

Discovery and Synthesis of C-Nucleosides as Potential New Anti-HCV Agents

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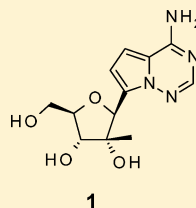
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Supporting Information

ABSTRACT: Nucleoside analogues have long been recognized as prospects for the discovery of direct acting antivirals (DAAs) to treat hepatitis C virus because they have generally exhibited cross-genotype activity and a high barrier to resistance. C-Nucleosides have the potential for improved metabolism and pharmacokinetic properties over their N-nucleoside counterparts due to the presence of a strong carbon–carbon glycosidic bond and a non-natural heterocyclic base. Three 2′CMe-C-adenosine analogues and two 2′CMe-guanosine analogues were synthesized and evaluated for their anti-HCV efficacy. The nucleotide triphosphates of four of these analogues were found to inhibit the NSSB polymerase, and adenosine analogue **1** was discovered to have excellent pharmacokinetic properties demonstrating the potential of this drug class.

KEYWORDS: C-Nucleoside, HCV, NSSB polymerase



EC₅₀ (cross genotype) = 0.4 - 1.6 μM
CC₅₀ > 100 μM

Dog PK: t_{1/2} = 7.6 h, F% = 96

Hepatitis C virus (HCV) is a major cause of chronic viral hepatitis that often leads to liver cirrhosis and hepatocellular carcinoma.^{1,2} In 2011 the protease inhibitors boceprevir and telaprevir became available to treat HCV infection with genotype 1 in combination with ribavirin and pegylated interferon.^{3,4} The recent approval of two new direct acting antivirals (DAAs), simeprevir and sofosbuvir,⁵ will significantly enhance the available armory, but there remains a need for new DAAs for use in safe and effective combination therapies.

Nucleoside and nucleotide inhibitors of HCV (NIs) are a class of DAA that tend to demonstrate broad activity across HCV genotypes and a high barrier to the emergence of viral resistance. Structural modifications have been made to both the carbohydrate (“sugar”) and heterocyclic base components of natural nucleosides to develop potent and selective antiviral analogues.⁶ The early promise of 2′CMe-7-deaza-adenosine (Figure 1) and, more recently, the excellent clinical success of sofosbuvir have validated the use of NIs as anti-HCV DAAs.^{5,7,8}

Structural analogues that have attracted relatively little attention to date for application to HCV are C-nucleosides,^{9–14} which comprise a sugar moiety and non-natural heterocyclic base connected by a carbon–carbon bond. This strong glycosidic bond makes C-nucleosides more resistant to enzymatic and hydrolytic cleavage than their N-nucleoside counterparts. This feature, combined with the broad structural

variation possible in the heterocyclic base, offers the potential for improved drug metabolism and pharmacokinetic properties. As part of an effort to investigate new nucleoside agents for HCV, C-nucleoside adenosine analogues **1–3** and guanosine analogues **4** and **5** were targeted. Reported herein are the synthesis and biological characterization of these compounds (Figure 1).

Pyrrolo[2,1-*f*][1,2,4]triazine-4-amine adenosine analogue, **1**, was first synthesized using the linear route shown in Scheme 1. This is a highly modified approach to that used by Patil et al., for the synthesis of the unsubstituted ribo-analogue (4-aza-7,9-dideazaadenosine).¹⁵ The 3,5-bis-dichlorobenzyl protected 2′CMe-ribose **7** was protected at the 2′-position to afford **8**, then hydrolyzed under acidic conditions to afford **9**. The key glycosylation reaction was performed using a procedure adapted from a method reported for the reaction of pyrroles with O-glycosylimidates.¹⁶ Riboside **9** was converted to the activated trichloroacetimidate **10**. Slow addition of pyrrole to **10** at low temperature in the presence of BF₃·Et₂O afforded an anomeric mixture of the pyrrolo-nucleoside, **11**. Modest improvement in β-selectivity was achieved at higher temper-

