Acknowledgments

We thank Lance Turtle for his insightful comments on our manuscript.

This study was supported by the Ministry of Science and Technology of China, the National Mega Project on Major Infectious Disease Prevention (no. 2017ZX10103005), the National Key Research and Development Program of China (no. 2018YFE0204500), the National Science Fund for Distinguished Young Scholars (no. 81525023), the National Science and Technology Major Project of China (nos. 2018ZX10713001-007, 2018ZX10201001-010, and 2017ZX10103009-005), the Special Fund for COVID-19 Research of Wuhan University, and the Fund for COVID-19 Research of Taikang Insurance Group Co., Ltd, and Beijing Taikang Yicai Foundation. No funders had any role in the study design; the collection, analysis, and interpretation of data; the writing of the article; or the decision to submit it for publication. The researchers confirm their independence from funders and sponsors.

H.Y. has received research funding from Sanofi Pasteur, GlaxoSmithKline, Yichang HEC Changjiang Pharmaceutical Company, and Shanghai Roche Pharmaceutical Company. None of that research funding is related to COVID-19. All other authors report no competing interests.

H.Y., Y. L., and Z.Y. supervised the study. Z.Y., J. N., H.H., Y.W., and L. Li recruited the patients; collected specimens; and collected demographic, clinical, and laboratory data. J.C., Q.W., and L. Liang plotted the figures; J.C., Q.W., W.L., Q.Z., and Y.L. analyzed the data; H.Y., Y.L., and Z.Y. interpreted the results; J.C. and Q.W. wrote the initial drafts of the manuscript. All authors read and approved the final report.

About the Author

Dr. Yao is a physician at the Wuhan Pulmonary Hospital, Wuhan, China. His research focuses on respiratory infections and critical care medicine.

References

- World Health Organization. Global tuberculosis report 2019 [cited 2020 Jun 11]. https://www.who.int/tb/publications/ global_report
- Houben RM, Dodd PJ. The global burden of latent tuberculosis infection: a re-estimation using mathematical modelling. PLoS Med. 2016;13:e1002152. https://doi.org/10.1371/journal. pmed.1002152
- Zhao Y, Xu S, Wang L, Chin DP, Wang S, Jiang G, et al. National survey of drug-resistant tuberculosis in China. N Engl J Med. 2012;366:2161–70. https://doi.org/10.1056/NEJMoa1108789
- Low JG, Lee CC, Leo YS, Low JG, Lee CC, Leo YS. Severe acute respiratory syndrome and pulmonary tuberculosis. Clin Infect Dis. 2004;38:e123–5. https://doi.org/10.1086/421396
- Liu W, Fontanet A, Zhang PH, Zhan L, Xin ZT, Tang F, et al. Pulmonary tuberculosis and SARS, China. Emerg Infect Dis. 2006;12:707–9. https://doi.org/10.3201/eid1204.050264

- Alfaraj SH, Al-Tawfiq JA, Altuwaijri TA, Memish ZA. Middle East respiratory syndrome coronavirus and pulmonary tuberculosis coinfection: implications for infection control. Intervirology. 2017;60:53–5. https://doi.org/ 10.1159/000477908
- World Health Organization. World Health Organization (WHO) information note: tuberculosis and COVID-19 [cited 2020 Jun 11]. https://www.who.int/docs/default-source/ documents/tuberculosis/infonote-tb-covid-19.pdf
- Guan WJ, Liang WH, Zhao Y, Liang HR, Chen ZS, Li YM, et al.; China Medical Treatment Expert Group for COVID-19. Comorbidity and its impact on 1590 patients with COVID-19 in China: a nationwide analysis. Eur Respir J. 2020;55:2000547. https://doi.org/10.1183/13993003.00547-2020

Address for correspondence: Hongjie Yu, School of Public Health, Fudan University, Key Laboratory of Public Health Safety, Ministry of Education, Shanghai, China; email: yhj@fudan.edu. cn; and Yingle Liu, State Key Laboratory of Virology, Modern Virology Research Center, College of Life Sciences, Wuhan University, Wuhan, China; email: mvlwu@whu.edu.cn

