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Preparation of a protected phosphoramidon precursor via an *H*-Phosphonate coupling strategy

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Abstract—The preparation of a phosphoramidon precursor is described using a phosphorus(III) coupling protocol. © 2006 Elsevier Ltd. All rights reserved.

Phosphoramidon (1) was isolated from Streptomyces tanashiensis in 1972 by a Japanese group at Keio University. It consists of a leucyl-tryptophan dipeptide coupled to α-L-rhamnose via a phosphoramidate linkage (Fig. 1). Phosphoramidon possesses an array of biological activity, including inhibition of the bacterial zinc protease thermolysin,² the mammalian plasma membrane zinc peptidase,³ and a metalloprotease endothelin converting enzyme. 4 Its ability to inhibit endothelin converting enzyme has rendered it a valuable tool in the study of hypertension,⁵ stroke,⁶ and diseases of the kidney.^{7,8} Phosphoramidon has been particularly scrutinized as a potent and specific inhibitor of a zinc metalloproteinase identified as an endothelin converting endopeptidase. As a result, it has been a model for the development of numerous novel inhibitors developed over the last decade. Phosphoramidon has also been used in the study of the upstream inhibition of endothelin conversion.

To support studies of this type, phosphoramidon is sold commercially. In practice however, supplies of phosphoramidon are subject to both high cost (ca. \$45/mg), and inconsistent availability. This report details our effort to produce gram quantities of phosphoramidon, which led ultimately to an improvement of the convergent coupling of the monosaccharide, phosphate ester, and dipeptide fragments of the natural product.

Our initial synthetic route to phosphoramidon followed the first report of its synthesis by De Nanteuil and coworkers in 1995 (Fig. 2),¹¹ in which three fragments served as key intermediates: triacylated L-rhamnose (2), phenyl dichlorophosphate (3), and the ethyl ester of leucyl-tryptophan (4). In their report, the three components were linked in a one-pot, double condensation reaction to afford the protected intermediate (11), which was saponified to afford 1 in 7.5% yield over two steps.

Toward that end, commercially available L-rhamnose 5 was peracylated under Steglich conditions to afford 6 in 99% yield (Scheme 1). ¹² Chemoselective deacylation of 6 at the anomeric position gave α -anomer 2 as a white solid in 77% yield. ¹³ The carbobenzyloxy protected

Figure 1. Phosphoramidon.

$$\mathbf{1} \Longrightarrow_{\mathsf{AcO}} \overset{\mathsf{OH}}{\underset{\mathsf{AcO}}{\bigvee}} \overset{\mathsf{OH}}{\underset{\mathsf{OAc}}{\bigvee}} \overset{\mathsf{OH}}{\underset{\mathsf{OPh}}{\bigvee}} \overset{\mathsf{Bu}}{\underset{\mathsf{DPh}}{\bigvee}} \overset{\mathsf{H}}{\underset{\mathsf{N}}{\bigvee}} \overset{\mathsf{CO}_2\mathsf{Et}}{\underset{\mathsf{N}}{\bigvee}}$$

Figure 2. Key fragments for the synthesis of phosphoramidon (De Nanteuil).

Keywords: Phosphoramidon; H-Phosphonate; Synthesis.

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Scheme 1. Unsuccessful coupling based on literature protocol.

dipeptide **10** was prepared in 98% yield via union of commercially available *N*-carbobenzyloxy-L-leucine (7) with the ethyl ester of L-tryptophan (9) (prepared in 89% yield by esterification of L-tryptophan **8** in refluxing ethanol with two equivalents of DL-camphorsulfonic acid) using dicyclohexylcarbodiimide (DCC) and 1-hydroxybenzotriazole (HOBT). ¹⁴ Palladium catalyzed hydrogenolysis of the Cbz group of **10** gave dipeptide **4** in 99% yield.

With ample quantites of 2 and 4 in hand, the literature coupling conditions involving phenyl dichlorophosphate 3 failed to provide any product 11. As the result of several attempts to effect the coupling, we successfully isolated glycosyl chloride 12 (anomeric proton, δ 5.87 (d, J = 2 Hz)). Furthermore, we noted the propensity for dipeptide 4 to transform slowly to cyclic dipeptide 13, even upon standing in purified form at low temperature. Through various modifications to the coupling protocol, we surmised that formation of the glycosyl chloride was both evidence that phosphorylation of the monosaccharide was occurring (leading to 12), as well as an indication that chloride displacement of the phosphate is competitive with dipeptide coupling. Therefore, we turned to a effective phosphorus lynchpin more for couplings.

To circumvent the failings of the doubly activated phosphorus reagent, the coupling route was redesigned to a step-wise approach. We hypothesized that acceleration of the dipeptide-glycosyl phosphate coupling might be achieved through the use of more

electrophilic phosphorus(III) H-phosphonates, for which several bench stable options can be found in the literature. 15 Phosphitylation of the anomeric alcohol of 2 with 14¹⁶ was followed by a hydrolysis protocol to afford H-phosphonate 15 in quantitative yield (Scheme 2).¹⁷ This step required chromatographic separation over silica gel to remove the salicylic acid. The subsequent coupling involves four distinct steps: (1) conversion of the tetracoordinate P(III) H-phosphonate to the tri-coordinate P(III) bis(silyl)phosphite, (2) oxidation of phosphorus with molecular iodine, (3) formation of the presumed phosphoryl pyridinium ion and (4) trapping with dipeptide 4 to afford 16. We found that this sequence furnished the desired aminophosphonate in 44% yield after silica gel chromatography. The anomeric center was assigned as α by comparison of ${}^3J_{C(1)H-P}$ (7.5 Hz) to known α linked phosphoramidates of rhamnose. The ^{31}P NMR (carbon and proton decoupled) showed a singlet at 3.83 ppm. As a practical note, the isolable compounds contain achiral phosphorus, thereby simplifying the spectral data. A total of 18.8 g of 16 was produced by this method, and the final saponification to phosphoramidon has been reported by De Nanteuil.¹¹

In summary, a two step coupling protocol utilizing salicylate phosphorus(III) reagent 14 allowed the efficient, multigram scale synthesis of phosphoramidon precursor 16. This synthesis should result in a more consistent and inexpensive supply of this highly employed, pharmacologically important agent as a tool in chemical biology.

Scheme 2. Phosphorus(III) coupling protocol.

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Supplementary data

Supplementary data associated with this article can be found, in the online version, at doi:10.1016/j.bmcl.2006.08.024.

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