



Since January 2020 Elsevier has created a COVID-19 resource centre with free information in English and Mandarin on the novel coronavirus COVID-19. The COVID-19 resource centre is hosted on Elsevier Connect, the company's public news and information website.

Elsevier hereby grants permission to make all its COVID-19-related research that is available on the COVID-19 resource centre - including this research content - immediately available in PubMed Central and other publicly funded repositories, such as the WHO COVID database with rights for unrestricted research re-use and analyses in any form or by any means with acknowledgement of the original source. These permissions are granted for free by Elsevier for as long as the COVID-19 resource centre remains active.

Journal Pre-proof

Prolonged Virus Shedding Even after Seroconversion in a Patient with COVID-19

Wang-Da Liu , Sui-Yuan Chang , Jann-Tay Wang , Ming-Jui Tsai , Chien-Ching Hung , Chia-Lin Hsu , Shan-Chwen Chang

PII: S0163-4453(20)30190-0
DOI: <https://doi.org/10.1016/j.jinf.2020.03.063>
Reference: YJINF 4537



To appear in: *Journal of Infection*

Accepted date: 31 March 2020

Please cite this article as: Wang-Da Liu , Sui-Yuan Chang , Jann-Tay Wang , Ming-Jui Tsai , Chien-Ching Hung , Chia-Lin Hsu , Shan-Chwen Chang , Prolonged Virus Shedding Even after Seroconversion in a Patient with COVID-19, *Journal of Infection* (2020), doi: <https://doi.org/10.1016/j.jinf.2020.03.063>

This is a PDF file of an article that has undergone enhancements after acceptance, such as the addition of a cover page and metadata, and formatting for readability, but it is not yet the definitive version of record. This version will undergo additional copyediting, typesetting and review before it is published in its final form, but we are providing this version to give early visibility of the article. Please note that, during the production process, errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

© 2020 The British Infection Association. Published by Elsevier Ltd. All rights reserved.

Highlights

- Prolonged virus shedding could be found among COVID-19 patients after clinical symptoms resolved and specific antibody emerged.
- Viral detection from throat gargling sample could be an alternative diagnostic method for mild COVID-19 patients with scarce sputum.
- Sputum remained the most sensitive specimen for viral detection after clinical symptoms resolved.

Journal Pre-proof

Dear editors,

Previous reports revealed that the emergence of the novel coronavirus (SARS-CoV-2) infection (COVID-19) had raised global concern.¹ Several studies discussing the clinical pictures of COVID-19 and specific antibody responding to SARS-CoV-2 have been published.²⁻⁴ However, the time sequences of clinical manifestations, virus shedding kinetics, contagiousness, and specific antibody reaction, which are essential for understanding pathophysiology and infection control strategy, have less been discussed. Here we present a COVID-19 patient with prolonged viral shedding and detailed time sequence of these parameters mentioned above.

A 50-year-old woman, who suffered from acute-onset fever up to 38.6°C four hours ago, visited the emergency department of a medical center in Taipei City, Taiwan. She denied of cough or other subjective discomforts, and disclosed that she lived in Wuhan, China and just traveled to Taiwan two days prior to hospital visit. Given the patient's travel history, she was transferred to a negative-pressure isolation room for suspecting COVID-19.

Chest radiography was normal. Hemogram revealed only mild thrombocytopenia (143 K/ μ L), while other blood tests were within normal limits. Sputum and throat swab specimens yielded positive results for SARS-CoV-2 by

real-time reverse transcription-polymerase chain reaction (RT-PCR). Fever subsided rapidly after admission. However, elevated temperature up to 38.2°C was noted four days later. She did not have obvious respiratory symptoms and remained good spirit. All blood tests, except mildly elevated C-reactive protein (CRP) (1.47 mg/dL) and mild thrombocytopenia, were normal. A once-daily low-grade fever persisted until ninth day after symptom onset. The CRP level reached to the peak on the tenth day after symptom onset and then normalized gradually.

Clinical specimens for SARS-CoV-2 testing were obtained in accordance with guidelines from the Centers for Disease Control and Prevention.⁵ During hospitalization, sputum and throat swab specimens were collected for SARS-CoV-2 RT-PCR and virus culture every other day. Throat wash by gargling using 10 mL normal saline were collected when sputum specimens were not available.⁶ Stool was collected for RT-PCR on the third, 14th and 20th day after symptom onset, and plasma was also sent within the first four days of hospitalization.

