BRIEF REPORT



Extended Remdesivir Infusion for Persistent Coronavirus Disease 2019 Infection

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Persistent severe acute respiratory syndrome coronavirus 2 infection is difficult to treat. Here, we report a case of 5-month persistent coronavirus disease 2019 in an immunocompromised patient who was successfully treated with 30 consecutive days of remdesivir. Prolonged remdesivir infusion with concurrent cycle threshold monitoring might provide a potential solution to cure these patients with difficult-to-treat infections.

Keywords. Ct value; persistent COVID-19; prolonged remdesivir; remdesivir mutation; remdesivir resistance.

Severe persistent severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) infections have been reported in immunocompromised patients [1, 2]. However, the treatment strategies for these patients remain unclear and mainly consist of supportive care beyond the initial antiviral and immunotherapy [3]. Additionally, remdesivir was the only antiviral available in the inpatient setting. Monoclonal antibodies, nirmatrelvir and ritonavir, and molnupiravir are not approved for use in inpatient settings in the United States. Here we report a case of a patient with persistent coronavirus disease 2019 (COVID-19) proven by viral culture requiring a total of 5 hospital admissions. He was eventually treated with extended remdesivir infusion guided by nasopharyngeal SARS-CoV-2 polymerase chain reaction (PCR) to prevent symptomatic recurrence.

CASE REPORT

A 44-year-old patient with underlying granulomatosis with polyangiitis and secondary hypogammaglobulinemia presented to

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our hospital in August 2020 with fever, chills, exertional dyspnea, and diarrhea. His comorbidities included hypogammaglobulinemia, chronic kidney disease, hypertension, obesity, deep venous thrombosis, and gout. At the time of admission, patient was taking allopurinol, warfarin, prednisone, and monthly intravenous immunoglobulin G (IVIG). Patient received rituximab for 2 years until 4 years prior to admission for granulomatosis. This was followed by mycophenolate and then high-dose prednisone with slow taper. At the time of admission for coronavirus disease 2019 (COVID-19), he was on prednisone 5 mg daily. COVID-19 was diagnosed by positive nasopharyngeal SARS-CoV-2 PCR on day 1. He received 5 days of remdesivir plus 10 days of dexamethasone and was discharged on room air. He presented with similar symptoms and was admitted from days 17 to 37. During that admission, he received convalescent plasma along with supportive care. He again developed fever, dyspnea, and hypoxia that required a third hospitalization from days 52 to 70. He had progressive chest imaging findings consistent with COVID-19 pneumonia. Extensive infectious diseases workup was all negative except elevated β-D-glucan. He was empirically treated 21 days for Pneumocystis pneumonia and discharged home on 2 L of oxygen and on prophylaxis with atovaquone. Five days later (day 75), he was admitted again (fourth admission day 75-103) with fever and worsening dyspnea, and repeat nasopharyngeal SARS-CoV-2 PCR remained positive. He received 10 days of remdesivir plus dexamethasone with significant improvement of clinical symptoms. On day 110, he presented with fever and dyspnea to the emergency department. Nasopharyngeal SARS-CoV-2 PCR was positive and SARS-CoV-2 immunoglobulin M and immunoglobulin G were negative.

Inhaled remdesivir has also been shown to reduce viral burden in a nonhuman primate model of SARS-CoV-2 infection [4]. Remdesivir has been shown to significantly reduce SARS-CoV-2 viral load in nasopharyngeal (NP) swabs in hospitalized patients with COVID-19 infection [5]. At this point, we ventured a strategy of extended remdesivir administration based partly on cycle threshold (Ct) value as a guide but primarily on the goal of negative nasopharyngeal SARS-CoV-2 PCR results, which we interpreted as a marker of complete eradication of viable virus during his fifth admission (days 110-142). He received a total of 30 days of remdesivir until he had repeatedly tested negative for nasopharyngeal SARS-CoV-2 PCR twice 24 hours apart. The patient did receive dexamethasone, convalescent plasma, and IVIG treatment throughout the course of illness and hospitalization but never received other immunomodulators such as tocilizumab or JAK inhibitors. The clinical course and administration

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of remdesivir are shown in Figure 1*A*. The Ct value during remdesivir infusion was monitored (Figure 1*B*). The patient was monitored for side effects throughout the duration of treatment with daily examinations, and 2-3 times weekly complete blood count and comprehensive metabolic profile. After the completion of treatment we did monthly telephonic visits with the patient for 3 months and then quarterly. The only minor side effect reported was some hair thinning, which reversed quickly. We are unsure if that was due to a medication side effect or overall chronic illness from prolonged infection. The patient was successfully cured with 30 days of remdesivir (without dexamethasone) using this strategy and, at 12-month follow-up, there is no indication of recurrence of symptoms (Table 1).

VIRAL CULTURE, WHOLE GENOME SEQUENCING, AND REMDESIVIR EFFECTIVENESS ESSAY

SARS-CoV-2 isolates from NP specimens were sent for culture, proven to be viable, and subjected to whole genome sequencing (WGS) (Supplementary Materials and Methods). Unfortunately, the day 0 NP specimen was not available for sequencing. WGS revealed the virus' lineage was B.1.369 and contained the P314L mutation in the RNA-dependent RNA polymerase (RdRP) gene, encoded at ORF1b. B.1.369 belongs to the same phylogenetic



Figure 1. Clinical response to standard and extended infusions of remdesivir with corresponding cycle threshold values. Abbreviation: HD, hospital day.

