative and absolute stereochemistry of this amino acid and that of the polycyclic framework of tetrazomine have not yet been determined. Here we report the determination of the absolute stereochemistry of the β -hydroxypipelicolic acid unit of tetrazomine.

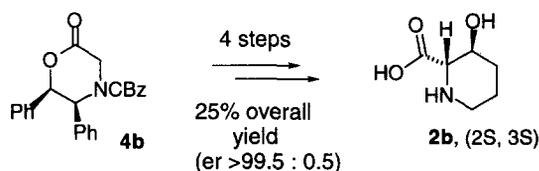


In connection with studies on the structure elucidation, total synthesis³ and mechanism of action of tetrazomine, we required stereochemically unambiguous syntheses of 2(R),3(R)- and 2(S),3(S)- β -hydroxypipelicolic acids **2a** and **2b** (Figure 2). The synthesis of the 2(R),3(S)-stereoisomer **3a** has been completed by two groups^{4,5} while the 2(S),3(R)- β -hydroxypipelicolic acid stereoisomer **3b** has not been previously reported. Several groups have recently reported syntheses of **2a** and **2b**⁶ and in this account, we describe an efficient asymmetric synthesis of the two *anti*-isomers of β -hydroxypipelicolic acid.

As shown in Scheme 1, the commercially available lactone **4a** (Aldrich) was converted into the corresponding boron enolate with di-*n*-butyl boron triflate.⁷ Diastereoselective aldol condensation with 4-pentenal (Lancaster) provided the desired *anti*- β -hydroxy aldol product **5a** in 69% yield. Ozonolysis of the olefin furnished the aldehyde **6a**.⁹ Mild catalytic hydrogenation of **6a** afforded bicyclic **7a**¹⁰ via sequential N-CBz deprotection and reductive amination. Finally, the amino acid 2(R),3(R) β -hydroxy pipercolic acid **2a**¹¹ was produced through catalytic hydrogenation over palladium-black.



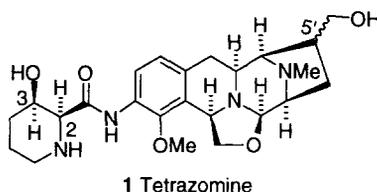
The corresponding 2(S),3(S) β -hydroxy pipercolic acid **2b** was synthesized in the same manner using the commercially available lactone **4b** in the aldol condensation (Scheme 2). Both enantiomers of β -hydroxy pipercolic acid were produced with enantiomeric ratios (er) of >99.5 : 0.5. The synthesis reported here is five steps from commercially available **4** with an overall yield of 25-28% and compares very favorably with existing syntheses reported in the literature which range from 9-15 steps in overall yields less than 25%.^{5,6}



To determine the relative and absolute stereochemistry of the β -hydroxy pipercolic acid of tetrazomine the amide of natural tetrazomine (Yamanouchi Pharmaceutical Co.) was subjected to hydrolysis. The hydrolysis product was compared to synthetic amino acids **2a**, **2b**, and **3a**.¹² Unfortunately, under basic hydrolysis conditions (2M LiOH, reflux 6 h) both *syn*- and *anti*- β -hydroxy pipercolic acids were detected in the crude hydrolysate by HPLC analysis indicating epimerization had occurred. Under acidic hydrolysis conditions, (4M HCl, 80 °C, 20 h) the only β -hydroxypipercolic acid detected by HPLC possessed the *syn*-relative stereochemistry. This amino acid was isolated from the hydrolysate by reverse-phase HPLC (20% MeOH/H₂O, Waters Resolve column, Waters 600 HPLC, uv 210nm) and was found to have physical data

(¹H-NMR, FAB HRMS) identical to that of the *syn*-β-hydroxy pipercolic acid **3a** which was prepared according to the procedure of Knight, et al., via Baker's yeast reduction of N-t-BOC 3-keto methyl pipercolate.¹² However, the β-hydroxy pipercolic acid obtained from the tetrazomine hydrolysis exhibited an optical rotation with the opposite sign to that of the authentic, synthetic sample of **3a**.¹³ Thus, the β-hydroxy pipercolic acid moiety of tetrazomine must correspond to **3b** with the (2*S*, 3*R*)-configuration.

Based on the structural similarity of tetrazomine to quinocarcin and the bioxalomycins, biogenetic and stereoelectronic² considerations lead us to propose an almost complete stereostructure for tetrazomine as depicted below. The only stereogenic center that is still reasonably suspect, is C5' which differs in relative configuration between quinocarcin and the bioxalomycins.



Efforts to elucidate the complete stereostructure of tetrazomine as well as defining the role that the β-hydroxy pipercolic acid unit plays in DNA recognition, transport and biological activity are under study in this laboratory.

