the formation of the pyruvate adduct of TDP and the substitution process in which a proton replaces CO2 to generate HETDP should be stereospecific¹⁷ as is the case for the El subunit of pyruvate dehydrogenase.^{5,14} The stereochemistry of these processes in enzymic and nonenzymic systems is the subject of ongoing studies.

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Supplementary Material Available: Listing of crystal data and tables of final atomic positional and thermal parameters for HETI (1 page). Ordering information is given on any current masthead page.

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Streptonigrin Biosynthesis. 8. Evidence for the Involvement of a New Shikimate Pathway Product and a New Route to Quinolines¹

W. Randal Erickson and Steven J. Gould*2

Department of Chemistry, Oregon State University Corvallis, Oregon 97331 Received September 15, 1986

We have previously reported data^{3,4} suggesting that a 4aminoanthranilic acid (1), D-erythrose-4-phosphate (2), and β methyltryptophan⁵ (3) are the key precursors in biosynthesis of the anticancer antibiotic streptonigrin (4). As shown in Scheme I, these can be combined in sequences that lead either to a 7aminoquinoline-2-carboxylic acid 5 (pathway A) or to a β -carboline 6 (pathway B) as the pivotal intermediate. We now report that pathway A is operative with 4-aminoanthranilic acid (1a) and 7-aminoquinoline-2-carboxylic acid (5a) as intermediates and that all three A-ring oxygenations occur at a later stage.

A fermentation in the presence of ¹⁸O₂ gas had yielded streptonigrin, labeled-among other positions-at C-5 and C-6 but not at C-8, suggesting that the C-8 oxygen was retained from a prearomatic precursor and that the hydroxylated compounds 1b and 5b were likely intermediates.⁶ However, we recognized that because C-8 is the carbonyl of a vinylogous ester, an oxygen atom may have been introduced by a metabolic oxidation but subsequently lost by exchange,^{7,8} either with the fermentation medium or during extractive workup. Indeed, when samples of authentic streptonigrin were stirred overnight in solutions of THF/H₂¹⁸O at pH 5.0 and at pH 10.5 and then reisolated and analyzed by ¹³C NMR, it was found that ¹⁸O had been incorporated to the extent of 15% and 30%, respectively, exclusively at C-8. Thus, neither the exact origin of the C-8 oxygen nor the oxidation level of the putative aromatic precursor(s) were certain at this point.

[4-15N]4-Aminoanthranilic acid (1c) was then synthesized in three steps (22% overall yield)9 utilizing H15NO3 (99% enriched).10

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A sample of the sodium salt of 1c in 0.05 M pH 8.5 phosphate buffer was added under sterile conditions to shaken fermentations¹¹ of Streptomyces flocculus. Pulse feedings of 61.1 mg/15 mL, 20.8 mg/10 mL, 20.4 mg/10 mL, and 20.2 mg/10 mL were divided among the flasks $(3 \times 500 \text{ mL})$ at 24, 36, 48, and 60 h, respectively. At the termination of the fermentation, standard workup afforded 27.6 mg of pure 4a. The ¹⁵N NMR spectrum of $4a^{12}$ exhibited a single resonance at 73.6 ppm¹³ that is attributable to the C-7 amine nitrogen of 4.5 Although the specific enrichment could not be calculated because of NOE due to proton decoupling, no resonance was detectable for the unenriched C-5' amine nitrogen.14

[4-²H]7-Aminoquinoline-2-carboxylic acid (5c) was next synthesized from the quinoline 7^{15} as shown in Scheme II. Reductive removal of chloride from 8 with deuterium gas afforded the labeled ester 9, and mild hydrolysis gave the amino acid 5c in 25% overall vield.

The sodium salt of 5c was fed by dividing pulses of 47.0, 44.5, 38.2, and 42.4 mg, each in 15 mL of buffer, among three 500-mL cultures at 28, 38, 48, and 58 h after inoculation, respectively. Standard workup afforded 20.6 mg of pure **4b** which was analyzed by ²H NMR.¹⁶ A singlet at $\delta 8.23^{17}$ was observed corresponding





to a deuterium label at C-4. By comparison with the natural abundance deuterium signal for solvent Me₂SO (also employed as internal chemical shift reference), incorporation was determined to be 1.4%.

