Supporting information for

The Design, Synthesis and Anti-viral Activity of Monofluoro and

Difluoro Analogues of 4'-Azidocytidine Against Hepatitis C Virus

Replication: The Discovery of 4'-Azido-2'-deoxy-2'-

fluoroarabinocytidine and 4'-Azido-2'-dideoxy-2',2'-difluorocytidine.

David B. Smith,[†] Genadiy Kalayanov,[§] Christian Sund,[§] Anna Winqvist,[§] Tatiana Maltseva,[§]

Vincent J-P. Leveque,[†] Sonal Rajyaguru,[†] Sophie Le Pogam,[†] Isabel Najera,[†] Kurt Benkestock,[§]

Xiao-Xiong Zhou,[§] Ann C. Kaiser,[†] Hans Maag,[†] Nick Cammack ,[†] Joseph A. Martin,[†] Steven Swallow,[†] Nils Gunnar Johansson,[§] Klaus Klumpp,[†] and Mark Smith*[†]

[†]Roche Palo Alto LLC, 3431 Hillview Avenue, Palo Alto, CA 94304, USA.

[§]Medivir AB, Department of Medicinal Chemistry, Lunastigen 7, SE-141, 44 Huddinge, Sweden.

*Corresponding author. Tel.: +1 650 855 5076; fax: +1 650 852 1132; e-mail Mark.Smith@Roche.com

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- S3 Analytical methods
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Biological methods

The HCV replicon assay was performed in the stable replicon cell line 2209-23 derived from Huh-7 cells stably transfected with a bicistronic HCV replicon (genotype 1b) expressing the Renilla luciferase reporter gene, as described.^{1,2}

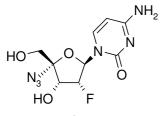
Analytical methods.

The analytical RP-HPLC system consisted of Waters 2695 Alliance separation module, Waters 996 photodiode array detector, and Micromass ZQ2000 mass detector (operated in +ESI). The columns used were an Atlantis dC18, 3×150 mm, 3μ m, 100A from Waters and Hypercarb, 50×3 , 3 µm, from Thermo Electron Corporation. The mobile phases were based on water/acetonitrile containing 5 mM ammonium acetate. LC-MS accurate mass measurements were performed using a HDMS Synapt instrument from Waters (UK) equipped with a lockspray interface, connected to a Waters Aquity system. The acquisition range was m/z 100 to 1000 with an acquisition time of 0.15 s (+ESI). Leucine enkephalin was used as lock mass. The reversed phase column was an YMC-UltraHT Pro C_{18} , 2.1 × 50 mm, 2µm, 120A from YMC (U.S.A) and the mobile phases were based on water/acetonitrile containing 0.2% formic acid. 1 H - and 13 C -NMR experiments were carried out on Varian spectrometer (UNITY INOVA) at magnetic field strength of 11.7 T operating at 499.84 MHz for ¹H and 125.67 MHz for ¹³C, unless otherwise stated. The spectrometer was equipped with ¹H, ¹³C, ¹⁵N 5 mm Indirect detected Cryo Probe. ¹H and ¹³C pulses were applied with 36.8 kHz and 15.7 kHz field, respectively. ¹³C decoupling was performed using GARP with 8.8 kHz field strength. To avoid the spinning artifacts, all spectra were measured on non-spinning samples. All experiments have been done at temperature 25 °C. The assignment of the ¹H - and ¹³C resonances have been based on homonuclear 2D COSY and NOESY experiments as well as on inverse heteronuclear experiments, gHSQC and gHMBC. For COSY and NOESY experiments, the data sets were recorded as 2 K × 256 real matrix with 4 (and 16) scans for each t₁ value and a spectral width of 14 ppm. For gHSQC and gHMBC, the

data sets were recorded as 2 K \times 256 real matrix with 4 scans for each t₁ value and a spectral width of 14 ppm in F2 and 230 ppm in F1 with the carrier 6 and 125 ppm, respectively. In all cases the recycle delay used was 2.0 s.

