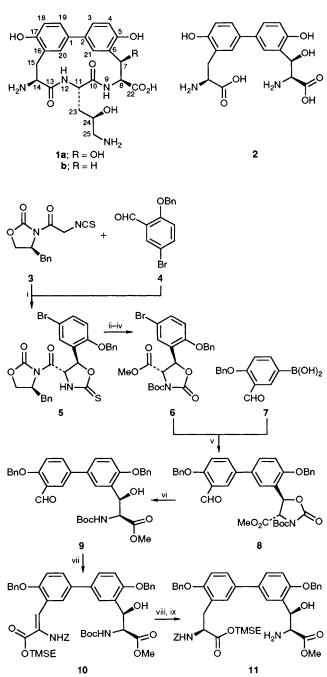
The Synthesis of Biphenomycin A¹

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Biphenomycin A, a highly potent antibiotic against Gram-positive, β -lactam-resistant bacteria, which was previously isolated from culture filtrates of *Streptomyces griseorubiginosus* No. 43708, has now been synthesized.

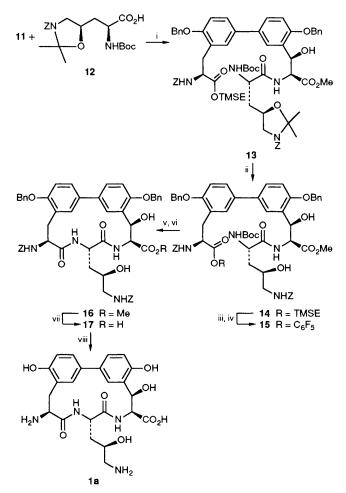
Biphenomycins A **1a** and B **1b** are cyclic tripeptides exhibiting high antibiotic activities against Gram-positive, β -lactamresistant bacteria and have previously been isolated from culture filtrates of *Streptomyces griseorubiginosus* No. 43708 in the ratio of 10:1 (**1a**: **1b**).^{2–4} These cyclopeptides, together with those of the vancomycin group, are unique in that they contain a biphenyl structural moiety. We have reported on the total synthesis of biphenomycin B^5 which contains the two non-proteinogenic amino acids (2S,4R)-hydroxyornithine and (S,S)-diisotyrosine. In place of the latter, the main metabolite biphenomycin A contains the corresponding hydroxy derivative **2** with an additional stereogenic centre *i.e.* the biphenyl



Abbreviations: Bn = benzyl; Boc = *tert*-butoxycarbonyl; TMSE = trimethylsilylethyl; Z = benzyloxycarbonyl

Scheme 1 Reagents and conditions: i, 3, Sn(OTf)₂, N-ethylpiperidine, tetrahydrofuran (THF), -78 °C, 1.5 h, 4, 2 h, -78 °C, 75%; ii, MeOMgBr, MeOH, 0 °C, 3 min, 90%; iii, (Boc)₂O, dimethylaminopyridine (DMAP), CH₂Cl₂, room temp., 20 min, 90%; iv, H₂O₂, formic acid, CH₂Cl₂, 0 °C, 30 min, 90%; v, THF, (Ph₃P)₄Pd, reflux, 1 h, 12 mol dm⁻³ Na₂CO₃, reflux 24 h, 50%; vi, CsCO₃, MeOH, room temp., 2 h, 80%; vii, N-benzyloxycarbonyl(dimethoxyphosphoryl) glycine trimethylsilylethyl ester, 1,8-diazabicyclo-[5.4.0]undec-7-ene (DBU), CH₂Cl₂, -78 °C, 15 h, 80%; viii, [Rh(1.5-cod) (dipamp)]⁺ BF₄⁻, H₂, MeOH, room temp., 72 h, quant.; ix, 6 mol dm⁻³ HCl, dioxane, room temp., 30 min, quant.

unit formally represents the dimer from (S)-o-hydroxyphenylalanine and (2S,3R)-o-hydroxyphenylserine. In this communication we describe the total synthesis of biphenomycin A. The key step for the construction of the biphenyl unit is the Pd⁰catalysed coupling⁶ of an arylboronic acid containing an unprotected aldehyde function with a protected 2-hydroxy-5-



Scheme 2 Reagents and conditions: i, N-(3-dimethylaminopropyl)-N'ethylcarbodiimide hydrochloride (EDC), hydroxybenzotriazole, CH₂Cl₂, -10 to 20 °C, 14 h, 80%; ii, AcOH-H₂O (9:1), 50 °C, 2 h, 70%; iii, Bu₄NF, dimethylformamide, room temp., 30 min; iv, C₆F₅OH, EDC, CH₂Cl₂, -15 to 20 °C, 14 h, 95%; v, 6 mol dm⁻³ HCl, dioxane, room temp., 30 min, evaporation, quant.; vi, CHCl₃– NaHCO₃ (aq.), 20 °C, 5 min, 70%; vii, 1 mol dm⁻³ LiOH, dioxane, room temp., 30 min, evaporation, quant.; viii, trimethylsilyl bromidethioanisole (1:1), CF₃CO₂H, 2 h, room temp., 60%

bromophenylserine. In the course of the synthesis it was necessary to mask two phenolic hydroxy groups, two amino groups, two hydroxy groups and one carboxy group with functions that are compatible with the protecting groups of the α -amino group of hydroxyornithine and the one carboxy group of the hydroxy diisotyrosine which needs to be cleaved during the peptide construction and ring closure steps.

