

# HCV viral kinetic analysis predicts shorter treatment duration with AT-527, a purine nucleotide prodrug with potent pan-genotypic antiviral activity in HCV-infected subjects regardless of cirrhosis status

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## Background

AT-527 is a novel modified guanosine nucleotide prodrug inhibitor of the hepatitis C virus (HCV) NS5B polymerase, with a favorable/highly differentiated profile compared to sofosbuvir (1,2). A clinical trial evaluating AT-527 administered as a single agent for 7 days was successfully completed, allowing for the HCV viral kinetic analysis presented herein (3).

## Methods

Subjects and Data: HCV RNA data obtained from 30 subjects in the 7-day study who received active AT-527 doses were used for the viral kinetic analyses (3):

- Part C: Treatment-naïve (TN) non-cirrhotic (NC) subjects with genotype (GT) 1b HCV (n=6/dose) received AT-527 escalating doses of 138, 275 or 550 mg QD for 7 days (expressed as approximate free base equivalent).
- Part D: TN NC GT3 subjects (N=6) received the 550 mg dose QD for 7 days.
- Part E: TN cirrhotic GT1b (n=3), GT2 (n=1) or GT3 (n=2) subjects received the 550 mg dose QD for 7 days.

### Key Baseline Characteristics

	Part C – GT1b NC			Part D – GT3 NC	Part E – Cirrhotic
	138 mg (N=6)	275 mg (N=6)	550 mg (N=6)	550 mg (N=6)	550 mg (N=6)
HCV RNA, mean (range) log <sub>10</sub> IU/mL	5.8 (5.2-6.5)	6.2 (5.4-6.9)	6.1 (5.5-6.5)	6.5 (5.6-7.2)	6.4 (5.7-7.3)
Genotype 1b/2/3, n	6/0/0	6/0/0	6/0/0	0/0/6	3/1/2

### HCV RNA Samples:

- Serial samples were collected prior to (0 h on study day 1) and post treatment initiation at 2, 4, 8, 12, 16, 24, 36, 48, 72, 96, 120 and 144 h (days 1-7 during dosing) and daily after completion of dosing on days 8 to 13, as well as on day 35.
- HCV RNA levels were quantified using COBAS® AmpliPrep TaqMan® v2.0, with a lower limit of quantitation (LLQ) of 15 IU/mL (or 1.18 log<sub>10</sub> IU/mL).

### Viral Kinetic Modeling and Simulation:

- HCV viral load decline under AT-527 treatment and rebound after treatment completion was assumed to follow the standard model as described by the differential equations below (4,5):

$$\frac{dT}{dt} = s - dT - \beta VT$$

$$\frac{dI}{dt} = \beta T_0 V - \delta I$$

$$\frac{dV}{dt} = (1 - \epsilon)pI - cV$$

- Where  $T$  is the number of target hepatocytes susceptible to infection,  $I$  is the number of productively infected hepatocytes and  $V$  is the viral load. Susceptible hepatocytes are produced at rate  $s$  and die at rate  $d$ . *De novo* infection of cells occurs at rate  $\beta$  and the infected cells are lost at rate  $\delta$ . HCV virions are produced at rate  $p$  per hepatocyte and cleared at rate  $c$  per virion.
- Effectiveness of AT-527 in blocking viral replication is defined by  $\epsilon$  ( $0 \leq \epsilon \leq 1$ ). Upon dosing completion, where  $\epsilon = 0$ , viral load starts to rebound.
- Parameter estimates and associated interindividual variability were obtained using a maximum-likelihood method by the stochastic approximation expectation-maximization (SAEM) algorithm implemented in MonolixSuite 2019R1 (Lixoft, Orsay, France).
- Simulation was performed using Simulx for the 550 mg daily dose to estimate time to reach LLQ and cure. The latter is defined as HCV RNA < 1 IU in the entire extracellular body fluid (roughly 13.5 L):  $7.5 \times 10^{-5}$  or  $-4.13 \log_{10}$  IU/mL.