Plasmid DNA containing the SARS-CoV-2 target sequences, including the envelope (E), nucleocapsid (N), and RNA-dependent RNA polymerase (RdRp) genes, was used to construct the standard curve to estimate the SARS-CoV-2 viral load by real-time RT-PCR. In addition, SARS-CoV-2 isolation was performed via Vero E6 and LLC-MK2 cell cultures. The full-length viral sequence was determined using

SARS-CoV-2 amplified from the sputum specimen collected on the ninth day after symptom onset, and was submitted to the GISAID (accession number is EPI_ISL_408489). Antibody response to SARS-CoV-2 viral N proteins was determined by western blotting using infected cell lysates.

The sequential changes of SARS-CoV-2 viral load in throat swab, sputum, and gargling water are presented in Figure. A drop of three logs of viral loads was observed among all specimens within one week after admission. However, SARS-CoV-2 persisted to be detectable till 63th day after symptom onset. Among respiratory specimens from different sites, specimen from sputum showed superior sensitivity to samples from throat swab and gargling wash. Initially, specimen from gargling wash showed considerable results compared with specimens from sputum or throat swab. Nevertheless, RT-PCR examinations from sputum still outweighed samples from gargling and throat swab after fever subsided. Among three target genes of the RT-PCR examinations, E gene of SARS-CoV-2 showed superior sensitivity to N and RdRp gene after clinical symptoms resolved. The stool specimen collected on the third day after symptom onset yielded positive results, but turned negative in the following specimens. The presence of SARS-CoV-2 could not be detected in the plasma samples while the SARS-CoV-2 titers in the respiratory specimen remained high.

SARS-CoV-2 could be isolated from cell cultures in throat swab collected upon admission, and all sputum specimens collected within 18 days after symptoms onset. RT-PCR continued to detect virus till the 63th day after symptom onset regardless virus could only be isolated from respiratory specimens collected within the first 18 days. Antibody to SARS-CoV-2 was firstly identified on the tenth day after symptom onset. In the meanwhile, no more fever above 37.5°C was noted, and CRP began to decline. Recovery of platelet count took place two days earlier than the emergence of SARS-CoV-2 antibody.

This case demonstrated that the virus shedding might continue even after clinical resolution and seroconversion. In addition, although SARS-CoV-2 virus could not be isolated after the 18th day of symptom onset, the positive RT-PCR results continued for more than 60 days. Because of the long interval between these two time points, it might be reasonable to infer that a small amount of viable virus, yet could not be detected by virus culture, remained present after the 18th day of disease course and last for more days. This implies the contagious period of COVID-19 might last more than one week after “clinical recovery”. Many of COVID-19 patients in Taiwan also had similar findings (unpublished data). Such prolonged virus shedding was also observed among asymptomatic pediatric patients in fecal specimen.⁷ However, this needs more studies to clarify since it would be a major issue in realizing and

controlling the COVID-19 epidemics.

Serial RT-PCR results in our case highlighted the importance of the clinical specimens sampled from lower respiratory tract for detecting SARS-CoV-2 virus⁸, which is different from the report from Zou L, et al.⁹ Additionally, our case demonstrated that throat wash by gargling could be an alternative method for COVID-19 diagnosis, since specimen from lower respiratory tract was barely available at the early stage of infection. Nevertheless, when the patient became afebrile, sputum still outweighed specimens from throat swab or gargling wash.

Our case highlighted the prolonged virus shedding course of COVID-19 even after the clinical symptoms resolved and seroconversion developed. It implies the possibility of a prolonged contagious period and should be investigated further to better control the epidemics. Our study also demonstrated viral detection from throat gargling sample could be an alternative diagnostic method for patients without sputum.

Declaration of Competing Interest

None to declare.

Acknowledgments

None to declare.

Journal Pre-proof

References

1. Tang JW, Tambyah PA, Hui DSC. Emergence of a novel coronavirus causing respiratory illness from Wuhan, China. *J Infect* **2020**;80(3):350-371.
2. Xiao DAT, Gao DC, Zhang DS. Profile of Specific Antibodies to SARS-CoV-2: The First Report. *J Infect* **2020** Mar 21. pii: S0163-4453(20)30138-9. doi: 10.1016/j.jinf.2020.03.012. [Epub ahead of print]
3. Guan WJ, Ni ZY, Hu Y, Liang WH, Ou CQ, He JX, et al. Clinical Characteristics of Coronavirus Disease 2019 in China. *N Engl J Med* **2020** Feb 28. doi: 10.1056/NEJMoa2002032. [Epub ahead of print]
4. Yang W, Cao Q, Qin L, Wang X, Cheng Z, Pan A, et al. Clinical characteristics and imaging manifestations of the 2019 novel coronavirus disease (COVID-19): A multi-center study in Wenzhou city, Zhejiang, China. *J Infect* **2020** Feb 26. pii: S0163-4453(20)30099-2. doi: 10.1016/j.jinf.2020.02.016. [Epub ahead of print]
5. Centers for Disease Control and Prevention. Interim Guidelines for Collecting, Handling, and Testing Clinical Specimens from Persons Under Investigation (PUIs) for Coronavirus Disease 2019 (COVID-19). Available at: <https://www.cdc.gov/coronavirus/2019-nCoV/lab/guidelines-clinical-specimens.html> Accessed 7 Mar 2020.

