

Studies on acyl pyrrolidine inhibitors of HCV RNA-dependent RNA polymerase to identify a molecule with replicon antiviral activity

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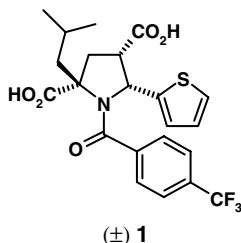
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Abstract—The SAR development is described for a series of *N*-acyl pyrrolidine inhibitors of the Hepatitis C virus RNA-dependent RNA polymerase, NS5B, from tractable $\Delta 21$ enzyme inhibitors to an example with antiviral activity in a cellular assay (HCV replicon).

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Hepatitis C virus (HCV), a positive strand RNA virus of the Flaviviridae family, is the major etiological agent of non-A, non-B hepatitis chronically infecting an estimated 2–3% of the world population.¹ Current treatments of HCV include interferon or pegylated interferon in combination with ribavirin which target stimulation of host defence mechanisms to result in viral clearance. Improvements in sustained response rate, which is presumed to be dependent on viral RNA load and genotype, are still needed.² Attempts to identify alternative options for the treatment of chronic hepatitis C virus infections are ongoing.



We recently described the identification of a series of *N*-acyl pyrrolidines (e.g. **1**) as inhibitors of the NS5B RNA-dependent RNA polymerase.³ A developable drug for the treatment of HCV must be able to penetrate infected cells to inhibit replication of the virus. In addition, favourable in vivo distribution, metabolism and pharmacokinetic properties are essential for efficacy in the patient.

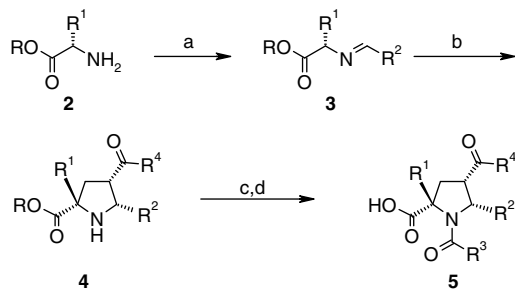
A crystal structure of NS5B of genotype 1b, isolate J4, with 21 amino acid C-terminal truncation ($\Delta 21$) showed that (+) **1** binds in the palm region of the polymerase enzyme.⁴ This supports the hypothesis that the polymerase enzyme inhibition of these *N*-acyl pyrrolidines is through binding in a specific pocket and not via a promiscuous or non-specific mechanism. The chemistry efforts described here were undertaken to improve the potency and replace one of the carboxylic acid functions to address the poor cell penetration and poor 'drug-like properties' of these tractable hit diacidic molecules. For efficiency, these two objectives were pursued in parallel.

Initially analogues were assayed for their inhibitory activity against the recombinant HCV NS5B enzyme of genotype 1b, isolate J4, with 21 amino acid C-terminal truncation ($\Delta 21$) using the biochemical assay described previously.⁵ As the programme progressed and

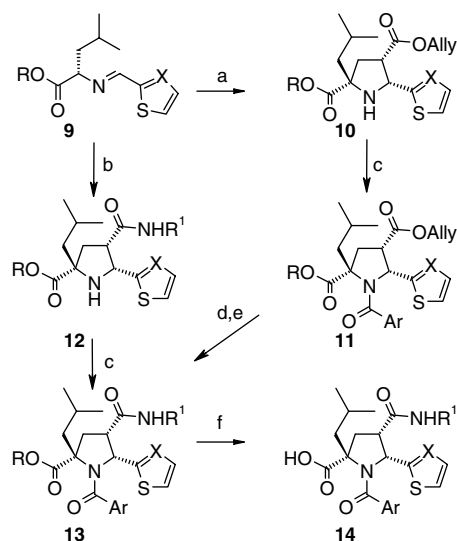
Keywords: Hepatitis C virus; RNA-dependent RNA polymerase; HCV NS5B; *N*-Acyl pyrrolidines; HCV replicon; SAR.

