

CORRESPONDENCE

Suppression of hepatitis C virus replication during COVID-19 infection

An outbreak of COVID-19 caused by Delta variant occurred in Southern Taiwan on May 2021. It is believed to have originated from two COVID-19 cases from Peru, and seventeen patients have been infected with Delta variant in the event. Here, we report a case suggesting that SARS-CoV-2 might suppress HCV during COVID-19 infection.

A 70-year-old woman presented with fever and cough for 2 days. She had tested positive for Delta variant of SARS-CoV-2 through nasopharyngeal swab real-time reverse transcriptase-polymerase chain reaction (RT-PCR) and virus culture; these tests are carried out at the Taiwan CDC laboratory or in a designated laboratory at hospitals. She was isolated in a negative pressure ward at a government hospital in Pingtung, Taiwan. The initial biochemical analysis results were abnormal for aspartate aminotransferase (AST) (47 IU/L), and the hepatitis C antibody (Anti-HCV) was positive. This was the first time she was diagnosed with HCV infection. The concurrent HCV RNA titer was low (140 IU/mL) (Table 1). Remdesivir was administered for COVID-19 infection and she was discharged after 2 weeks (Figure 1). HCV RNA titer was rechecked 8 weeks after COVID-19 infection and HCV RNA titer elevated more than 10,000 times (1,675,038 IU/mL), but the serum AST and alanine aminotransferase

(ALT) was normal. HCV Genotype 1b was detected and she received direct antiviral agents (DAA).

In late 2019, COVID-19 emerged in Wuhan, China, and resulted in a formidable outbreak that expanded globally and caused huge health and economic burden. COVID-19 causes damages to many extrapulmonary organs. The first case of COVID-19 in Taiwan was diagnosed on January 2020 and the mortality rate was 1.3% at that time.¹ Taiwan had few domestic cases until small outbreaks occurred in May 2021.² The dominant variants were Alpha and Beta, and Delta variant cluster emerged last June 2021. Delta variant emerged recently in India and spread gradually to other countries. It has now become the dominant cause of COVID-19 in the world.

SARS-CoV-2 might cause hepatitis B virus (HBV) reactivation³; however, the interaction between HCV and SARS-CoV-2 has rarely been described. One case report described no significant change in HCV RNA titer after COVID-19 infection in a nonhospitalized patient.⁴ Here, we report a case that HCV RNA might be significantly elevated after COVID-19 infection in hospitalized patients, and HCV viremia flare up might occur after COVID-19 infection. In the previous literature, the mean change of viral load was less than 1 log₁₀ per year and rarely exceeded 2 log₁₀.⁵ This case changed more than 4 log₁₀ within

TABLE 1 The characteristics of the COVID-19 patient before and after the COVID-19 infection

	Pretreatment	Posttreatment ^a
AST (IU/L, mean [SD])	47	29
ALT (IU/L, mean [SD])	30	14
Bilirubin (mg/dL)	1.6	1.5
Platelet count (×10 ³ U/L, mean [SD])	39	70
Albumin (g/dL)	3.1	3.4
Creatinine (mg/dL)	1.0	1.0
HBsAg	Negative	Negative
HCV RNA (IU/mL)	140	1,675,038

Note: COVID-19: coronavirus disease 2019; AST: aspartate aminotransferase.

Abbreviations: ALT, alanine aminotransferase; HBsAg, hepatitis B surface antigen; HCV, hepatitis C virus.

^aEight weeks after the admission.