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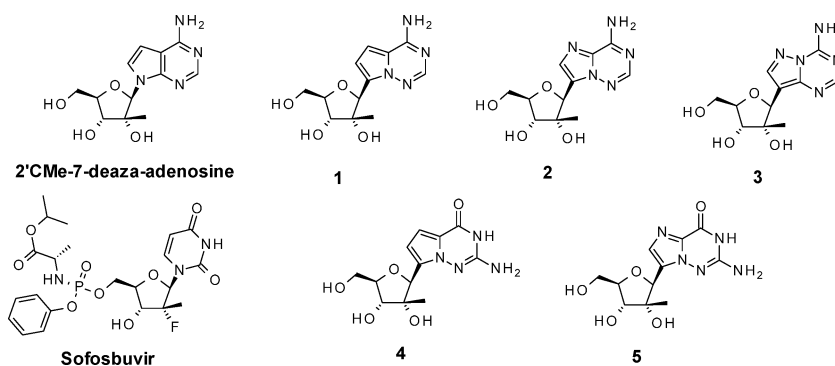
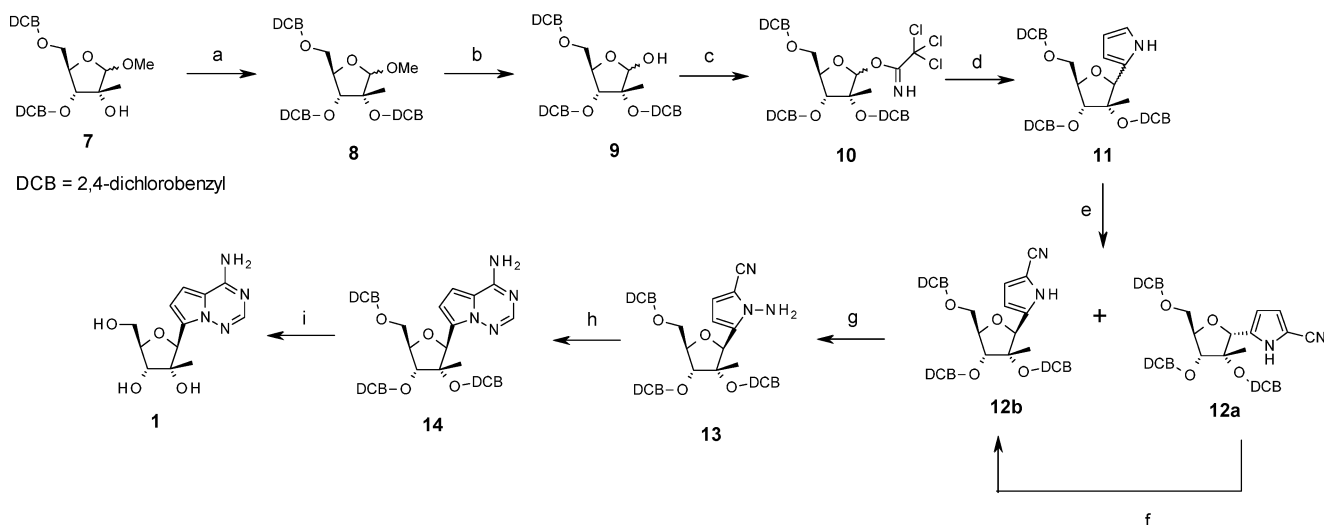


Figure 1. Structures of sofosbuvir, 2'CMε-7-deaza-adenosine, and target C-nucleoside analogues.