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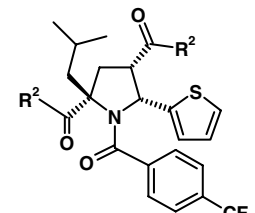
From initial studies varying the ester function, it rapidly became apparent that the 2-carboxylic acid is very sterically hindered and therefore difficult to modify. Construction of the pyrrolidine ring with the required 2-carboxyl substitution already in place on the amino acid avoided this difficulty. This strategy of introducing



Conversion of either, or both, of the carboxylic acid groups of (\pm) **1** to an ester resulted in the loss of inhibition of NS5B, see compounds **15–18** (Table 1). Thus it appeared necessary to retain hydrogen bond donors in the carboxylic acid replacements which suggested examining amides or primary alcohols. Although the primary amide at position 4 (**19**) or the corresponding 2-amide (**20**) showed reduced NS5B inhibition, they did retain some activity. However a series of secondary amides at position 2 all showed less enzyme inhibition than the primary amide, compounds **21–25**. The primary alcohol analogue, **7**, retained some weak activity (51% at 10 μ M) but **8** was inactive (0% at 10 μ M). These data suggest that any change to the 2-carboxyl group essentially suppresses inhibitory activity.



Scheme 3. Reagents and condition: (R = resin for solid-phase or *t*-Bu for solution synthesis): (a) allyl acrylate, AgNO₃ or LiBr, NEt₃, MeCN or THF; (b) acrylamide, AgNO₃ or LiBr, NEt₃, MeCN or THF; (c) R³COCl, base; (d) Pd^[0], PPh₃; (e) amine (R¹NH₂), HOAt, DIC, DIEA; (f) TFA.

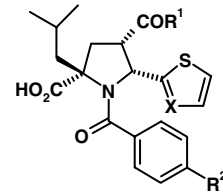
Table 1. NS5B inhibitory activity of *N*-acyl pyrrolidine esters


The chemical structure shows a pyrrolidine ring with a 4-tert-butylbenzoyl group at the 2-position and a 5-thiazolyl group at the 3-position. The nitrogen of the pyrrolidine is substituted with an acyl group R¹. The thiazole ring is substituted with an acyl group R² at the 4-position. A CF₃ group is attached to the benzene ring of the 4-tert-butylbenzoyl group.

Compound	R ¹	R ²	NS5B J4 Δ21 IC ₅₀ (μM)
1	OH	OH	0.3
15	OMe	OH	>20
16	OCH ₂ Ph	OH	>20
17	OH	OMe	>20
18	OMe	OMe	>20
19	OH	NH ₂	4.4
20	NH ₂	OH	8.1
21	NHMe	OH	>20
22	NHEt	OH	>20
23	NHCH ₂ Ph	OH	>20
24	NH <i>i</i> -Pr	OH	>20
25	NHPh	OH	>20

Meanwhile, studies to expand the SAR of the series had continued with examination of alternate 1-substitution. Benzyl and sulphonamide analogues of compound **1** were inactive while its urea derivative showed reduced potency (IC₅₀ = 2.1 μM).

More rewarding was the continued investigation of additional *N*-acyl groups and 5-heterocycles performed in a combinatorial manner with small arrays. This uncovered a synergistic effect from the combination of the 1-*tert*-butylbenzoyl and 5-thiazoyl groups **28** compared with **1**, **26** and **27** (Table 2). Indeed, when applied to the amide series this effect was also observed (**32** cf. **30** and **31**) and resulted in a significant advance for the study, identifying the mono-acid **32** with similar potency to its di-acid analogue **28** capable of showing measurable inhibition of full-length enzyme (NS5B RB01 FL).

Table 2. NS5B inhibitory activity of *N*-acyl pyrrolidines


The chemical structure shows a pyrrolidine ring with a 4-tert-butylbenzoyl group at the 2-position and a 5-thiazolyl group at the 3-position. The nitrogen of the pyrrolidine is substituted with a COR¹ group. The thiazole ring is substituted with an X group at the 4-position. A CF₃ group is attached to the benzene ring of the 4-tert-butylbenzoyl group.

Compound	X	R ¹	R ²	NS5B J4 Δ21 IC ₅₀ (μM)	NS5B RB01 FL IC ₅₀ (μM)
1	C	OH	CF ₃	0.33	20
26	N	OH	CF ₃	0.45	
27	C	OH	<i>t</i> -Bu	0.75	
28	N	OH	<i>t</i> -Bu	0.07	3.8
29	C	NH ₂	CF ₃	4.40	
30	C	NH ₂	<i>t</i> -Bu	0.33	
31	N	NH ₂	CF ₃	2.05	
32	N	NH ₂	<i>t</i> -Bu	0.08	4.4