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References and Footnotes

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4. Knight, D.W.; Lewis, N.; Share, A.C.; Haigh, D., *Tetrahedron Asymmetry* **1993**, *4*, 625~628. This paper does not describe the synthesis of the free amino acid but the protected version: N-t-BOC (2*R*, 3*S*)-3-hydroxy-methyl pipercolate which has been converted to **3a** (this work, ref. 11).
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8. **5a**: ¹H-NMR (300 MHz, d₆-DMSO vs TMS 393K) 1.69-1.85 (2H, m), 2.14-2.28 (2H, m), 4.17 (1H, m), 4.84 (1H, d, J=2.25 Hz), 4.95-5.18 (4H, m), 5.25 (1H, d, J = 3.18Hz), 5.47 (1H, d, J= 5.34 Hz), 5.82-5.88 (1H, m), 6.53 (1H, d, J = 3.16 Hz), 6.60 (2H, d, J=7.24 Hz), 6.99-7.39 (13 H, m). Anal calcd. for C₂₉H₂₈NO₅: C, 73.87; N 2.97; H 6.23 found: C, 74.04; N,2.85; H, 6.14. $[\alpha]_D^{20} = +13.7$ (c=1.3, CHCl₃). **5b**: $[\alpha]_D^{20} = -12.4$ (c=1.1, CHCl₃).

9. **6a**: ¹H-NMR (300MHz, d₆-DMSO vs TMS, 393 K) δ 2.04 (2H, m), 2.29 (2H, m), 4.53 (1H, m), 4.99 (1H, d, J = 2.52 Hz), 5.03 (4H, m), 5.24 (2H, m), 6.71 (1H, d, J = 3.24 Hz), 7.02 (15H, m) 9.70 (1H, s). HRMS (FAB) calcd. for C₂₈H₂₈NO₆ (MH⁺): 474.1917. found: 474.1914. $[\alpha]_D^{20} = + 3.1$ (c = 1.25, CHCl₃). **6b**: $[\alpha]_D^{20} = - 3.7$ (c = 1.25, CHCl₃).

10. **7a**: ¹H-NMR (300MHz, CDCl₃ vs TMS) δ 1.20 (1H, m), 1.70 (2H, m), 2.06 (2H, m), 2.86 (1H, d, J=11.4Hz), 3.01 (1H, d, J=8.7Hz), 3.96 (1H, ddd, J=2.1, 3.9, 4.8Hz), 4.24 (1H, d, J=3.9Hz), 4.9 (1H, s, D₂O exchangeable), 6.17 (1H, d, J=3.9Hz), 7.02-7.30 (10H, m). ¹³C-NMR (75.47 MHz) δ 22.89, 32.42, 52.04, 62.95, 66.02, 62.95, 66.02, 69.11, 83.29, 125.3, 125.4, 127.6, 128.0, 128.2, 129.9, 130.7, 135.6, 172.1. HRMS (FAB) calcd. for C₂₀H₂₂NO₃ (MH⁺): 324.1600. found: 324.1586. $[\alpha]_D^{20} = +21.4$ (c = 1.9 CHCl₃) **7b** $[\alpha]_D^{20} = -20.9$ (c = 1.6 CHCl₃).

11. **(2R,3R)-β-Hydroxypipelicolic acid (2a)**. ¹H-NMR (300 MHz, D₂O) δ 1.55 (2H, m), 1.81 (2H, m), 2.91 (1H, m), 3.18 (1H, m), 3.44 (1H, d, J = 6.96 Hz), 3.97 (1H, m). HRMS (FAB) calcd. for C₆H₁₂NO₃ (MH⁺): 146.0817, found 146.0822. The optical purity of the final amino acids **2a** and **2b** were determined to be > 99.5 : 0.5 er by chiral HPLC analysis (Daicel Chiral Pak WH, column temperature 40 °C using 0.25 mM CuSO₄ on a Waters 600 HPLC with a UV detector at 210 nm).

12. **(2R,3S)-β-Hydroxypipelicolic acid (3a)**. This amino acid was synthesized using the Baker's yeast method of Knight, et al., (see ref. 4). ¹H-NMR (300 MHz, D₂O) δ 1.85 (2H, m), 2.12 (2H, m), 3.07 (1H, ddd, J= 12.9, 3.6, 2.7 Hz), 3.47(1H, ddd, J= 12.6, 2.1, 1.8 Hz), 3.75 (1H, d, J = 1.5 Hz), 4.57 (1H, s, broad). ¹³C-NMR (75.47 MHz vs. d₄ MeOD) δ 16.89, 29.72, 44.56, 63.20, 65.04, 173.06. HRMS (FAB) calcd. for C₆H₁₂NO₃ (MH⁺) 146.0817 found: 146.0821. The final product reported by Knight, et al., (ref. 4) N-t-BOC (2R, 3S)-3-hydroxy-methyl pipicolate, was converted to the free amino acid by sequential treatment with TFA in CH₂Cl₂ followed by removal of the solvent and treatment of the amino methyl ester with KOH in water at room temperature. Acidic Dowex-50WX2-100 ion exchange resin treatment (eluted within 2% ammonium hydroxide) afforded the free amino acid **3a** (81% for two steps). To our knowledge, physical data for the free amino acid (**3a**) have not been previously reported .

13. The isolated amino acid (**3b**) exhibited an $[\alpha]_D^{20} = -72.3^\circ$ (c=0.10, 1M HCl) whereas, **3a** (see ref. 12) exhibited an $[\alpha]_D^{20} = +82.1^\circ$ (c=0.12, 1M HCl).