Scheme 1. Synthesis of 1^α



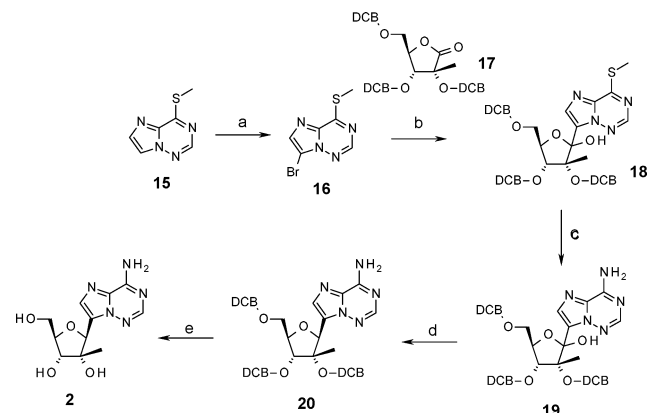
^αReagents and conditions: (a) NaH, 2,4-dichlorobenzyl chloride; (b) TFA/water 9:1; (c) Cl₃CCN, Cs₂CO₃; (d) (i) 4 Å mol. sieves, DCM, 2 h, rt, (ii) pyrrole, BF₃·OEt₂, -50 °C (60% yield, β/α 2:1) or -78 °C (80% yield, β/α 1:1), (iii) NH₃ in MeOH, -78 °C; (e) ClSO₂NCO, CH₃CN/DMF, 0 °C; (f) BF₃·OEt₂, DCM, reflux; (g) (i) NaH, THF, 0 °C, (ii) Ph₂P(O)ONH₂, THF, 0 °C; (h) formamidine acetate, DMA, 140 °C, 1–2 h; (i) H₂, Pd–C (10%), 45 °C, 18 h, NaOAc, MeOH, HOAc.

ature (-50 °C, α/β = 1:2, 60% yield), but at the cost of yield (-78 °C, α/β = 1:1, 80% yield). Because pyrrolo-nucleoside 11 decomposed slowly on standing, it was immediately converted to the more stable pyrrolonitrile nucleoside 12 by reaction with chlorosulfonyl isocyanate in DMF. The α/β mixture was separated at this stage by flash chromatography, and more of the desired β-anomer (12b) recovered after BF₃·Et₂O catalyzed anomerization of the purified α-anomer (12a). The β-stereochemistry was established at this stage by the observation of strong NOE between the H4' and the H1' protons. Electrophilic amination of the pyrrole nitrogen, ring closure with formamidine acetate, followed by buffered hydrogenolysis afforded the free adenosine analogue 1.

An analogous linear route toward the pyrrolo[2,1-f][1,2,4]-triazine-4-amine guanosine analogue 4 was attempted. Evidence was obtained that this may be achieved via hydrolysis of the 2,6-diamino analogue of 1 under strongly basic conditions at high temperature. However, this approach was abandoned in favor of the convergent synthesis described below.

Imidazo[2,1-f][1,2,4]triazine-4-amine adenosine analogue, 2, was synthesized using a convergent approach, which utilized the direct coupling of a bicyclic heterocycle, as shown in Scheme 2. After selective bromination of 15, the bromo-base 16 was coupled with ribolactone 17. Amine displacement of the

Scheme 2. Convergent Synthesis of A-Analogue 2^α



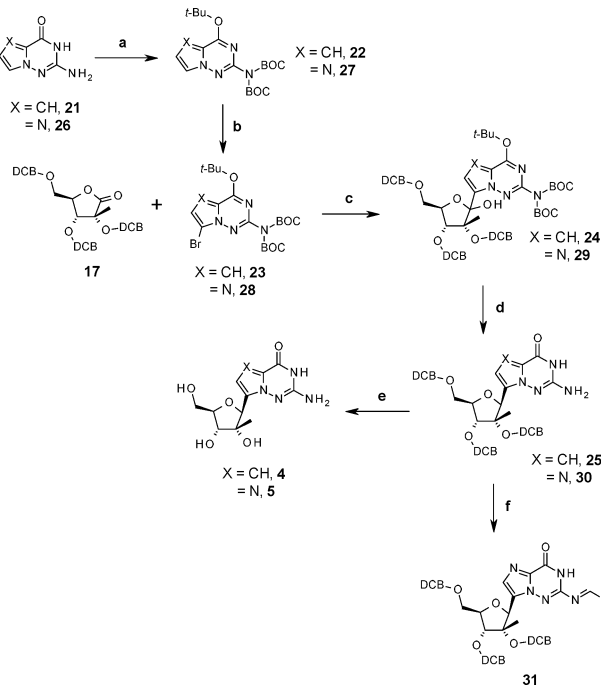
^αReagents and conditions: (a) NBS, DMF, 86 °C, 1 h; (b) (i) *n*BuLi, -78 °C, THF, (ii) 17, -78 °C; (c) NH₃ in MeOH, RT to reflux; (d) BF₃·OEt₂, Et₃SiH, DCM, -78 °C to RT, 3 h; (e) H₂, Pd–C (10%), 60 °C, 18 h, NaOAc, MeOH/DCM (9:1), AcOH.

