(±)-7-DEAZAARISTEROMYCIN LACKING THE HYDROXYMETHYL SUBSTITUENT

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Abstract: As part of a plan to uncover new nucleoside based inhibitors of S-adenosyl-L-homocysteine (AdoHcy) hydrolase that are incapable of undergoing competing nucleotide formation, a derivative of 7-deazaaristeromycin (carbocyclic tubercidin) (5) has been prepared that lacks the C-5' hydroxymethyl side-chain. This compound did not exhibit appreciable cytotoxicity or activity against a number of DNA and RNA viruses and was not inhibitory towards AdoHcy hydrolase.

A common feature of many agents currently under consideration as anti-human cytomegalovirus agents is their ability to inhibit the virally induced DNA polymerase¹ after conversion to their 5'- or "5'-like" monophosphate and, in turn, triphosphate derivatives. A prominent example that follows this sequence of events with a clinically useful anti-HCMV outcome is 9-[(1,3-dihydroxy-2-propoxy)methyl]guanine (DHPG or ganciclovir).²⁻⁵ Unfortunately, undesirable side-effects⁶ are associated with DHPG that are the likely consequence of the phosphorylation sequence. A potentially more favorable approach to designing other nucleoside derived anti-HCMV agents would be to study compounds less likely to undergo the initial phosphorylation. Among a number of possibilities in this direction, we considered shortening⁷ or eliminating⁸ the C-5' hydroxymethyl moiety of aristeromycin (1). This led to C-5'-nor aristeromycin (2) as a potent, non-toxic anti-HCMV agent that was also effective against a number of other viruses.⁷ These properties for 2 were attributed⁷ to inhibition of S-adenosyl-L-homocysteine (AdoHcy) hydrolase and revealed a new enzymic target for anti-HCMV agent design.

As part of our effort to develop antiviral agents based on 7-deaza carbocyclic nucleosides, we turned our attention to similar modifications of carbocyclic 7-deazaadenosine (3, 7-deazaaristeromycin), which has AdoHcy

hydrolase inhibitory properties ¹⁰ but shows cytotoxicity that, by analogy to tubercidin, is likely the result of phosphorylation at its C-5' center. Following the same side-chain alteration rationale described above that led to 2, compounds 4 and 5 arose as desirable targets. Compound 4 has been reported by us elsewhere. ¹¹ Compound 5 is reported here.

Thus, $(1\alpha,2\alpha,3\beta)$ -3-amino-1,2-cyclopentanediol hydrochloride (6)¹² was neutralized and allowed to react with 4,6-dichloro-5-(2,2-diethoxyethyl)pyrimidine (7)¹³ in the presence of triethylamine to give an intermediate (assumed to be 8). This latter product was not fully characterized but was cyclized to (\pm) - $(1\alpha,2\alpha,3\beta)$ -3-(4-chloro-7*H*-pyrrolo[2,3-*d*]pyrimidin-7-yl)-1,2-cyclopentanediol (9)¹⁴ upon treatment with acid. Ammonolysis of 9 yielded the desired (\pm) - $(1\alpha,2\alpha,3\beta)$ -3-(4-amino-7*H*-pyrrolo[2,3-*d*]pyrimidin-7-yl)-1,2-cyclopentanediol (5).¹⁵

Using AD-169 and Davis strains in HEL cells, 7,17 5 did not show any anti-HCMV activity (MIC₅₀ \geq 100 μ g/mL) and displayed no appreciable effect on the growth viability (MIC₅₀ 130 μ g/mL) in the host cells. As expected, in view of these results, 5 was not inhibitory towards AdoHcy hydrolase (IC₅₀ > 100 μ M) under conditions where neplanocin A, carbocyclic 3-deazaadenosine (C-c³Ado), 3-deazaneplanocin A and (S)-DHPA proved inhibitory to the enzyme (IC₅₀ 0.0072, 0.018, 0.031, and 0.28 μ M, respectively).

The effect of compound 5 on a number of other viruses was evaluated and no activity was found against (MIC₅₀ in µg/mL, host cell line): HSV-1 (KOS), TK- HSV-1 (B2006), HSV-2 (G), vaccinia, and vesicular

stomatitis (> 200, E₆SM); vesicular stomatitis, polio-1, and coxsackie B4 (> 200, HeLa); respiratory syncytial (Long) (> 100, HeLa); coxsackie B4, reo-1, Semliki forest, and Sindbis (> 200, Vero); parainfluenza 3 (VR-93) (150, Vero); influenza A (Ishikawa) and influenza B (Singapore) (> 100, MDCK); and, HIV-1 (III_B/LAI) and HIV-2 (ROD) (> 130, MT-4). Interestingly, 5 did show some activity towards Junin and Tacaribe (MIC₅₀ in μg/mL, 84 and 64, respectively, in Vero cells).

In analyzing the toxicity properties of 5 towards cell lines other than HEL, the following MIC₅₀ values were obtained: morphology $E_6SM(>200)$, morphology HeLa (>200), morphology Vero (≥ 200), morphology MDCK (≥ 100), and growth/viability MT-4 (> 130).

The virus strains, cell systems, and assays used to determine antiviral activity and cytotoxicity have been described previously.^{7,17-20} In all of these assays the reference compounds (BVDU, ribavirin, neplanocin A, and carbocyclic 3-deazaadenosine) showed the expected antiviral activity.⁷

From these results, it is clear that removal of the hydroxmethyl side-chain from 7-deazaaristeromycin has a significant impact on its antiviral properties when considered in parallel with the same modification on the parent aristeromycin.⁸

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- 14. Compound 9: off-white solid; mp 159-161 °C, ¹H NMR (DMSO- d_6) δ 1.1–2.6 (m, 4 H, H-4' and H-5'), 3.3 (s, 1 H, OH), 4.0 (s, 1 H, OH), 4.3 (m, 1 H, H-3'), 4.5-5.2 (m, 2 H, H-1' and H-2'), 6.7 (d, J = 3.8 Hz, 1 H, H-5), 7.9 (d, J = 3.8 Hz, 1 H, H-6), 8.6 (s, 1 H, H-2); ¹³C NMR (DMSO- d_6) δ 27.2, 29.9, 60.9, 72.5, 77.9, 98.6, 99.6, 130.7, 149.5, 150.5, 150.9.
- 15. Compound 5¹⁶: off-white solid; mp 205-207 °C, ¹H NMR (DMSO- d_6) δ 1.1-2.6 (m, 4 H, H-4' and H-5'), 4.2 (m, 1 H, H-3'), 4.4-5.7 (m, 4 H, H-1', H-2', and 2 x OH), 6.8 (d, J = 2.9 Hz, 1 H, H-5), 7.5 (d, J = 2.9 Hz, 1 H, H-6), 7.9 (s, 1 H, H-2), 8.2 (s, 2 H, NH₂); ¹³C NMR (DMSO- d_6) δ 28.5, 30.1, 60.3, 72.4, 78.5, 101.5, 104.0, 124.1, 148.0, 150.0, 154.2.
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