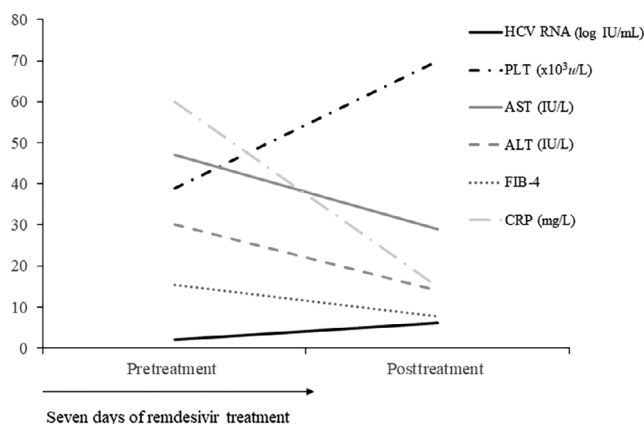


FIGURE 1 The pretreatment and posttreatment biochemical data of the patient. ALT, alanine aminotransferase; AST, aspartate aminotransferase; CRP, C-reactive protein; FIB-4, fibrosis-4 index; HCV, hepatitis C virus; PLT, platelet count

This is an open access article under the terms of the Creative Commons Attribution-NonCommercial-NoDerivs License, which permits use and distribution in any medium, provided the original work is properly cited, the use is non-commercial and no modifications or adaptations are made.

© 2022 The Author. *The Kaohsiung Journal of Medical Sciences* published by John Wiley & Sons Australia on behalf of Kaohsiung Medical University.


2 months, and it was unlikely to be a natural fluctuation. The patient received 7 days of remdesivir treatment, and the HCV RNA was checked after 2 days of remdesivir treatment. There was no evidence that remdesivir would suppress HCV in the current literature. We suppose that SARS-CoV-2 might suppressed HCV during COVID-19 infection, but further study is needed. The limitation to our case was that the baselines status of HCV RNA before COVID-19 infection was unknown.

ACKNOWLEDGMENTS

This study was supported by the Ministry of Health and Welfare, Pingtung Hospital and Kaohsiung Medical University Hospital. The authors thank the secretaries of the Hepatobiliary Division in Kaohsiung Medical University Hospital.

CONFLICT OF INTEREST

All authors declare no conflict of interest.

Tyng-Yuan Jang^{1,2} 

¹*Hepatobiliary Division, Department of Internal Medicine, Kaohsiung Medical University Hospital, Kaohsiung Medical University, Kaohsiung, Taiwan*

²*Department of Internal Medicine, Pingtung Hospital, Ministry of Health and Welfare, Pingtung, Taiwan*

Correspondence

Tyng-Yuan Jang, Hepatobiliary Division, Department of Internal Medicine, Kaohsiung Medical University Hospital, 100 Tzyou Road, Kaohsiung City 807, Taiwan.
Email: sls0902000@gmail.com

ORCID

Tyng-Yuan Jang  <https://orcid.org/0000-0003-2961-130X>

REFERENCES

1. Lin CY, Wang WH, Urbina AN, Tseng SP, Lu PL, Chen YH, et al. Importation of SARS-CoV-2 infection leads to major COVID-19 epidemic in Taiwan. *Int J Infect Dis.* 2020;97:240–4.
2. Wang CJ, Ng CY, Brook RH. Response to COVID-19 in Taiwan: big data analytics, new technology, and proactive testing. *JAMA.* 2020;323(14):1341–2.
3. Liu J, Wang T, Cai Q, Sun L, Huang D, Zhou G, et al. Longitudinal changes of liver function and hepatitis B reactivation in COVID-19 patients with pre-existing chronic hepatitis B virus infection. *Hepatol Res.* 2020;50(11):1211–21.
4. Villar LM, de Paula VS, Pinto LCM, Marques BCL, da Costa VD, da Silva LL, et al. Clinical and laboratory characteristics of hepatitis C and COVID-19 coinfection: prolonged RNA shedding in nonhospitalized case. *Clin Case Rep.* 2021;9(8):e03877.
5. Fanning L, Kenny-Walsh E, Levis J, Choudhury KR, Cannon B, Sheehan M, et al. Natural fluctuations of hepatitis C viral load in a homogeneous patient population: a prospective study. *Hepatology.* 2000;31(1):225–9.