

racemate, behaves as a strong, competitive inhibitor of IS, with a K_i value of $0.36 \pm 0.05 \mu\text{M}$ (compare to chorismate $K_m = 7.0 \pm 1.3 \mu\text{M}$). In contrast, **5** is a weak competitive inhibitor of AS (K_i of $195 \pm 15 \mu\text{M}$; chorismate $K_m = 5.4 \pm 0.3 \mu\text{M}$). The affinity of **5** for IS is not proof that the isomerization catalyzed by this enzyme involves the transition state **4** nor does it suggest whether such a species is cationic or anionic in nature; however, inhibition by **5** is not supportive of a mechanism which requires covalent attachment of the substrate to the enzyme or transient lactonization. That compound **5** is more effective as an inhibitor of IS than of AS, in spite of the homology of the two proteins, may reflect the specificity of AS for ammonia over water as cosubstrate and suggests that the 6-amino analogue of **5** may be a selective inhibitor of AS.

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Supplementary Material Available: Complete experimental details for the synthesis and enzymatic evaluation of **5** (8 pages). Ordering information is given on any current masthead page.

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Biosynthesis of Blastidicin S from Cytosylglucuronic Acid (CGA). Isolation of Cytosine/UDPglucuronosyltransferase and Incorporation of CGA by *Streptomyces griseochromogenes*

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Blastidicin S, **1**, an antifungal antibiotic produced by *Streptomyces griseochromogenes* used commercially for the control of *Piricularia oryzae* (rice blast), was first isolated by Takeuchi et al. in 1958.¹ Its structure and absolute stereochemistry were elucidated by chemical means²⁻⁵ and confirmed by X-ray diffraction.^{6,7} Seto et al.⁸ established that **1** is biosynthesized from cytosine, **2**, D-glucose, **3**, L- α -arginine, and L-methionine. This group has also reported the characterization of a number of structurally related metabolites from *S. griseochromogenes*.⁹⁻¹³

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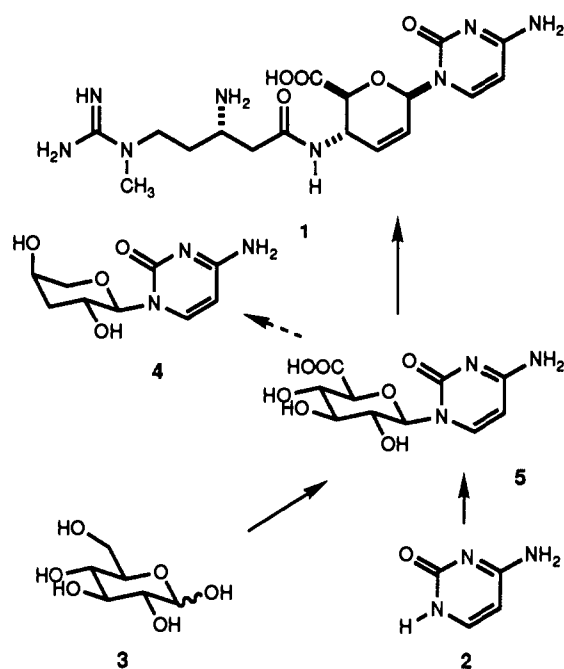
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Scheme I



We have shown that L-arginine is converted to L- β -arginine and have established the stereochemistry of the arginine-2,3-amino-mutase reaction.¹⁴ We now report the identity of the first intermediate committed to the biosynthesis of the remainder of **1** and the isolation of the responsible enzyme.

For our further studies we adopted an approach of altering the fermentation conditions to block potential biosynthetic steps and accumulate intermediates.¹⁵ When aminooxyacetic acid (a transaminase inhibitor)¹⁶ was included in a fermentation with a chemically defined medium,⁹ increases in the concentration of pentopyranine C, **4** (54 mg, 5.70-fold), and of another, unidentified metabolite ("UK", 247 mg, 82.4-fold) as well as a 33% decrease in the concentration of **1** (345 mg) were observed by HPLC (Waters Assoc. C₁₈ RadialPak column, 97% H₂O, 3% CH₃CN, 0.1% TFA, detection at 278 nm). Alternatively, inclusion of arginine hydroxamate (an inhibitor of arginine biosynthesis)¹⁷ (1.2 g/L) and cytosine (500 mg/L) decreased production of **1** 85% and increased production of **4** and "UK" 25-fold and 163-fold, respectively, while inclusion of L-ethionine (a methyltransferase inhibitor)¹⁸ (400 mg/L) and cytosine (1 g/L) only reduced **1** 10%, but increased **4** 12-fold and "UK" 443-fold (to 1.33 g/L!).