thiomethyl group followed by selective lactol reduction yielded the desired β-nucleoside 20 and subsequently the free nucleoside 2 after hydrogenolysis. A similar convergent

synthesis of the pyrrolo analogue **1** was developed for large scale synthesis. Convergent methods for **1** and **2** have also been reported by others since we completed this work.¹²

Guanosine analogues **4** and **5** were prepared using an analogous approach, as shown in Scheme 3. Compounds **21**

Scheme 3. Convergent Synthesis of G-Analogues **4 and **5**^α**

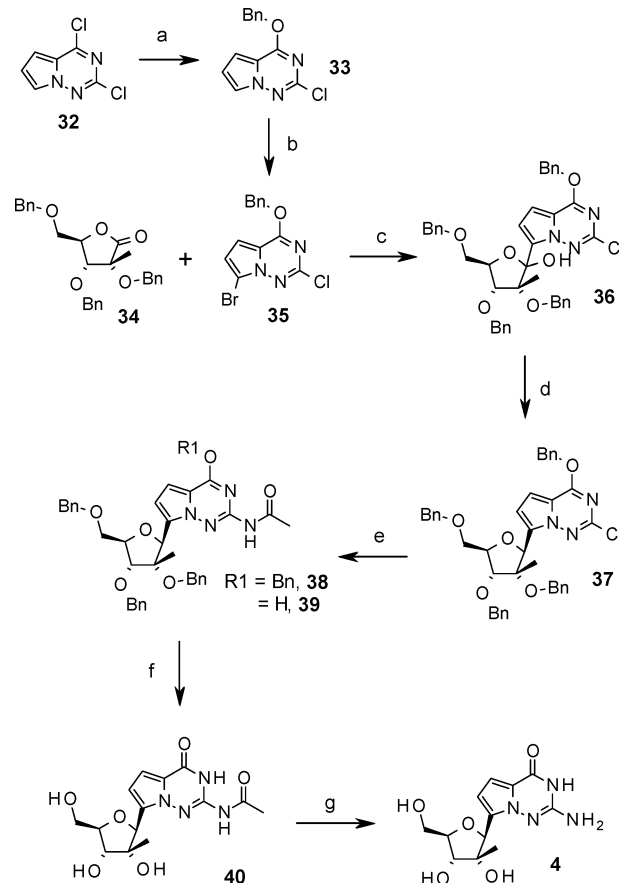


^αReagents and conditions: (a) TEA, DMAP, Boc₂O, MeCN, RT, 46 h; (b) NBS, DCE, -10 to 0 °C, 1 h; (c)(i) *n*BuLi, THF -100 °C, **23** or **28** (ii) **17**; (d) BF₃·OEt₂, Et₃SiH, MeCN, -78 °C to RT, 3 h; (e) H₂ (60 psi) Pd-C (10%), 60 °C, 18 h, NH₄OAc, MeOH/EtOAc (13:2) or BBr₃, -78 °C to -30 °C (**30** to **5**); (f) DMF-dimethylacetal, DCM/MeCN, RT to 60 °C.