Detection of SARS-CoV-2 in Hemodialysis Effluent of Patient with COVID-19 Pneumonia, Japan

Ayako Okuhama, Masahiro Ishikane, Daisuke Katagiri, Kohei Kanda, Takato Nakamoto, Noriko Kinoshita, Naoto Nunose, Takashi Fukaya, Isao Kondo, Harutaka Katano, Tadaki Suzuki, Norio Ohmagari, Fumihiko Hinoshita

Author affiliations: National Center for Global Health and Medicine, Tokyo, Japan (A. Okuhama, M. Ishikane, D. Katagiri, K. Kanda, T. Nakamoto, N. Kinoshita, N. Nunose, T. Fukaya, I. Kondo, N. Ohmagari, F. Hinoshita); National Institute of Infectious Diseases, Tokyo (H. Katano, T. Suzuki)

DOI: https://doi.org/10.3201/eid2611.201956

We report detection of severe acute respiratory syndrome coronavirus 2 RNA in hemodialysis effluent from a patient in Japan with coronavirus disease and prolonged inflammation. Healthcare workers should observe strict standard and contact precautions and use appropriate personal protective equipment when handling hemodialysis circuitry from patients with diagnosed coronavirus disease. Since December 2019, coronavirus disease (CO-VID-19), caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), has been a major health threat worldwide (1). Reports have been published on COVID-19 among patients receiving hemodialysis (2), but none have evaluated whether HD effluent is infectious. In addition, handling of hemodialysis circuitry is not mentioned in US Centers for Disease Control and Prevention (CDC) guidelines for COVID-19 infection control and prevention in dialysis facilities (3). We report detection of SARS-CoV-2 RNA in hemodialysis effluent from a patient with COVID-19 pneumonia and prolonged inflammation.

The patient, a 79-year-old man with end-stage renal disease (ESRD) due to IgA nephritis, had been receiving maintenance hemodialysis 3 times per week for 12 years. Six days before admission, he started having a fever and cough. Four days later, he had a nasal swab test for SARS-CoV-2 RNA. Quantitative reverse transcription PCR (qRT-PCR) (4) of the patient's specimen was positive, and he was admitted to the hospital. At admission, his body temperature was 37.7°C and oxygen saturation was 98% on room air. Multiple bilateral patchy ground glass opacities (GGO) were observed on the patient's chest computed tomography (CT) scan (Figure, panel A). Blood test results showed C-reactive protein (CRP) of 8.8 mg/dL and leukocyte count of 4,470 cells/ μ L. Although we started him on hydroxychloroquine (200 mg 2 ×/d) and azithromycin (500 mg, $1 \times /d$), he had a fever (>38.0°C) on day 2 of his hospitalization. A follow-up chest CT on hospitalization day 5 showed worsening COVID-19 pneumonia and expanding GGO areas (Figure, panel B).

During the patient's hospitalization, we administered hemodialysis by using a polysulfone membrane dialyzer in a private depressurized room with dedicated machines. We tested hemodialysis effluent for SARS-CoV-2 on day 2. PCR results showed SARS-CoV-2 RNA of 157.9 copies/µL with cycle threshold (C) values of 38.3 at 1 hour after starting hemodialysis but were negative on effluent collected at 2 hours. Because the patient's fever persisted and CRP levels remained high, on hospitalization days 9, 11, and 15 we performed direct hemoperfusion by using a $\beta 2$ microglobulin adsorbent column (Lixelle-DHP) to absorb cytokine. On hospitalization day 10, the patient became afebrile and CRP began decreasing until it reached 5.9 mg/dL on hospitalization day 15. On hospitalization day 16, chest CT showed markedly improved pneumonia (Figure, panel C), and the patient was discharged (Table).