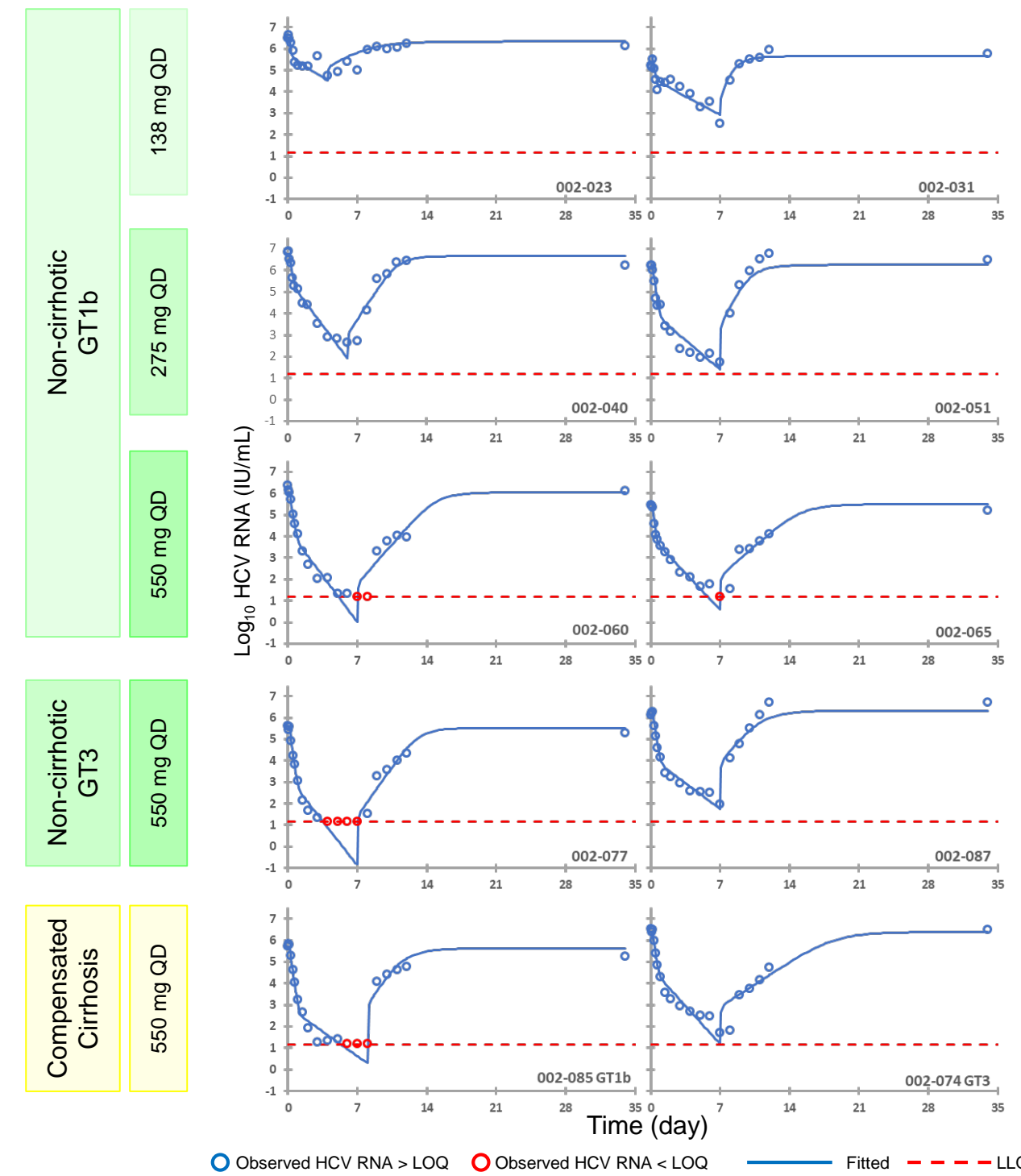
## Results

### Viral Kinetic Modeling

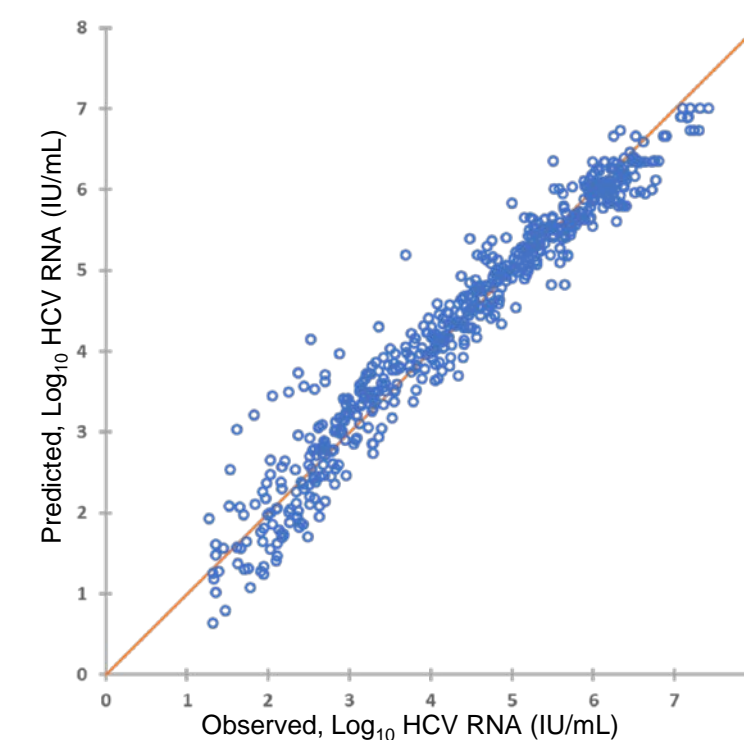
A model incorporating pharmacologic delay ( $\tau$ ) as well as AT-527 dose as a covariate of ( $\delta$ ) was found to best describe the observed data.

