

Synthesis of the Antibiotically Active Part of Agrocin 84

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Abstract. Phosphitylation of bis-O-silylated *threo*-2,3-dihydroxy-4-methylpentanamide and condensation of the resulting N-acylphosphorodiamidite with 2'-O-acetyl-3'-deoxyarabinoadenosine led, after oxidation and deprotection, to the isolation of the title compound. © 1998 Elsevier Science Ltd. All rights reserved.

Nearly two decades ago, Kerr et al.¹ reported that the biological control of crown gall disease can be achieved by inoculation of susceptible plants with the non-pathogenic strain 84 of Agrobacterium radiobacter. Structural elucidation revealed that the controlling agent, so-called agrocin 84 (1), consists of the 3'-deoxy-D-arabinoadenosine core unit in which the N^6 -exocyclic amino function of the adeninyl and HO-5' of arabinosyl units are connected via phosphoramidate bonds to the anomeric centre of D-glucofuranose and D-threo-2,3-dihydroxy-4-methylpentanamide, respectively. Structure-activity studies showed that fragment 2, formed by brief heating (see Scheme 1) of agrocin 84 (1), exhibits a non-specific antibiotic activity for pathogenic strains. Moreover, subjection of agrocin 84 (1) to mild basic conditions gives fragment 3 displaying no antibiotic activity. However, the N^6 -D-glucofuranosylphosphoramidate function in 3 proved to be essential for strain specificity of the antibiotic. Thus far, synthetic efforts in preparing agrocin 84 (1) have been limited to analogues of fragment 3 in which adenosine, instead of 3-deoxyarabinoadenosine, is either linked via a P-N(6)-phosphoramidate bond to HO-6 of methyl- β -D-glucopyranoside² or, as in agrocin 84 (1), to HO-1 (β) of D-glucofuranose.³

Reagents and conditions. i. 110 °C, pH 7.5, 10 min; ii. 25 °C, 1M NH₄OH, 3 h.

We here report for the first time a convenient route to the synthesis of the biologically active fragment 2. A straightforward preparation of target compound 2 is hampered by two main factors. First of all, the P-O(5') bond is readily cleaved (cf. conversion of 1 into 3) under mild basic conditions. Secondly, N-O phosphoryl

Reagents and conditions: *i* DCC, DMAP, BnOH, 89 %; ii (a) LDA, PhSeBr (b) H $_2$ O $_2$, 74%; iii AD-mix-β, 90%; iv TBS-Tf, 2,6-lutidine, CH $_2$ Cl $_2$, 0 °C, 6 h, 85%; v. (a) H $_2$, Pd/C, t-BuOH/THF, 10 min; (b) BOP,DIPEA, NH $_3$, t-BuOH/THF, 20 min, 78 % (two steps);

migration of the N-acylphosphoramidate bond in 2 readily occurs¹ under mild acidic conditions. It was anticipated that protection of the diol function in the 2(S), 3(R)-dihydroxy-4-methylpentanamide unit with tert-butyldimethylsilyl (TBS) groups, as in compound 9 in Scheme 2, would not only be compatible with 2, but also with the removal of the base labile groups required for transient protection of HO-2' in the 3'-deoxyarabinoadenosine as well as the N-acylphosphoramidate function. The preparation of key intermediate 9 is depicted in Scheme 2, and commences with the esterification of commercially available 4-methylpentanoic acid (4). Thus, condensation of 4 with benzyl alcohol under the influence of DCC/DMAP gave the benzyl ester 5, which was converted into the corresponding E-alkene derivative 6 involving phenyl selenation followed by

Scheme 3

Reagents and conditions: *i.* DIPEA, CH $_2$ Cl $_2$, 100%; *ii.* (a) TMS-Cl, Pyridine, 20 min; (b) BzCl, Pyridine, 2 h; (c) NH $_3$ •H $_2$ O, Pyridine 0 °C 30 min; *iii.* DMT-Cl, Pyridine, 4h; *iv.* NH $_3$ /MeOH, 60 h; *v.* Ac $_2$ O, Pyridine, 4 °C, 32 h; *vi.* DCA/CH $_2$ Cl $_2$, HSnBu $_3$, 0 °C, 30 min; *vii.* (a) 1-MTT (6 eq.), CH $_3$ CN, 1 h; (b) I $_2$ (0.2 M in pyridine)/THF/H $_2$ O, 0 °C; *viii.* NH $_3$ •H $_2$ O/MeOH (2/1), 15 h; *ix.* 1M TBAF in DMF, 6 min.