Our case highlights 3 things. First, inflammation and clinical symptoms of COVID-19 can persist in patients on hemodialysis. COVID-19 is thought to progress in a 2-stage manner: viral replication and hyperinflammation (1). Hyperinflammation starts 7-10 days after symptom onset and involves extensive lung areas. This patient's fever persisted for >13 days, with pneumonia and CRP worse at 11 days after fever onset. Hyperinflammation appeared to progress slower and be maintained longer than in patients who are not receiving hemodialysis, which might be related to immune system dysfunction in patients with ESRD (5). Second, although SARS-CoV-2 RNA has been detected in various clinical specimens (6,7), our case demonstrates it also can be detected in hemodialysis effluent, even though we did not detect SARS-CoV-2 RNA in blood, as

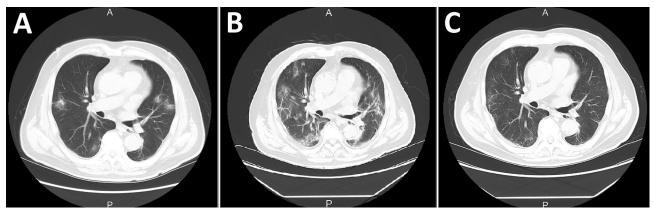


Figure. Chest computed tomography (CT) scan of a patient on hemodialysis diagnosed with positive reverse transcription PCR for severe acute respiratory syndrome coronavirus 2 in hemodialysis effluent, Japan. A) Chest CT at day 1 of hospitalization showing bilateral patchy ground glass opacities (GGO). B) Chest CT from day 5 of hospitalization showing worsening coronavirus disease 2019 (COVID-19) pneumonia with GGO expansion. C) Chest CT on hospitalization day 16 showing improvement of COVID-19 pneumonia; the patient was discharged on this day. A, anterior; P, posterior.

		, - I						Specimens tested for SARS-CoV-2 by qRT-PCR‡			
Day after									Effluent	, time	
symptom	Hospitalization,	Temperature,		Med	lication†	CRP,		Nasal		collected	
onset	d	°C	Event	AZM	Hydroxy	mg/dL	Dialysis	swab	Blood	1 h	2 h
1		37.3									
2		37.2									
3		37.3									
4		37.3	Clinic					18.8 (NA‡)			
5		37.7									
6		39.0									
7	1	38.8	Chest CT	Ν	N	8.8	-				
8	2	38.4		Y	Y	9.0	Y	29.6	ND§	38.3	ND
								(1,080.6)§		(157.91)	
9	3	38.7		Y	Y	-	-	_			
10	4	38.7		Y	Y	14.0	-	_			
11	5	37.4	Chest CT	Ν	Y	15.0	Y	_			
12	6	37.0		N	Y	_	_	_			
13	7	37.2		Ν	Y	-	-	_			
14	8	37.0		N	Y	_	_	_			
15	9	36.9		Ν	Y	14.4	Lixelle- DHP	-			
16	10	37.0		Ν	Ν			34.3 (NA‡)			
17	10	36.9		N	N	_	_ Lixelle-	ND			
17		50.5		IN	IN IN	_	DHP				
18	12	36.9		N	N	13.7	_	ND			
19	13	36.8		Ν	N	_	-	_			
20	14	36.6		N	N	_	_	_			
21	15	36.7		Ν	Ν	5.9	Lixelle- DHP	-			
22	16	36.7	Chest CT, discharge	Ν	Ν	-	_	-			

 Table. Clinical course and quantitative reverse transcription PCR results for severe acute respiratory syndrome coronavirus 2 RNA in patient receiving hemodialysis, Japan*