### Goodness-of-Fit

- Representative individual fitted vs. observed viral kinetics



### Predicted vs. observed HCV RNA



### Population Viral Kinetic Parameter Estimates

Parameter	Log <sub>10</sub> V <sub>0</sub> (IU/mL)	$\delta$ (d <sup>-1</sup> )	$c$ (h <sup>-1</sup> )	$\tau$ (h)	$\epsilon$
Estimate (±SE)	6.07±0.079	1.02±0.067	0.340±0.019	4.03±0.26	0.994±0.001

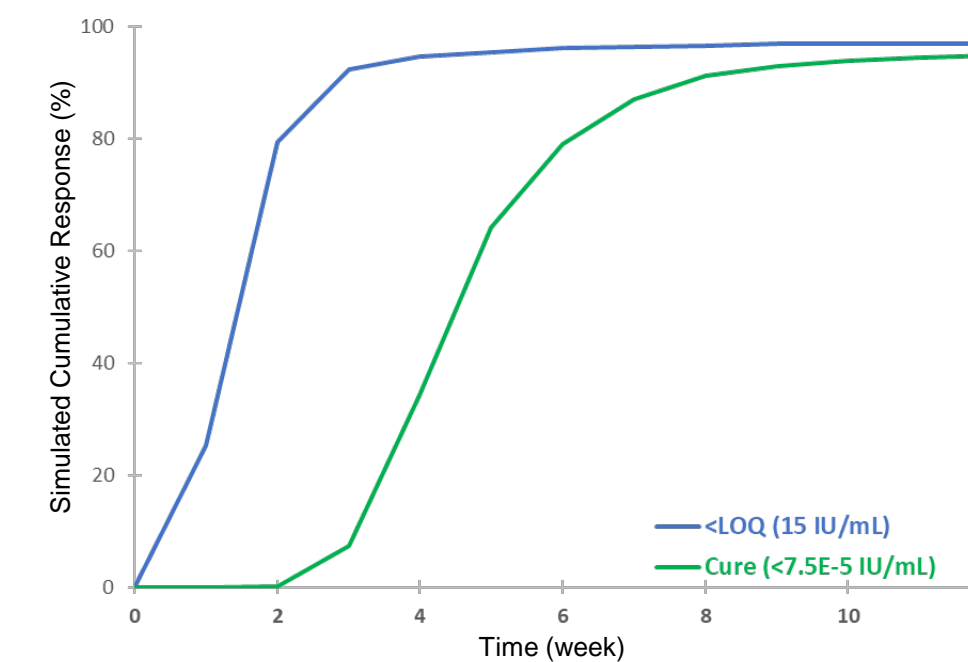
### Summary (mean±SD) Individual Viral Kinetic Parameters by Cohort/Dose

Cohort	N	Log <sub>10</sub> V <sub>0</sub> (IU/mL)	$\delta$ (d <sup>-1</sup> )	T <sub>1/2</sub> cell (d)	$c$ (h <sup>-1</sup> )	T <sub>1/2</sub> HCV (h)	$\tau$ (h)	$\epsilon$
138 mg GT1b NC	6	5.75±0.45	1.22±0.50	15.0±4.33	0.394±0.059	1.79±0.25	4.08±0.12	0.888±0.042
275 mg GT1b NC	6	6.12±0.34	1.22±0.44	16.0±8.24	0.335±0.030	2.09±0.18	3.84±0.19	0.976±0.013
All 550 mg	18	6.16±0.40	0.98±0.23	18.2±5.48	0.331±0.061	2.14±0.43	4.10±0.17	0.992±0.006
Overall	30	6.07±0.44	1.07±5.67	17.1±0.32	0.344±0.053	2.06±0.32	4.04±0.17	

- Viral kinetic parameter estimates were consistent across cohorts, except for  $\epsilon$ , the effectiveness of AT-527 in blocking viral production: the 550 mg dose resulted in the highest  $\epsilon$  which is in agreement with up to a mean of 2.4 log<sub>10</sub> reduction of HCV viral load observed in the first 24 h after dosing in the 550 mg cohorts, regardless of genotypes or cirrhosis status (3).
- The estimated loss rate of infected hepatocytes ( $\delta$ ), about 1 d<sup>-1</sup>, with AT-527 alone was faster than 0.35 day<sup>-1</sup> with sofosbuvir (4). This higher loss rate with AT-527 leads to an estimated infected cell half-life of approximately 17 h, as compared to 48 h with sofosbuvir.
- The estimated clearance rate constant of HCV virions ( $c$ ) and half-life were ~0.35 h<sup>-1</sup> and 2.0 h respectively across cohorts. These results are in agreement with previously published data (5).
- The pharmacologic delay ( $\tau$ ) was estimated to be around 4 h. This is consistent with AT-527 being a nucleotide prodrug that requires multistep activation to its active triphosphate. This delay is in agreement with the plasma T<sub>max</sub> (4-5 h) of AT-273, nucleoside surrogate of the intracellular active triphosphate of AT-527.

