

## **SUPPLEMENTARY TABLES**

*Supplementary Table 1. Primer sequences used in this study.*

<i>Name</i>	<i>Sequence (5'-3')</i>	<i>Type</i>
sJFH_8202 <sup>a</sup>	GTG GAG TAT CTC TTG AAA GCA TGG	Outer sense
sJFH_8256	TAT GAT ACC CGA TGC TTC GAC TCA	Inner sense
sJFH_9010	ATT GAG AGG TTA CAC GGG CTT GA	Inner antisense
sJFH_9049	ACA TAC TCT CAC CAC GAA CTG AC	Outer antisense

<sup>a</sup> *Primer numbers refer to the positioning on the genome using absolute numbering of nucleotides of the reference strain H77.*

*Supplementary Table 2. Detection limit of nRT-PCR as determined by serial endpoint dilutions of JFH-1 transcripts.*

<i>RNA copies/ reaction</i>	<i>No. Positive results/ Total PCRs</i>
6	8/8
3	6/8
0.6	2/8