*AZM, azithromycin; CRP, C-reactive protein; CT, computed tomography; Hydroxy, hydroxychloroquine; Lixelle-DHP, direct hemoperfusion using a β2 microglobulin adsorbent column; NA, not available; ND, not detected; qRT-PCR, quantitative reverse transcription-PCR; –, not done. †We prescribed azithromycin, 500 mg 2 times/d from day 1 to 3 because it was 1 of the potentially effective treatment regimens at the time. We also prescribed hydroxychloroquine 200 mg 2 times/d and initially planned to use it for 10 d in total, but the patient's liver function tests (LFTs) became elevated during the course. We suspected side effects of hydroxychloroquine and stopped it on day 9. His LFTs returned to normal afterwards. ‡Results for SARS-CoV-2 shown as cycle threshold values (Viral load, copies/μL). Viral loads were not available because PCR was performed at an outside commercial laboratory where they did not report these results. The same PCR method was used (4) at both National Institute of Infectious Diseases (NIID), Japan, and the outside laboratory. HD effluent was collected at 1 hr and 2 hr into hemodialysis. §PCR test was performed at NIID, Japan where they report viral loads.

noted in a previous case (6). We hypothesized that only a small amount of fragmented RNA might pass through the dialysis membrane at the start of hemodialysis, but no marked fragments remain in the blood as a session progresses. Third, our case suggests Lixelle-DHP can have therapeutic effects for patients on hemodialysis. Although we did not measure the patient's predialysis and postdialysis cytokine levels, use of a blood purification technique might alleviate the effects of cytokine in COVID-19 pathophysiology due to its proven effect in reducing plasma cytokine levels in general (8).

Our report has several limitations. First, we did not confirm the duplicability of PCR results of hemodialysis effluent. We performed PCR only once and did not reevaluate the same specimen, even though the C_t was high. Second, the infectiousness of hemodialysis effluent is unclear. Its viability should be quantified by endpoint titration on authorized cell lines, as previously reported (9). Third, this is a single case report. Despite these limitations, we cannot underestimate the infectiousness of hemodialysis effluent. We performed dialysis in a private room with dedicated machines. We also conducted strict standard and contact precautions when handling HD circuitry, following CDC recommendations for preventing transmission of hepatitis B virus infection among patients on HD (*10*).

In conclusion, we report positive qRT-PCR results for SARS-CoV-2 RNA from hemodialysis effluent in a patient receiving renal dialysis. The clinical course of our patient was characteristic of the persistent inflammation of COVID-19 and shows the potential effectiveness of Lixelle-DHP as a treatment in patients on hemodialysis. Our case indicates that strict standard and contact precautions are essential when handling hemodialysis circuitry of patients with COVID-19. As more patients on hemodialysis contract SARS-CoV-2,

RESEARCH LETTERS

we expect further studies on infection control and prevention in dialysis facilities and on the effectiveness of Lixelle-DHP in treating patients with COVID-19.

Acknowledgments

We thank all the clinical staff at our hospital for their dedication to patient care and the patient, who provided written informed consent to have his clinical details presented in this report.

About the Author

Dr. Okuhama is a clinical senior resident at the Disease Control and Prevention Center, National Center for Global Health and Medicine, Tokyo, Japan. Her research interests include emerging infectious diseases and public health.