### Simulation

- Simulation was performed with subjects receiving a 550 mg QD dosing regimen of AT-527 as a single agent to assess treatment duration required to achieve a cure, defined as plasma HCV RNA < 1 IU/mL (-4.13 log<sub>10</sub> IU/mL). Time to reach plasma HCV RNA < LLQ was also evaluated.

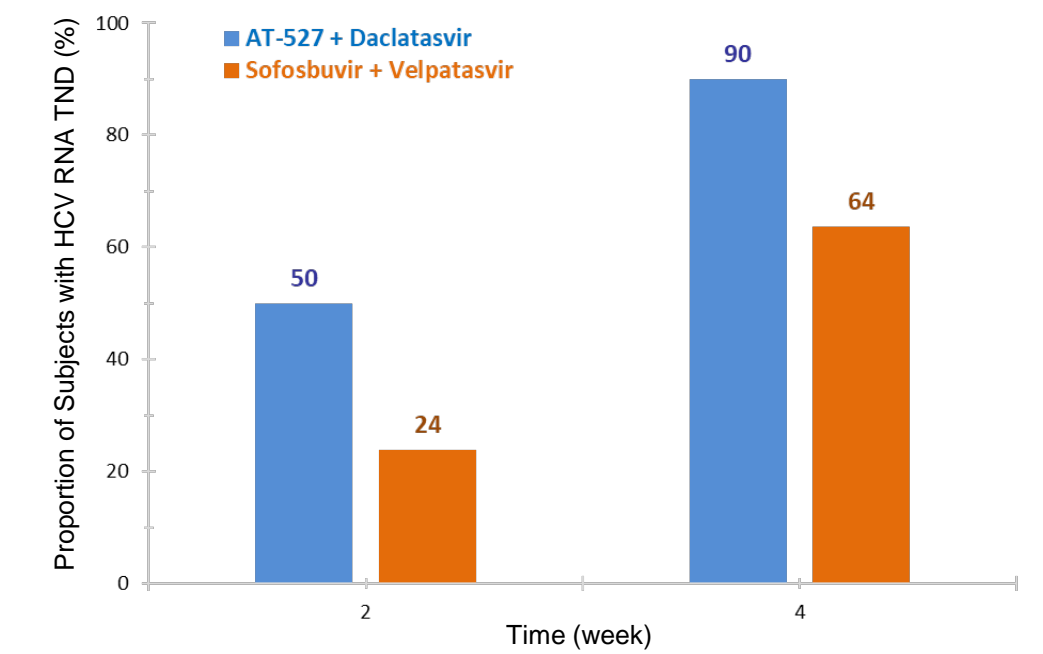


Simulation results showed that with AT-527 as a single agent at 550 mg/day:

- About 80% and 95% of subjects would have plasma HCV RNA < LLQ by wk 2 and 4 respectively.
- Approximately 80% and >90% of subjects would achieve a cure with 6 and 8 wks of treatment, regardless of HCV genotype or cirrhosis status.

### On-Treatment Response from an On-Going Phase II Study of AT-527 plus Daclatasvir

- AT-527 plus daclatasvir was well-tolerated for 8-12 weeks in HCV GT1 treatment-naïve subjects (N=10).
- High proportions of subjects achieved HCV RNA target not detected (TND) at week 2 (50%) and week 4 (90%), with all subjects achieving TND at end of treatment. These on-treatment response rates are much higher than those observed with sofosbuvir plus velpatasvir (6).
- The very rapid early viral kinetics with AT-527 plus daclatasvir allowed for an 8 week treatment duration in all but one subject. SVR results are pending.



## Conclusions

- Viral kinetic analysis based on data generated from 7-day dosing of AT-527 as a single agent suggests that >90% of all patients should be cured with 8 weeks of treatment.
- Interim data from an on-going pilot study with AT-527 plus daclatasvir, the only commercially available standalone NS5A inhibitor, suggest that high cure rates and shorter treatment duration may be achieved with a AT-527 containing regimen, as compared to sofosbuvir-containing regimens.
- A highly effective single-tablet once daily pan-genotypic short treatment regimen with less drug-drug-interaction potential will greatly simplify the care for chronic hepatitis C especially in patients with cirrhosis.

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