Communications to the Editor

The Oxidative Macrocyclization of Phenolic Peptides. A Biomimetic Approach to the Synthesis of the Vancomycin Family of Antibiotics

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The family of glycopeptide antibiotics, as represented by vancomycin (1), is widely used in the treatment of staphylococcal infections. These important compounds express their antibiotic activity by inhibiting bacterial cell wall biosynthesis by selectively binding to the C terminal D-Ala-D-Ala residues of peptidoglycan precursors. The specific mode of binding of these compounds is under investigation by a number of research groups; however, due to the architectural complexity of these molecules, only the syntheses of simple analogues have been accomplished to date. lef In accord with our objectives, which have been directed at the development of general approaches to the synthesis of vancomycin and other members of this family of antibiotics, we report our preliminary investigations on a biomimetic oxidative macrocyclization approach to the bicyclic C, D, E phenyl ether fragment of vancomycin.

In initial studies, we chose to construct two model peptides, one with which to assess the C-D ring macrocyclization and the other with which to assess the analogous D-E ring closure. The synthesis of the C-D ring cyclization precursor began with the *N*-Boc methyl ester 2, prepared according to previously described methodology

Scheme Ia

 $^{\it a}$ (a) TFA, thioanisole; (b) N-Boc-O-benzyl-4-hydroxyphenylglycine, diisopropylcarbodiimide, hydroxybenzotriazole (HOBt); (c) 1-[3-(dimethylamino)propyl]-3-ethylcarbodiimide hydrochloride (EDC), HOBt; (d) Bu_3SnH, Pd(II).

Scheme IIa

 a (a) NaN₃, DMSO; (b) LiOOH; (c) β -cyanoalanine (trimethylsilyl)ethyl ester, EDC, HOBt; (d) SnCl₂, MeOH; (e) N-Boc-N-methylleucine, EDC, HOBt; (f) n-Bu₄NF; (g) N-Boc-3,5-dibromo-4-hydroxyphenylglycine N-methyl amide, EDC, HOBt; (h) Bu₃SnH, Pd(II).

from this laboratory (Scheme I).³ Removal of the Boc protecting group from 2 with trifluoroacetic acid followed by coupling to the

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Scheme III

racemization-prone D-N-Boc-O-benzyl-4-hydroxyphenylglycine4 provided dipeptide 3 in 90% overall yield with less than 1% racemization as determined by HPLC analysis. After removal of the Boc protecting group of dipeptide 3 followed by base extraction, the resulting free amine was coupled to the D-N-Boc-phenylglycine 45 to yield tripeptide 5 in 81-90% yield with less than 4% racemization as determined by HPLC analysis. Treatment of tripeptide 5 with Bu₃SnH and Pd(PPh₃)₂Cl₂⁶ as catalyst removed the allyl protecting group without any concomitant hydrodehalogenation, to provide the tripeptide cyclization precursor 6 in 89-93% yield.

The tetrapeptide that was used to model the D-E ring closure was synthesized from the α -bromo carboximide 7, prepared according to methodology previously established in this laboratory (Scheme II).8 Reaction of 7 with sodium azide in dimethyl sulfoxide followed by removal of the oxazolidinone by treatment with lithium hydrogen peroxide⁹ provided the α -azido acid 8

(4) Prepared as described by Kamiya et al.: Kamiya, T.; Hashimoto, M.;

(86%), which was then coupled to the (trimethylsilyl)ethyl (TMSE) ester of L-β-cyanoalanine¹⁰ to give the dipeptide 9 in 84% Reduction of the azide with stannous chloride in methanol¹³ followed by coupling to D-N-Boc-N-methylleucine then provided tripeptide 10 in 92% yield. The removal of the (trimethylsilyl)ethyl protecting group was accomplished in 99% yield by treatment with tetrabutylammonium fluoride in DMF.14 Coupling the resulting acid 11 with the N-methyl amide allyl ether of the D-phenylglycine 4¹⁵ followed by removal of the allyl protecting groups with Bu₃SnH and Pd(PPh₃)₂Cl₂ afforded the D-E cyclization precursor 12 in 81% overall yield. 16

Nakaguchi, O.; Oku, T. Tetrahedron 1979, 35, 323-328. (5) Prepared in 87% overall yield by bromination of D-hydroxyphenyl-glycine with bromine in acetic acid, followed by protection with tert-butyl pyrocarbonate in dioxane/water at pH 11. HPLC analysis (Pirkle column) of the dimethylated derivative (prepared by treatment with diazomethane) revealed that less than 1% racemization had occurred during the above re-

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⁽⁷⁾ Established methods for the removal of the allyl group, such as employing RhPPh3Cl to isomerize the allyl ether to the enol ether, proved unsuccessful due to the unreactivity of the resultant enol ether toward acidic hydrolysis or mercuric acetate oxidation.

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⁽¹⁰⁾ Prepared in 96% yield by treatment of Cbz-β-cyanoalanine with dicyclohexylcarbodiimide and (dimethylamino)pyridine in the presence of (trimethylsilyl)ethanol, followed by hydrogenolysis of the Cbz protecting group with Pd/C as catalyst. The enantiomeric purity of the resulting amino ester was determined to be 96% by GC analysis of the α -methoxy- α -(trifluoromethyl)phenylacetamide derivative.