and **26** were chosen as appropriate starting heterocycles. Under standard conditions for Boc-protection, N-Boc protection was achieved for both starting materials, together with *t*Bu-protection of the 4-oxygen, a surprising transformation that has been observed for other guanosine analogues.¹⁷ After selective bromination, the direct glycosylation of the bromo-bases **23** and **28** with the lactone **17** was achieved at very low temperature (-100 °C). It was noted that the anions of these guanosine base analogues were much less stable than the anions of the corresponding adenosine base analogues. The coupling thus required careful, slow addition of the lactone to the organolithium base to maintain a low temperature and minimize anion quenching. The protecting groups on the base were removed in the subsequent stereoselective lactol reduction to afford DCB-protected β -nucleosides **25** and **30**. The β -stereochemistry of **25** was established at this stage by the observation of an interaction between H1' and H4' in the NOESY spectrum. Such unambiguous signals were not observed for **30**; so, the formamidine analogue **31** was prepared under mild conditions. The observation of a strong interaction between H1' and H4' in the NOESY spectrum of this compound provided evidence of β -stereochemistry. The free guanosine nucleoside analogues **4** and **5** were obtained upon hydrogenolysis of the 2,4-dichlorobenzyl groups under buffered conditions.

Because of the anion instability of the intermediate organolithium base, formed from **23**, the synthesis of **4** was only possible in small, low milligram quantities using the method outlined in Scheme 3. A more robust procedure was therefore developed, which is shown in Scheme 4. The

Scheme 4. Improved Synthesis of **4^α**



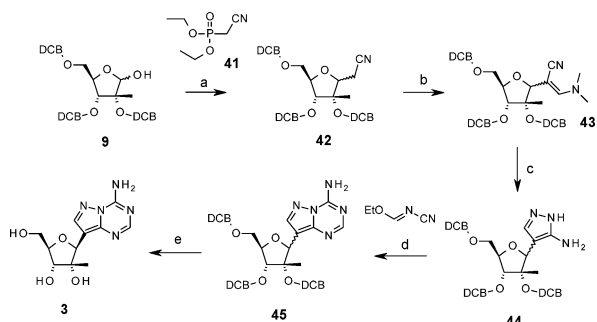
^αReagents and conditions: (a) *n*BuLi, BnOH, -10 °C to RT, THF; (b) NBS, DCM, -10 °C to RT; (c)(i) *n*BuLi, -100 °C, 2-Me-THF, (ii) **34** containing C7 isomer, -100 °C to RT; (d) BF₃·OEt₂, Et₃SiH, DCM, -78 °C, 15 min; (e) acetamide, Pd₂(dba)₃, xantphos, Cs₂CO₃, 130 °C, 1 h; (f) H₂, MeOH, Pd-C (10%), RT to 50 °C, 70 h; (g) NaOMe, RT to 80 °C.

heterocycle **35** was chosen for glycosidic coupling and subsequent transformation to nucleoside **4** based on observations made during the initial synthesis of **4**. These included the relative instability of anions on heterocycles bearing a protected nitrogen at the 2-position, the requirement for O4 protection to enable good metalation, and the susceptibility of the 2,4-dihalides to uncontrolled hydrolysis. Heterocycle **35** was prepared from the bis-chloro analogue **32** by displacement of the 4-chloro with BnOLi followed by bromination with NBS in DCE to yield an inseparable mixture of **35** along with its C7 isomer (3:1). As the C7 isomer was found to be inert in the glycosylation reaction, the mixture was used without separation. Glycosylation of the per-benzylated lactone **34** was achieved in 60% yield based on **35**. After selective reduction of the lactol **36** to afford β -nucleoside **37**, the 2-amino was installed by applying cross-coupling chemistry,¹⁸ which also resulted in partial deprotection of the 4-oxygen to afford a mixture of **38** and **39**. The benzyl groups were cleaved under hydrogenolysis and

the acetamide group cleaved under basic conditions to afford the free guanosine nucleoside **4**.