The synthesis of a protected (R)-hydroxy-(S,S)-diisotyrosine 11 is illustrated in Scheme 1. The aryl bromide 6 represents a completely protected 2-hydroxy-5-bromophenylserine and can be constructed via 5 in excellent enantioselectivity and diastereoselectivity by the Evans method.7 The Pd⁰-catalysed coupling of 6, with the boronic acid 7 containing an unprotected aldehyde function gives rise to the biphenyl derivative 8. After opening of the oxazolidinone, the highly diastereoselective8 construction of the Z-dehydroamino acid derivative 10 is achieved by condensation of 9 with an appropriate phosphorylglycine9 and subsequent completely diastereoselective hydrogenation using the [Rh(1,5-cod)-(dipamp)]+ BF₄- hydrogenation catalyst {cod = cycloocta-1,5-diene; dipamp = (R,R)-1,2-bis[2-(2-methoxyphenyl)phenylphosphino]ethane}¹⁰ gives rise to the biphenyl unit 11. The further steps are illustrated in Scheme 2. The construction of the linear substrate 13 for the ring closure reaction is achieved starting from the hydroxyornithine derivative 125,11 in analogy to the synthesis of biphenomycin B. The only exception is that the free carboxy group is now protected as a methyl ester (in place of a benzyl ester in the biphenomycin B synthesis). The ring closure reaction of the ω -aminopentafluorophenyl ester in the two phase system CHCl₃-NaHCO₃ (aq.) occurs within 5 min to give a 70% yield without dilution.¹² After saponification of the methyl ester 16, the four benzylic protecting functions are cleaved by the method of Yajima¹³ using trimethylsilyl bromide-thioanisole in trifluoroacetic acid. The synthesized product is separated by reversed phase chromatography and lyophilized. It is identical in every respect with the natural compound.

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References

- 1 For Part 82 of the series Amino Acids and Peptides see U. Schmidt and J. Schmidt, J. Chem. Soc., Chem. Commun., 1992, 529.
- 2 Isolation and structure: M. Ezaki, M. Iwami, M. Yamashita, S. Hashimoto, T. Komori, K. Umehara, Y. Mine, M. Kohsaka, H. Aoki and H. Imanaka, J. Antibiot., 1985, 38, 1453; I. Uchida, N.

Shigematsu, M. Ezaki, M. Hashimoto, H. Aoki and H. Imanaka, J. Antibiot., 1985, 38, 1462; I. Uchida, M. Ezaki, N. Shigematsu and M. Hashimoto, J. Org. Chem., 1985, 50, 1342.

- Configuration: R. Kannan and D. H. Williams, J. Org. Chem., 1987, **52**, 5435.
- 4 Configuration and conformation: J. C. Hempel and F. K. Brown, J. Am. Chem. Soc., 1989, 111, 7323; F. K. Brown, J. C. Hempel, J. S. Dixon, S. Amato, L. Mueller and P. W. Jeffs, J. Am. Chem. Soc., 1989, 111, 7328.
- 5 U. Schmidt, R. Meyer, V. Leitenberger, A. Lieberknecht and H. Griesser, J. Chem. Soc., Chem. Commun., 1991, 275.
- 6 N. Miyaura, T. Yanagi and A. Suzuki, Synth. Commun., 1981, 11, 513.
- 7 D. A. Evans and A. E. Weber, J. Am. Chem. Soc., 1986, 108, 6757.
- 8 U. Schmidt, H. Griesser, V. Leitenberger, R. Mangold, R. Meyer and B. Riedl, Synthesis, 1992, 487
- 9 U. Schmidt, A. Lieberknecht and J. Wild, Angew. Chem., 1982, 94, 797; Angew. Chem., Int. Ed. Engl., 1982, 21, 770; Angew. Chem. Suppl., 1982, 1682; U. Schmidt, A. Lieberknecht and J. Wild, Synthesis, 1984, 53.
- 10 W. S. Knowles, M. J. Sabacky, B. D. Vineyard and D. J. Weinkauff, J. Am. Chem. Soc., 1975, 97, 2567; B. D. Vineyard, W. S. Knowles, M. J. Sabacky, G. L. Bachman and D. J. Weinkauff, J. Am. Chem. Soc., 1977, 99, 5946.
- 11 U. Schmidt, R. Meyer, V. Leitenberger, F. Stäbler and A. Lieberknecht, Synthesis, 1991, 409.
- 12 U. Schmidt, A. Lieberknecht, H. Griesser and J. Talbiersky, J. Org. Chem., 1982, 47, 3261; in a two-phase system: U. Schmidt, R. Utz, A. Lieberknecht, H. Griesser, B. Potzolli, J. Bahr, K. Wagner and P. Fischer, Synthesis, 1987, 236; U. Schmidt, M. Kroner and H. Griesser, *Tetrahedron Lett.*, 1988, **29**, 4407. 13 H. Yajima, N. Fujii, S. Funakashi, T. Watanabe, E. Murayama
- and A. Otaka, Tetrahedron, 1988, 44, 805.