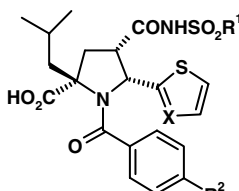
Separation of the racemic mixtures of thiazole di-acid **28** and acid amides **29** and **32** by chiral HPLC readily provided the pairs of enantiomers which exhibited data similar to that described for compound **1** where the inhibition resided in the (+) isomer (Table 3) and in agreement with the enantiomer bound in the enzyme crystal structure.¹⁰ Although this was a significant advance in enzyme potency, the amides demonstrated only marginal, if any, detectable cellular activity in line with low permeability similar to that seen for the di-acids (as measured in an artificial membrane permeability assay, <0.3 nm/s).¹¹

Following the progress described above, further investigation of substituted amides was considered. Initially amides which retained an acid proton such as acyl sulphonamides were examined. These acid isosteres retained a good level of enzyme inhibition, particularly when combined with the 4-*tert*-butylbenzoyl acyl group (Table 4). However, as these do not address the permeability issue of di-acids (artificial membrane permeability assay for compound **36**, <0.3 nm/s) so attention turned to examining non-acidic amide analogues.

A series of 4-*N*-substituted amide analogues of **32** was examined whilst keeping the other groups constant with the aim of improving cell permeability and retaining potency (Table 5). The benzyl **39** and methyl **40** amides showed a significant drop in potency; however the phenyl amide **41** retained significant inhibition of the Δ21 and

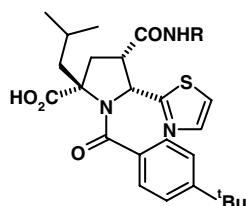
Table 3. NS5B inhibitory activity of separated enantiomers

Compound	NS5B J4 Δ21 IC ₅₀ (μM)	NS5B RB01 FL IC ₅₀ (μM)
(+) 1	0.30	11.9, 20
(−) 1	18	
(+) 28	0.03	2.1 (<i>n</i> = 6)
(−) 28	2.36	
(+) 29	2.23	
(−) 29	>30	
(+) 32	0.07	3.8 (<i>n</i> = 6)
(−) 32	2.50	

Table 4. NS5B inhibitory activity of *N*-acyl sulphonamides


The chemical structure shows a pyrrolidine ring with a 4-tert-butylbenzoyl group at the 2-position and a 5-thiazolyl group at the 3-position. The nitrogen of the pyrrolidine is substituted with a CONHSO₂R¹ group. The thiazole ring is substituted with an X group at the 4-position. A CF₃ group is attached to the benzene ring of the 4-tert-butylbenzoyl group.

Compound	X	R ¹	R ²	NS5B J4 Δ21 IC ₅₀ (μM)	NS5B RB01 FL IC ₅₀ (μM)
33	C	Me	CF ₃	0.23	
34	C	CF ₃	CF ₃	0.20	
35	N	Me	CF ₃	0.41	
36	N	Me	<i>t</i> -Bu	0.04	
37	N	Ph	<i>t</i> -Bu	0.50	3.15
38	N	CH ₂ Ph	<i>t</i> -Bu	0.55	4.23

Table 5. NS5B inhibitory activity of substituted amides

Compound	R	NS5B J4 $\Delta 21$ IC ₅₀ (μ M)	NS5B RB01 FL IC ₅₀ (μ M)	HCV replicon IC ₅₀ (μ M)
39	CH ₂ Ph	8.29	68	
40	Me		22	
41	Ph	0.16	3.1 ($n = 4$)	45 ^a , 77 ^a
(+) 41	Ph		0.5 ($n = 4$)	47 ^a , 30 ^a
(–) 41	Ph		>100	>200 ^a

^a CC₅₀ > 200 μ M, some cell toxicity seen at the highest dose.

full-length enzymes. Encouraged by detectable permeability in the artificial membrane permeability assay (2.2 nm/s), compound **41** was tested in the cell based HCV replicon assay. Indeed a weak antiviral effect could be detected which was judged sufficient to encourage further study.¹² Once again, after separation by chiral HPLC, the (+) enantiomer showed superior activity in both the enzyme and replicon assays (Table 5).

Medicinal chemistry studies towards an antiviral agent for the treatment of HCV infection are described. Hit molecule **1** had been identified from the follow-up of a high-throughput screening combinatorial mixture hit.³ The identification of a *N*-acyl pyrrolidine **41** which demonstrated an antiviral effect in the replicon cell based assay provided a lead for further medicinal chemistry optimisation to identify a candidate molecule for the treatment of HCV infections.¹²

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- The absolute stereochemistry of (+) **1** was determined by small molecule X-ray crystallography to be 2*S*, 4*S*, 5*R* (unpublished results).
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