References

- Huang C, Wang Y, Li X, Ren L, Zhao J, Hu Y, et al. Clinical features of patients infected with 2019 novel coronavirus in Wuhan, China. Lancet. 2020;395:497–506. https://doi.org/ 10.1016/S0140-6736(20)30183-5
- Wang R, Liao C, He H, Hu C, Wei Z, Hong Z, et al. COVID-19 in hemodialysis patients: A report of 5 cases. Am J Kidney Dis. 2020;76:141–3. https://doi.org/10.1053/ j.ajkd.2020.03.009
- US Centers for Disease Control and Prevention. Interim guidance for infection prevention and control recommendations for patients with suspected or confirmed COVID-19 in outpatient hemodialysis facilities 2020 Apr 21 [cited 2020 Apr 22]. https://www.cdc.gov/ coronavirus/2019-ncov/healthcare-facilities/dialysis.html
- Shirato K, Nao N, Katano H, Takayama I, Saito S, Kato F, et al. Development of genetic diagnostic methods for novel coronavirus 2019 (nCoV-2019) in Japan. Jpn J Infect Dis. 2020;73:304–7. PubMed https://doi.org/10.7883/ yoken.JJID.2020.061
- Lisowska KA, Dębska-Ślizień A, Jasiulewicz A, Heleniak Z, Bryl E, Witkowski JM. Hemodialysis affects phenotype and proliferation of CD4-positive T lymphocytes. J Clin Immunol. 2012;32:189–200. https://doi.org/10.1007/s10875-011-9603-x
- Katagiri D, Ishikane M, Ogawa T, Kinoshita N, Katano H, Suzuki T, et al. Continuous renal replacement therapy for a patient with severe COVID-19. Blood Purif. 2020;11:1–3. https://doi.org/10.1159/000508062
- 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;323:1843–4. https://doi.org/10.1001/jama.2020.3786
- Tsuchida K, Yoshimura R, Nakatani T, Takemoto Y. Blood purification for critical illness: cytokines adsorption therapy. Ther Apher Dial. 2006;10:25–31. https://doi.org/10.1111/ j.1744-9987.2006.00342.x
- van Doremalen N, Bushmaker T, Morris DH, Holbrook MG, Gamble A, Williamson BN, et al. Aerosol and surface stability of SARS-CoV-2 as compared with SARS-CoV-1. N Engl J Med. 2020;382:1564–7. https://doi.org/10.1056/ NEJMc2004973
- US Centers for Disease Control and Prevention. Recommendations for preventing transmission of infections among chronic hemodialysis patients. MMWR Recomm Rep. 2001 Apr 27;50(RR-5):1–43.

Address for correspondence: Masahiro Ishikane, Disease Control and Prevention Center, National Center for Global Health and Medicine 1-21-1 Toyama, Shinjuku-ku, Tokyo 162-8655, Japan; email: ishikanemasahiro@gmail.com

Seroprevalence of SARS-CoV-2–Specific Antibodies, Faroe Islands

Maria Skaalum Petersen, Marin Strøm, Debes Hammershaimb Christiansen, Jógvan Páll Fjallsbak, Eina Hansen Eliasen, Malan Johansen, Anna Sofía Veyhe, Marnar Fríðheim Kristiansen, Shahin Gaini, Lars Fodgaard Møller, Bjarni á Steig, Pál Weihe

Author affiliations: University of the Faroe Islands, Tórshavn,
Faroe Islands (M.S. Petersen, M. Strøm, E.H. Eliasen,
M. Johansen, A.S. Veyhe, M.F. Kristiansen, S. Gaini, P. Weihe);
The Faroese Hospital System, Tórshavn (M.S. Petersen,
E.H. Eliasen, M. Johansen, A.S. Veyhe, P. Weihe); Faroese Food and Veterinary Authority, Tórshavn, (D.H. Christiansen,
J.P. Fjallsbak); National Hospital of the Faroe Islands, Tórshavn,
(M.F. Kristiansen, S. Gaini, B. Steig); COVID-19 Task Force,
Ministry of Health, Tórshavn (M.F. Kristiansen, B. Steig); Odense
University Hospital, Odense, Denmark (S. Gaini); University of
Southern Denmark, Odense (S. Gaini); Chief Medical Officer
Office, Tórshavn (L.F. Møller)

DOI: https://doi.org/10.3201/eid2611.202736

We conducted a nationwide study of the prevalence of severe acute respiratory syndrome coronavirus 2 infection in the Faroe Islands. Of 1,075 randomly selected participants, 6 (0.6%) tested seropositive for antibodies to the virus. Adjustment for test sensitivity and specificity yielded a 0.7% prevalence. Our findings will help us evaluate our public health response.

The magnitude of the coronavirus disease (CO-VID-19) pandemic is unknown because of a relatively large proportion of presumably asymptomatic persons (1–3). Reported infection rates, which mostly rely on PCR-based testing of symptomatic persons, may underestimate underlying infection rates. Analysis of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)-specific antibodies is required to