On the basis of these data it appears that streptonigrin is biosynthesized via pathway A with R = H (Scheme I), and unless there is a metabolic grid, it is unlikely that 1b is also an intermediate. The evidence suggests that compound 1a represents a new metabolite of the shikimate pathway,19 while the involvement of 5a reveals a fundamentally new biosynthetic pathway to the quinoline ring system.²⁰ This may be viewed (Scheme III) as

(14) In earlier work both the C-7 amine and C-5' amine peaks were of equal intensity in a natural abundance ¹⁵N NMR spectrum. See ref 5. The C-7 amine nitrogen was also observed in the enriched and natural abundance samples by using a refocused decoupled INEPT sequence. This gave a sig-nal-to-noise ratio approximately 3 times greater than that of the standard experiment described in ref 12. The C-5 amine nitrogen was not observed in this case, presumably due to rapid proton exchange eliminating the possibility of efficient polarization transfer.

(15) Heindel, N. D.; Bechara, I. S.; Lemke, T. F.; Fish, V. B. J. Org. Chem. 1967, 32, 4155

(16) Spectrum taken on 10.0 mg in 0.4 mL of Me₂SO with a Bruker AM 400 spectrometer at 61.4 MHz (sweep width = 639 Hz, data points = 2K zero filled to 8K, Hz/pt = 0.16, acquisition time = 1.604 s, pulse width = 45°, no. of scans = 33993).

(17) The previous assignments of H-3 and H-4 were inadvertently reversed.¹⁸ The correct assignments were confirmed by a 2-D $^{13}C/^{1}H$ HETCOR experiment on authentic 4.

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⁽¹⁾ Presented by WRE at the 41st Northwest Regional ACS Meeting, Portland, OR, June 16-18, 1986.

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⁽¹¹⁾ Gould, S. J.; Chang, C. C. J. Am. Chem. Soc. **1980**, 102, 1702. (12) Spectrum taken on 22.0 mg in 0.4 mL of Me_2SO-d_6 with a Bruker AM 400 spectrometer at 40.5 MHz (sweep width = 2778 Hz, data points = 4K zero filled to 8K, Hz/pt = 0.68, acquisition time = 0.737 s, pulse width = 38°, line broadening = 2.0 s, relaxation delay = 1.2 s, no. of scans = 22305)

⁽¹³⁾ Relative to external [15N]aniline, 56.5 ppm, obtained from MSD Isotopes

Scheme I

<u>PATHWAY A</u>



Scheme II



Reagents: a) xs POCl_3, \triangle , 4h b) 3.3 eq SnCl_2+2H₂O, con. HCl, 3h, 0°-25° c) ²H₂, 10% Pd/C, 1.1 eq. KOH, MeOH, 1h , 25° d) 1N NaOH, 0.5h , 25°

Scheme III



analogous to the biosynthesis of tryptophan from anthranilic acid and ribose diphosphate.21

In order to further probe this remarkable pathway, testing of 1b, 7-amino-8-hydroxyquinoline-2-carboxylic acid (5b), and 7-

(20) For examples of quinoline biosynthesis from anthranilic acid and "acetate", see: Herbert, R. B. The Biosynthesis of Secondary Metabolites; Chapman and Hall: New York, 1981; pp 131-132. For examples from tryptophan metabolism, see: Luckner, M. Secondary Metabolism in Microorganisms, Plants, and Animals; Springer-Verlag: New York, 1984; pp 404-406.

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amino-5-hydroxyquinoline-2-carboxylic acid as potential later intermediates is currently under investigation.

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Intermediates in Nucleophilic Aromatic Substitution

Radu Bacaloglu,* Clifford A. Bunton,* and Giorgio Cerichelli

> Department of Chemistry, University of California Santa Barbara, California 93106

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Bimolecular nucleophilic aromatic substitution by anions in polar hydroxylic solvents is generally written as rate-limiting formation of a Meisenheimer, or σ , complex,^{1,2} but π -complexes³ are also postulated reaction intermediates.⁴

Unexpectedly, reported rate constants for formation of Meisenheimer complexes from OH⁻ and a nitroarene or quinazoline

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