oxidation⁴. Sharpless asymmetric dihydroxylation⁵ of 6 afforded 7 (95% ee), the *threo* configuration of which was firmly established by its conversion into known *threo*-2,3-dihydroxy 4-methylpentanoic acid $\{[\alpha]_D +12.8 \text{ (c } 0.36, \text{EtOH)}; [\alpha]_D \text{ lit.}^1 +13.6\}$. Silylation of 7 with *tert*-butyldimethylsilyl triflate (TBS-Tf) in the presence

of 2,6-lutidine proceeded smoothly⁶ to yield the fully protected derivative **8**. Transformation of **8** into the required building unit **9** could be realized most effectively by executing the following three-step one-pot procedure. Thus, reductive debenzylation of **8**, and subsequent *in situ* activation of the free carboxylic acid group with BOP⁷ in the presence of diisopropylethylamine (DIPEA), followed by bubbling dry gaseous ammonia through the reaction mixture gave **9** in a yield of 39% over the seven steps.

The 2'-O-acetyl-3'-deoxyarabinoadenosine building block 15 was readily accessible (Scheme 3) from known⁸ 3'-deoxyarabinoadenosine (10) by the following well established protecting group manipulations. Tritylation of the N^6 -benzoyl derivative 11, prepared from 10 by the Jones N-acylation procedure⁹, with 4,4'-dimethoxytrityl chloride (DMT-Cl) followed by ammonolysis of the N^6 -benzoyl group of 12 led to the partially protected derivative 13. Selective O-acylation¹⁰ of 13 with acetic anhydride at 4 °C gave, after detritylation of 14 with dichloroacetic acid (DCA) in the presence of the effective cation scavenger tributyltin hydride, the 2'-O-acetyl-3'-deoxyarabinoadenosine 15 in an overall yield of 33% (based on 10).

In line with our previous experiences¹¹, introduction of the *N*-acylphosphoramidate bond between the two building units **9** and **15** could be readily accomplished *via* the two-step phosphitylation protocol depicted in Scheme 3. Thus, phosphitylation of **9** with the bifunctional reagent **16** in the presence of DIPEA gave, after purification by silica gel chromatography, the monofunctional phosphorodiamidite **17** in a near quantitative yield. Selective *O*-phosphitylation of **15** with **17** using an excess of the rather acidic activating reagent 1-methyltetrazoline-5-thione¹² (1-MTT) gave, after *in situ* oxidation of the transiently formed *N*-acylphosphoramidite intermediate with iodine in aqueous pyridine, the fully protected compound **18**¹³ in a near quantitative yield. Removal of the two base-labile protecting groups (*i.e.* acetyl and cyanoethyl) with aqueous ammonia for 15 h at 20 °C yielded, as gauged by ³¹P NMR spectroscopy, the partially deprotected product **19**. Subsequent desilylation of **19** with tetrabutylammonium fluoride (TBAF) in dry DMF for 6 min, followed by

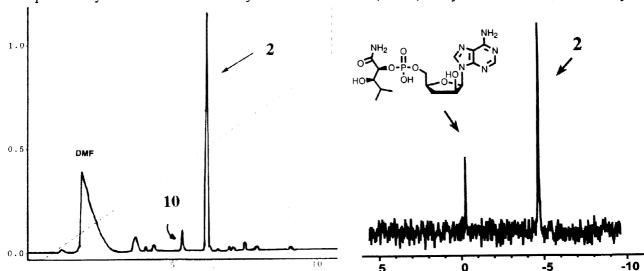


Figure 1. HPLC trace of crude 2.

Figure 2. 31P NMR spectrum of 2 after mild acid treatment

quenching with triethylammonium acetate (TEAA) buffer (pH 7, 1 M) afforded crude 2 which was *inter alia* contaminated (see Fig. 1) with a small amount (4.6%) of 3'-deoxyarabinoadenosine (10). Purification of the crude product by RP HPLC [(MeCN/TEAA (20 mM, pH 7.5)] gave homogeneous 2 in 35% yield. The homogeneity and structure of 2, fully ascertained by analytical as well as spectroscopic data, was also independently confirmed by the following chemical evidence. Subjection of 2 to H₂O-HOAc (9/1, v/v) for 2.5 h at ambient temperature led to a product having the same retention time (HPLC) and ESI-MS spectrum as the starting product. However, the ³¹P NMR spectrum showed two distinct resonances (see Fig. 2). The major

resonance at δ_P -4.8 ppm coincided with the signal observed for **2** and the minor resonance at δ_P -0.2 ppm may be attributed to the earlier by Kerr *et al.*¹ proposed N \rightarrow O phosphoryl migration product of **2**.