⁽¹¹⁾ We chose to employ β -cyanoalanine as a protected derivative of asparagine in order to avoid the potential side reactions during activation and peptide coupling that are inherent to asparagine-containing peptides. nonpolar nature of β -cyanoalanine also greatly facilitated the manipulation and purification of the peptides that contained this amino acid.

⁽¹²⁾ Previous studies have shown that the α -azido acids do not racemize

under standard peptide coupling conditions (ref 2b).
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⁽¹⁴⁾ Carpino, L. A.; Sali, A. C. J. Chem. Soc., Chem. Commun. 1979,

⁽¹⁵⁾ Prepared by coupling the pentafluorophenol ester of 4 (DICI, pentafluorophenol) with N-methylamine. The enantiomeric purity was determined to be 93% by HPLC analysis of the α -methoxy- α -(trifluoromethyl)phenylacetamide methyl phenyl ether derivative.

Our optimized conditions for promoting the oxidative cyclization of substrates 6 and 12 (Scheme III) are derived from a related transformation initially reported by Yamamura¹⁷ and later modified by Inoue. ^{18,19} We have found that the optimal protocol for the cyclization of 6 is oxidation with 10 equiv of thallium(III) nitrate trihydrate (TTN) (excess TTN is necessary to ensure complete reaction) in 5:1 THF/methanol at 1 mM concentration with 3 equiv of pyridine/equiv of TTN to serve as an acid scavenger. Increasing the ratio of THF to methanol results in incomplete reaction, while increasing the ratio of methanol to THF results in a lower yield. The reduction of the resulting para-quinol 13 is accomplished in situ by the addition of excess CrCl₂.²⁰ We have found these conditions to be superior to the zinc/acetic acid reduction described by Yamamura¹⁹ as they avoid the isolation of the unstable intermediate para-quinol methyl ether. Under the conditions described above, the cyclic product 14 is isolated in 42% overall yield from the cyclization precursor 6.

Model peptide 12 was cyclized and subsequently reduced under analogous conditions except that 1:1 CH2Cl2/methanol was employed as the solvent. When these conditions were employed, the macrocyclic diphenyl ether 16 was obtained in 48% overall yield. One notable difference between the two macrocyclizations is the displacement of bromine by methoxide in the formation of para-quinol 15.21 Presumably, this substitution occurs in the cyclization of 12 and not in the cyclization of 6 due to a more sterically crowded environment at the para position of the intermediate leading to 15. These observations thus dictate the order of assemblage of the macrobicyclic diether 19.

In order to evaluate the oxidative coupling strategy to provide the C, D, E bicyclic phenyl ether vancomycin synthon, the monocyclic diphenyl ether 14 was treated with trifluoroacetic acid to remove the Boc protecting group, and the resulting amine was coupled to tripeptide 11 with disopropylcarbodiimide and hydroxybenzotriazole to provide the hexapeptide 17 in 72-78% overall yield (Scheme III).16 The allyl group was then removed as described previously in 92-93% yield to provide hexapeptide 18. The optimal conditions for the cyclization of 18 were found to be 5 equiv of TTN in 30:1 methylene chloride/methanol at 1 mM concentration (4 h, -23 °C). After in situ reduction of the resulting para-quinol ether with excess CrCl₂, the dicyclic compound 19 was obtained in 40% overall yield.

These studies clearly demonstrate the feasibility of pursuing a total synthesis of vancomycin and related antibiotics via biomimetic oxidative phenolic coupling.

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Supplementary Material Available: Spectral data for all compounds and detailed experimental procedures for the oxidative macrocyclizations as well as for the syntheses of 2-6, 17, and 18 (15 pages). Ordering information is given on any current masthead page.

Aristolochene Biosynthesis and Enzymatic Cyclization of Farnesyl Pyrophosphate

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Aristolochene (1) is a bicyclic sesquiterpene belonging to the eremophilane group of hydrocarbons. The (-) enantiomer of 1 was first isolated in 1970 by Govindachari et al. from the plant Aristolochia indica. It is also reported to occur in Bixa orellana leaf oil and in the defensive secretions of Syntermes soldier termites.^{2,3} The (+) enantiomer 1 was recently isolated in our laboratory from the mycelial extracts of the fungus Aspergillus terreus.⁴ The (+) enantiomer is also the probable biosynthetic precursor of PR toxin produced by Penicillium roquefortii.5 Recently, Hohn and co-workers have isolated aristolochene synthase from P. roquefortii⁶ and purified the enzyme to homogeneity.7

The proposed mechanism for the formation of aristolochene from farnesyl pyrophosphate (FPP) (2), the universal precursor of sesquiterpenes, 10 is shown in Scheme I. Cell-free extracts of A. terreus prepared from mycelia harvested between 45 and 60 h after inoculation showed terpenoid cyclase activity.¹¹ Preparative incubation of [1-3H]FPP (2a)12 with crude cell-free extracts produced radioactive hydrocarbon 1a,13 which was found to cochromatograph with synthetic (±)-aristolochene¹⁴ on TLC (SiO₂, AgNO₃-SiO₂) as well as by radio-GC analysis (FFAP). Dilution with carrier (±)-aristolochene and oxidation with MCPBA followed by hydrolysis with HClO₄ gave the corresponding diol 3a, which was recrystallized to constant activity, thereby confirming

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