Application of the convergent chemistry used for the pyrrolo[2,1-*f*][1,2,4]triazine and imidazo[2,1-*f*][1,2,4]triazine bases was unsuccessful for the synthesis of pyrazolo[1,5-*a*]-1,3,5-triazine adenosine **3**. Scheme 5 shows its successful linear

Scheme 5. Synthesis of Pyrazolo-Nucleoside 3^α



^αReagents and conditions: (a) NaH, DME, **41**, 0 °C to RT; (b) (CH₃)₃COCH[N(CH₃)₂]₂, DMF, 60 °C, 15 h; (c) NH₂NH₂·HCl, EtOH, 105 °C, 2 h; (d) ethyl *N*-cyanoformimidate, 85 °C, toluene; (e) H₂, MeOH, Pd-C (10%), 45 °C, 17 h.

synthesis, which adopts similar methodology reported by Klein et al. for the synthesis of the unsubstituted riboside analogue.^{19,20} Condensation of 2-diethoxyphosphoryl-acetonitrile, **41**, with the tris-benzyl riboside **9** afforded the C-nucleoside intermediate **42**. Separation of the anomers was possible at this stage but unnecessary, as interchange between anomers was observed under reaction conditions for the dimethylamine adduct **43** and for the amino-pyrazole intermediate **44**. In situ condensation of **44** with ethyl cyanoimidoformate afforded the protected nucleoside **45**. The anomers were separated, and the β-stereochemistry was established by the observation of an interaction between H1' and H4' in the NOESY spectrum. The pyrazolotriazine adenosine analogue **3** was obtained after hydrogenolysis.

The in vitro anti-HCV results for the parent C-nucleosides **1**–**5** are shown in Table 1. The low IC₅₀ values observed for the pyrrolo-triazine and imidazo-triazine adenosine and guanosine nucleoside triphosphates show they are well

Table 1. Inhibitory Potency (EC₅₀) of Nucleosides on HCV RNA Replication in Replicon Assay; Cytotoxicity (CC₅₀); and Inhibitory Potency (IC₅₀) of Corresponding Nucleoside-Triphosphate (NTP) on NSSB Polymerase Enzyme (WT GT1b(Con1))

compd	GT1b EC ₅₀ (μM) ^a	CC ₅₀ (μM) ^b	NTP IC ₅₀ (μM) ^c
2'CMe-7-deaza-A	0.20, 0.3 (lit.) ²¹	>100, >100 (lit.) ²¹	0.20, 0.07 (lit.) ²¹
pyrrolo-A (1)	1.6, 1.98 (lit.) ¹²	>100, 85 (lit.) ¹²	0.37, 0.31 (lit.) ¹²
pyrrolo-G (4)	>100	>100	0.21
imidazo-A (2)	1.3, 1.28 (lit.) ¹²	34, >89 (lit.) ¹²	0.55, 2.1 (lit.) ¹²
imidazo-G (5)	>100, >89 (lit.) ¹²	>100, >89 (lit.) ¹²	0.31, 0.19 (lit.) ¹²
pyrazolo-A (3)	>100	>100	33

^aMeasured using subgenomic replicon cell lines described by Blight et al.²² See Supporting Information for methods. ^bMeasured using Huh-7 derived replicon cell lines with MTT as the vital dye indicator. ^cSee Supporting Information for methods.

recognized by the NSSB enzyme. The adenosine analogues **1** and **2** were furthermore active in the replicon assay, although the imidazo-triazine analogue, **2**, also showed some cytotoxicity. The corresponding guanosine analogues **4** and **5** were inactive in the cell-based replicon assay, a result consistent with the relatively poor activity reported for N-nucleoside guanosine analogues 2'CMe-G (EC₅₀ = 2–4 μM) and 7-deaza-2'CMe-G (EC₅₀ > 50 μM).^{7,21} The activity of the corresponding NTP suggests they were not efficiently phosphorylated by host kinases. In contrast, the poor enzyme activity observed for the pyrazolotriazine adenosine analogue, **3**, is consistent with its poor activity in the replicon assay. On the basis of these initial results, **1** was selected for further investigation as a potential HCV DAA.

Compound **1** displayed excellent activity against HCV genotypes 1a, 2a, 2b, 3a, 4a, 5a, 6a, and 7a. EC₅₀ and IC₅₀ values ranged from 0.39 to 1.0 μM (see Supporting Information for details) demonstrating its potential suitability as part of a pan-genotypic HCV therapy.