The results presented in this paper clearly show that the *N*-phosphorodiamidite derivative 17 is an effective building unit in the synthesis of the biologically active part of agrocin 84 (1). In addition, the partially protected compound 18 promises to be of great value in the future preparation of agrocin 84 (1) and analogs thereof.

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- 13. Compound **18** (2:1 mixture of diastereoisomers at phosphorus): ³¹P (80.7 MHz, CDCl₃) -1.57, -1.63. ¹H NMR ¹H-resonances marked with asterisk showed apparent splitting due to presence of the minor P-diastereoisomer (200 MHz, CDCl₃) δ 8.33 (s 1H H-8), 8.13* (s 1H H-2), 7.92* (br. d 1H NH J_{PH} 13), 6.45 (d 1H H-1' J_{1'2'} 5.2); 5.72 (br. s 2H NH₂); 5.53 (m 1H H-2'); 4.44 (m 5H H-4', H-5', CH₂CH₂O); 4.12* (d 1H H-2" J 3.6); 3.56 (dd 1H H-3" J₃₂ 3.3, J₃₄ 6.5); 2.63 (m 3H H-3'a, CH₂CH₂O); 2.34 (m 1H H-3b); 1.84* (s 3H CH₃ Ac); 1.64 (br. m 1H H-4"); 0.98, 0.95, 0.93, 0.91 (m 24H H-5", CH₃C TBS); 0.12, 0.05, 0.04, -0.01 (s 12H CH₃Si). ¹³C NMR (75.5 MHz, CDCl₃) δ 173.7 (CONH), 169.1 (C=O, Ac); 155.5 (C-6); 152.7 (C-2); 149.4 (C-4); 139.7 (C-8); 118.9 (C-5); 116.3 (CN); 84.3, 83.9 (C-1'); 78.6 (C-3"); 75.5, 75.4 (C-4', C-2"); 71.7, 71.6 (C-2'); 68.8, 68.4 (CH₂, CNE); 62.3, 62.1, 62.0 (C-5'); 32.3, 32.0 (C-3'); 30.7 (C-4"); 25.7, 25.5, 20.3, 19.8 (CH₃, tBu); 20.3 (CH₃, Ac); 19.8 (C-5") 19.6, 19.4 (CH₂CN); 18.9 (C-5a"); 17.9 (C_q, tBu); -5.3, -5.0, -4.5 (CH₃, MeSi).

 Compound 2: ³¹P (243 MHz, D₂O) -4.7. ¹H NMR (600 MHz, D₂O, NH₄*- form) δ 8.53 (1H s H-8), 8.13 (1H s H-2), 6.26 (1H d H-1' J_{1'2'} 5.2); 4.76 (m 1H H-2'); 4.41 (m 1H H-4'); 4.17 (m 2H H-2", H-5'); 4.09 (m 1H H-5'a); 3.44 (dd 1H H-3" J₁ 1.3, J₂ 9); 2.46 (ddd 1H H-3"a J₁ 6.7, J₂ 6.7, J_{gem} -13.4); 2.09 (ddd 1H H-3"b J₁ 8.5, J₂ 8.5, J_{gem} -13.3); 1.77 (m 1H H-4"); 0.92 (d 3H H-5" J₅₄ 6.7); 0.83 (s 3H H-5"a J₅₄ 6.9). ¹³C NMR (151 MHz, D₂O, NH₄*- form) δ 178.6 (C-1"); 157.8 (C-6); 151.2 (C-2); 149.3 (C-4); 142.5 (C-8); 119.2 (C-5); 86.0 (C-1'); 78.4 (C-3"); 77.8 (C-4', ³J_{CP} 8.6); 73.0 (C-2", ³J_{CP} 6.6); 71.5 (C-2'); 66.8 (C-5', ²J_{CP} 4.3); 32.8 (C-3'); 30.6 (C-4"); 19.2, 18.9 (C-5", C-5a"). ESI-MS (m/z) 459.0 (M-H)⁻.