Compound	Mw (Da)	Purity (%)
10	286.1	98.9
17	286.1	99.6
20	286.1	94.2
28	286.1	97.6
34	305.1	99.8

1-(4'-Azido-2'-deoxy-2'-fluoro-β-D-ribofuranosyl)cytosine (10).





Data File C:\EZXDATA\GENADIY\11-08\201108-MSV12111-03374.D Sample Name: MSV121

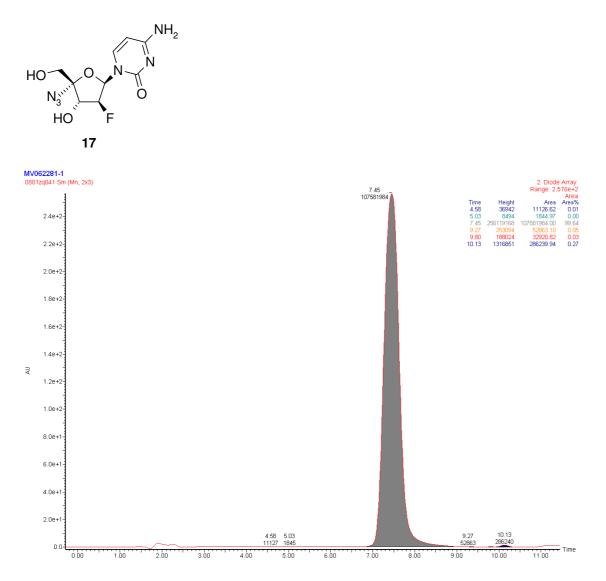
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Acq. Instrument		
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		Inj Volume : 5 µl
Acq. Method		C:\Chem32\1\METHODS\ROCHE5.M
Last changed	:	2008-11-20 12:47:10 by Genadiy Kalayanov (modified after loading)
Analysis Method	÷	C:\Chem32\1\METHODS\PN0599A3.M
Last changed		2008-11-18 11:26:26 by Christian Sund
2		(modified after loading)
Method Info	3	STANDARD METHOD (pos and neg ionizaion); 5 to 99 % ACN in 3 min
		Flow: 0.8 ml/min. UV=210-400 nm, ACE C8 3 * 50 mm
		Mobile phase A: 10 mM NH4Ac in 90% H2O, B: 10 mM NH4Ac in 90% ACN
Sample Info	:	a
1		Agilent Easy-Access Method: 'Roche 5 min'
		*** No target masses specified ***
	-30	5,190 Ref=off (C:\EZXDATA\GENADIY\11-08\201108-MSV12111-03374.D)
mAU 🗄	-30	up
		A
		/4
1000 -		
1000		
750		
750 - 500 -		8 / 5
750 500 250		1.358 4.613
750 - 500 -	÷	

Sample Name: MSV121

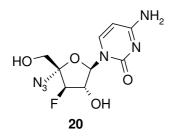
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Sorted By	: Signal							
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Use Multiplier &	Dilution Factor with ISTDs							
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Signal has been Peak RetTime Typ # [min]	modified after loading from rawdata file! • Width Area Height Area							
Signal has been Peak RetTime Typ # [min] 	modified after loading from rawdata file! • Width Area Height Area [min] [mAU*s] [mAU] % •							
Signal has been Peak RetTime Typ # [min] 1 1.358 BB	modified after loading from rawdata file! Width Area Height Area [min] {mAU*s] [mAU] %							