Because the nucleotide triphosphate is the compound responsible for anti-HCV activity, it is important to measure the ability of relevant cells to anabolize the parent nucleoside. The in vitro anabolism of **1** was studied in adherent Huh-7 cells and plated primary human hepatocytes. Following incubation of Huh-7 cells with 10 μM initial concentration of **1** for 24 h, the intracellular concentration of **1**-NTP reached 9.3 pmol/million cells. Surprisingly, primary human hepatocytes generated much higher (~50-fold) intracellular levels of **1**-NTP (487 pmol/million cells) after 24 h with the same starting concentration, offering the promise of enhanced HCV antiviral activity in humans. Intracellular levels of **1** were found to be comparable in Huh-7 cells and hepatocytes, but the mono- and diphosphate species were about 10-fold lower in the replicon cell line suggesting that the first and second phosphorylation steps are more efficient in hepatocytes. The decay half-life (*t*_{1/2}) of **1**-NTP displays a biphasic profile in both replicon cells and primary human hepatocytes: a rapid initial decay *t*_{1/2} (0.7 and 3.3 h, respectively), followed by a prolonged elimination phase (*t*_{1/2} ≥ 35 h).

The pharmacokinetic properties of **1** were determined in male Sprague–Dawley rats, beagle dogs, and cynomolgus monkeys. Compound **1** displayed excellent cross-species pharmacokinetics with elimination *t*_{1/2} ranging from 4 to 7.6 h and high bioavailabilities (33–96%). On the basis of these data, the drug may have sufficient PK properties for a once daily dosing regimen in humans taking into account the long intracellular half-life of its triphosphate in hepatocytes.

Compound **1** was evaluated in a range of in vitro safety assays to assess its suitability for progression into animal toxicity studies. In a mitochondrial toxicity study over 14 days in HepG2 cells, **1** showed no increase in lactate production or decrease in mtDNA up to 10 μM. Low levels of cytotoxicity were observed in rat cells (CC₅₀ = 190 μM for liver epithelial cells and CC₅₀ > 250 μM for kidney fibroblasts and heart myoblasts). Compound **1** did not inhibit erythroid and myeloid progenitor proliferation in human bone marrow cells up to 100 μM and was negative in an AMES mutagenicity test. In addition, compound **1** triphosphate did not inhibit human DNA polymerases (IC₅₀ >100 μM for α, β, and γ enzymes). Despite this promising in vitro profile, unacceptable adverse effects, including death, were observed during single oral dose toxicity studies in rats (dosed up to 400 mg/kg). This result caused us to focus development efforts on derivatives of **1** with

a potentially improved safety margin. Since the completion of this work, the results of safety pharmacology and toxicology studies of **1** have been published.¹¹

In summary, the synthesis and anti-HCV activity of a series of C-nucleoside purine analogue inhibitors of HCV NSSB polymerase have been investigated. Four of the five nucleosides showed excellent enzyme activity as their triphosphates, with two adenosine analogues showing activity in the cell-based replicon assay. The enzyme active guanosine analogues were, however, inactive in the whole cell assay, presumably due to inefficient anabolism to their nucleotide triphosphates. The pyrrolotriazine adenosine analogue **1** was chosen for further profiling because it showed excellent cross-genotype activity, low in vitro toxicity, good anabolism, and excellent pharmacokinetic properties in three species. Compound **1** incorporates the key structural features of a 2'-CMe-ribose moiety with a pyrrolotriazine 7-deazaadenine base mimic, connected by a highly stable C–C bond. Unfortunately, the observation of adverse effects in rat safety studies caused us to re-evaluate our pursuit of **1** as a drug candidate. Despite this, these results demonstrated the potential for C-nucleosides to be used in new antiviral HCV therapies provided a sufficient therapeutic margin can be established.

■ ASSOCIATED CONTENT

■ Supporting Information

Experimental procedures, analytical data, and the description and results of in vitro assays and in vivo pharmacokinetics. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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Author Contributions

All authors have given approval to the final version of the manuscript.

Notes

The authors declare no competing financial interest.

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