6. Wang WK, Chen SY, Liu IJ, Chen YC, Chen HL, Yang CF, et al. Detection of SARS-associated coronavirus in throat wash and saliva in early diagnosis. *Emerg Infect Dis* **2004**;10:1213-9.
7. Xu Y, Li X, Zhu B, Liang H, Fang C, Gong Y, et al. Characteristics of pediatric SARS-CoV-2 infection and potential evidence for persistent fecal viral shedding. *Nat Med* 2020. doi:10.1038/s41591-020-0817-4. [Epub ahead of print]
8. Wang W, Xu Y, Gao R, Lu R, Han K, Wu G, et al. Detection of SARS-CoV-2 in Different Types of Clinical Specimens. *JAMA*. 2020. doi: 10.1001/jama.2020.3786. [Epub ahead of print]
9. Zou L, Ruan F, Huang M, Liang L, Huang H, Hong Z, et al. SARS-CoV-2 Viral Load in Upper Respiratory Specimens of Infected Patients. *N Engl J Med* **2020** Feb 19. doi: 10.1056/NEJMc2001737. [Epub ahead of print]

Figure Legends

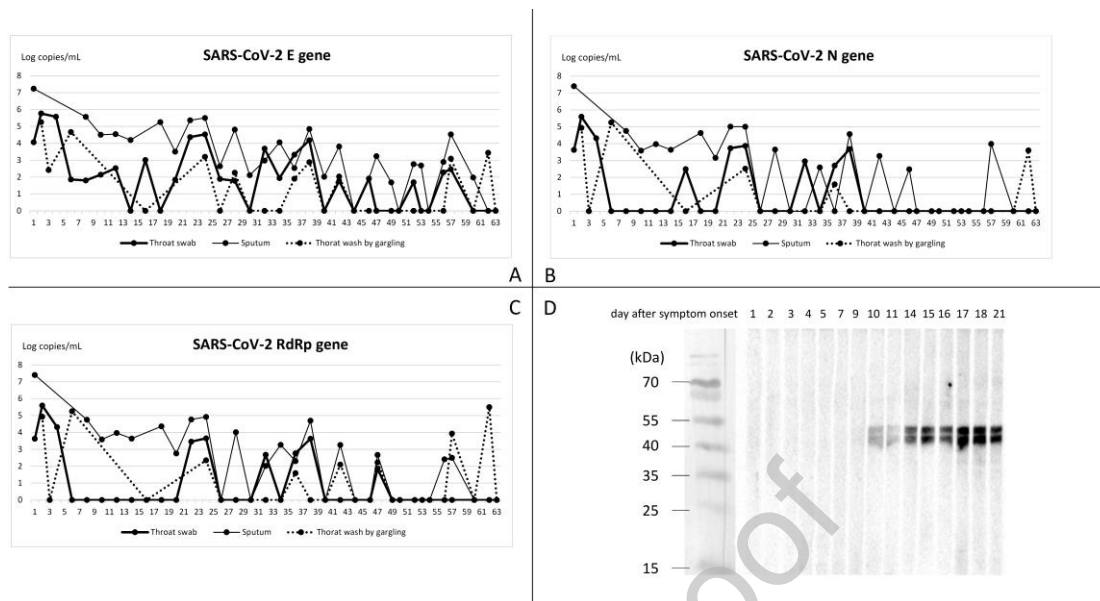


Figure 1. Profile of viral load and antibody response to SARS-CoV-2 in the case

patient. (A)(B)(C) Sequential changes of SARS-CoV-2 viral load in the throat swab, sputum, and throat wash by gargling. The SARS-CoV-2 viral load was determined using the protocol provided by the WHO (<https://virologie-ccm.charite.de/en>).

Plasmids containing partial E, N, and RdRp fragments were used respectively as the standards to calculate the amount of SARS-CoV-2 viral load in the specimens. (D)

Antibody responses of the present case. Antibody responses of the present case to viral proteins extracted from SARS-CoV-2 infected Vero-E6 cells were determined by the western blot.