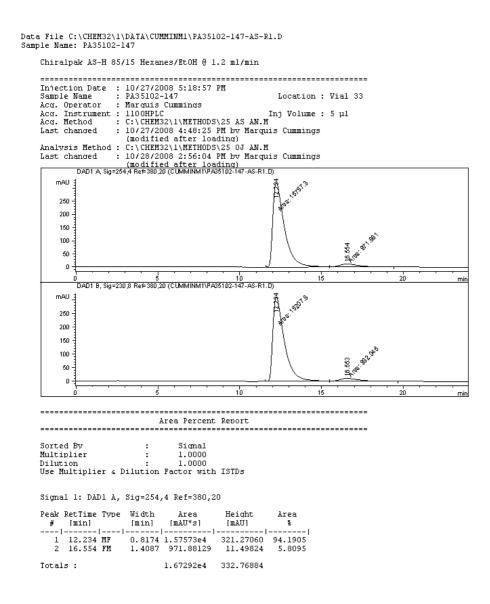
Totals : 1.52035e4 1467.01941

1-(4'-Azido-2'-deoxy-2'-fluoro-β-D-arabinofuranosyl)cytosine (17).



1-(4'-Azido-3'-deoxy-3'-fluoro-β-D-xylofuranosyl)cytosine (20).

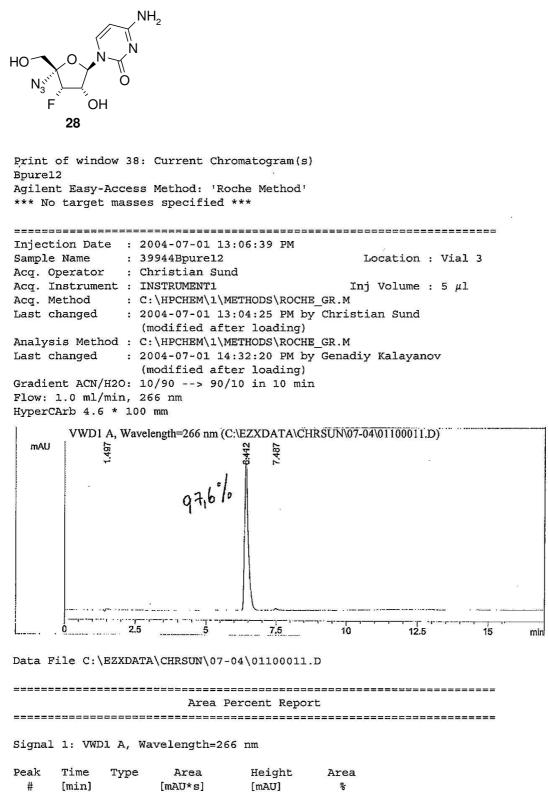




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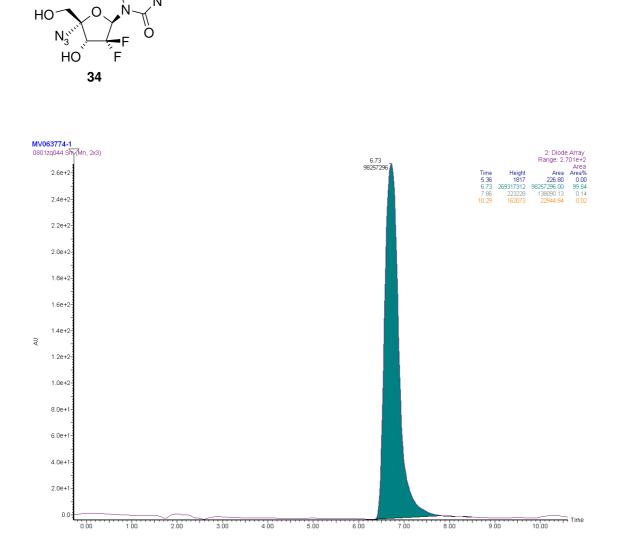
1-(4'-Azido-3'-deoxy-3'-fluoro-β-D-ribofuranosyl)cytosine (28)



1	1.497	PB	27.44311	2.81430	1.4286
2	6.412	BB	1874.70911	212.52504	97.5971
3	7.487	VB	18.71299	2.19273	0.9742

1-(4'-Azido-2'-dideoxy-2',2'-difluoro-β-D-ribofuranosyl)cytosine (34).

NH₂



References.

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