The unknown metabolite was determined to be cytosylglucuronic acid (CGA) **5** (Scheme I).¹⁹ Seto et al. had isolated **5** as a minor fermentation product and named it pentopyranic acid.¹¹ A cell-free extract of *S. griseochromogenes* was next prepared and incubated with cytosine and either UDPglucose, UDPgalactose, UDPgalacturonic acid, or UDPglucuronic acid.²⁰ In only the last incubation was substrate consumed and a product generated, which corresponded to **5** by HPLC; a preparative-scale incubation (100 mL) yielded enough **5** to obtain a ¹H NMR

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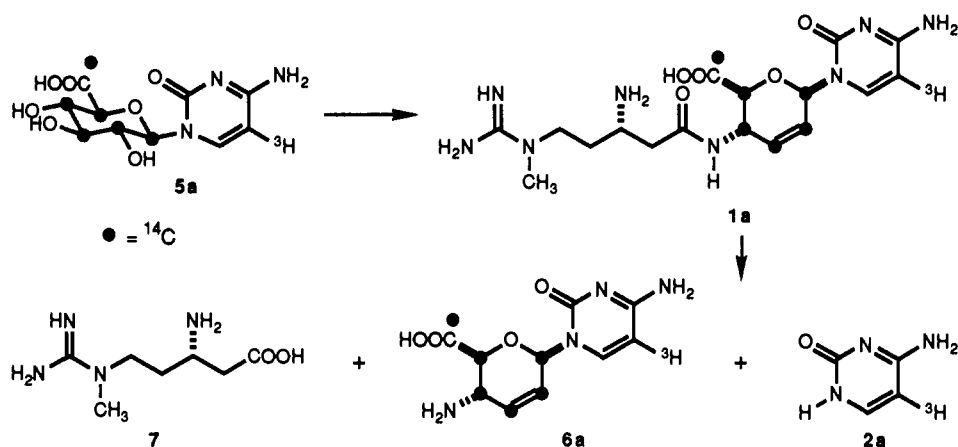
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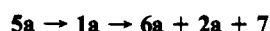
(19) The structure of **5** was established by ¹H NMR, ¹³C NMR, ¹H-¹H COSY, and ¹H-¹³C HETCOR experiments and mass spectrometric analysis.

(20) We had previously found that incorporation of [1-¹⁴C]galactose into **1** was 8.14-fold higher than incorporation of [1-¹⁴C]glucose. In addition, although XDPglucosyltransferases had been demonstrated in prokaryotes,^{21,22} UDPglucuronosyltransferase had not.²³

Scheme II



spectrum. As is common for nucleosides, **5** was not readily taken up by *S. griseochromogenes* mycelia (HPLC analysis), and efforts to prepare a cell-free extract or protoplasts that could metabolize **5** to **1** were unsuccessful. In order to test **5** as a biosynthetic intermediate, a sample, **5a** (³H/¹⁴C = 4.25, 1.29 × 10⁸ dpm/mmol ¹⁴C), was prepared enzymatically from [5-³H]cytosine (52.45 μCi, 19.9 mCi/μmol, Sigma Chemical Co.) and UDP[U-¹⁴C]-D-glucuronic acid (10.95 μCi, 250 μCi/μmol, ICN). A portion of this (10.0 mg, 4.49 × 10⁶ dpm ¹⁴C) was fed to a 200-mL fermentation in the standard complex medium¹⁴ 39 h after inoculation with a seed culture. Ninety-eight hours later workup¹⁴ yielded 455 mg of labeled blasticidin S **1a** (³H/¹⁴C = 6.75, 1.01 × 10⁴ dpm/mmol ¹⁴C). Most of **5a** was recovered from the broth unchanged (88.4%). On the basis of the unrecovered material, the ¹⁴C incorporation was 1.8%. The change in ³H/¹⁴C could be explained if only 0.14% of the **5a** fed had been hydrolyzed to [5-³H]cytosine, **2a**, since **2** had previously been incorporated almost quantitatively.⁸ To test the specificity of the incorporation, 300 mg of **1a** were hydrolyzed^{2,3} yielding labeled cytosinine **6a** (29.4 mg, ³H/¹⁴C = 6.26, 9.70 × 10³ dpm/mmol ¹⁴C, 96% retention of ¹⁴C and 89% retention of ³H from **1a**), labeled cytosine **2a** (21.4 mg, 5.84 × 10⁴ dpm/mmol ³H, 86% retention of tritium), and blasticidic acid, **7** (not isolated, but unlabeled vide supra). Some exchange of tritium had occurred during the hydrolysis. These results demonstrate unequivocally the specific intact incorporation of **5a** exclusively into the cytosinine portion of blasticidin S, **1**.



The biosyntheses of a number of nucleoside antibiotics have been studied.²⁴⁻²⁷ The biosynthesis of blasticidin S appears to be the first instance where the formation of a novel nucleoside has been demonstrated at the cell-free level to be the first committed step in the secondary pathway. In addition, although UDPglucuronosyl transferases are common in mammalian xenobiotic metabolism and at least two have been reported in fungi (also eukaryotes),^{28,29} this is the first report of such an enzyme

from a prokaryotic organism.³⁰

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(30) CGA is likely to also prove to be the first committed intermediate in the biosynthesis of arginomycin,³¹ the bagougeramines,³² gougerotin,³³ and possibly Sch 36605.³⁴

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Oligoamines as Simple and Efficient Catalysts for RNA Hydrolysis

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Recent interest has focused on the molecular design of artificial nucleases, in which catalytic residues are attached to sequence-recognizing moieties.¹ Developments have been made in oxidative fission of specific ribose residues in DNA.² However, cleavage of DNA and RNA via hydrolysis of the phosphodiester linkage has been scarcely accomplished.³⁻⁶ We report here remarkable

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