Table 1. Admission Details, Clinical Presentation, Radiological Features, and Treatment

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DIS RA CRP: 0.8 mg/dL at 12 mo after discharge

cluster as the Iota (B.1.526), Beta (B.1.351), and Mu (B.1.621) variants (Supplementary Table 1A and Supplementary Figure 1A). This correlates well with the time these viruses emerged in the population. The likelihood that the sequenced viruses were different on day 0 (ie, reinfection vs persistent infection) because of missing sample is unlikely because it was within 90 days of the first presentation. We believe the chances of reinfection within 90 days during that phase of the pandemic and probably even today are highly unlikely. The guidance from the Centers for Disease Control and Prevention at that time was not to recollect NP specimen in a recent positive as it likely to be positive; hence, day 18 NP specimen was not collected. An in vitro remdesivir susceptibility assay demonstrated that our patient's isolate (B.1.369) was effectively inhibited by remdesivir, thus showing that the P314L nsp12/RdRp mutation seen by WGS had no effect on remdesivir activity (Supplementary Table 1B and Supplementary Figure 1B and 1C). No other nsp12/RdRp substitution was identified (Supplementary Table 1B). In a previous case of persistent SARS-CoV-2 viremia in an immunodeficient patient, a 76-year-old woman with a postrituximab B-cell immunodeficiency for whom remdesivir therapy failed, the authors found a new RdRP mutation, D484Y, that did not exist pretreatment. However, the lack of susceptibility testing did not conclusively prove association of treatment failure and resistance because of the reported new point mutation [6]. Another report did identify a new point mutation nsp12 E802D during treatment of a patient with acquired B-cell deficiency who had developed an indolent persistent SARS-CoV-2 infection and failed remdesivir therapy but ultimately responded after treatment with casirivimabimdevimab. In vitro experiments confirmed an approximately 6-fold increase in remdesivir inhibitory concentration 50 (IC₅₀) associated with the mutation, which also resulted in a fitness cost in the absence of remdesivir [7]. The same E802D mutation was previously identified in an in vitro remdesivir resistance selection experiment, and found to confer an approximately 2.5-fold increase in IC_{50} to the drug [8]. In our case, we were able to extend remdesivir treatment for a total of 30 days in this immunocompromised patient with persistent COVID-19 to overcome the lack of sustained virologic response to standard duration of 10 days and achieved clinical cure. We did not observe any side effects or laboratory abnormalities due to the extension. Remdesivir was (and still is) the only antiviral treatment option for hospitalized patients shaping this therapeutic decision for our patient. This is the first case report of extended remdesivir infusion beyond the standard 10 days of recommended therapy by the manufacturer. This supports further study of extended remdesivir therapy in immunocompromised patients with persistent infection and symptoms, as an alternative treatment management along with other reported option of monoclonal antibodies

[7, 9]. Our patient has remained symptom-free at 12 months of follow-up.

Supplementary Data

Supplementary materials are available at *Open Forum Infectious Diseases* online. Consisting of data provided by the authors to benefit the reader, the posted materials are not copyedited and are the sole responsibility of the authors, so questions or comments should be addressed to the corresponding author.

Notes

Author contributions. M. A. M. and T.-Y.C. contributed to data collection and manuscript writing. D. N., H. C., and M. H. contributed to polymerase chain reaction (PCR) and whole genome sequencing (WGS). L. H. and M. G. contributed to viral culture and remdesivir effective assay. G. C. contributed to technical review and manuscript writing, C. J. contributed to PCR, WGS, and manuscript writing.

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Patient consent. A signed informed consent from the patient was obtained, and the institutional review board of the Central Texas Veterans Health Care System, Temple, Texas, exempted this case report as per local policy.

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Potential conflicts of interest. G. C. is an employee and shareholder of Gilead Sciences. All other authors report no potential conflicts.

All authors have submitted the ICMJE Form for Disclosure of Potential Conflicts of Interest. Conflicts that the editors consider relevant to the content of the manuscript have been disclosed.

References

- Choi B, Choudhary MC, Regan J, et al. Persistence and evolution of SARS-CoV-2 in an immunocompromised host. N Engl J Med 2020; 383:2291–3.
- Aydillo T, Gonzalez-Reiche AS, Aslam S, et al. Shedding of viable SARS-CoV-2 after immunosuppressive therapy for cancer. N Engl J Med 2020; 383:2586–8.
- Helleberg M, Niemann CU, Moestrup KS, et al. Persistent COVID-19 in an immunocompromised patient temporarily responsive to two courses of remdesivir therapy. J Infect Dis 2020; 222:1103–7.
- Vermillion MS, Murakami E, Ma B, et al. Inhaled remdesivir reduces viral burden in a nonhuman primate model of SARS-CoV-2 infection. Sci Transl Med 2022; 14: eabl8282.
- Biancofiore A, Mirijello A, Puteo MA, et al. Remdesivir significantly reduces SARS-CoV-2 viral load on nasopharyngeal swabs in hospitalized patients with COVID-19: a retrospective case-control study. J Med Virol 2022; 94:2284–9.
- Martinot M, Jary A, Fafi-Kremer S, et al. Emerging RNA-dependent RNA polymerase mutation in a remdesivir-treated B-cell immunodeficient patient with protracted coronavirus disease 2019. Clin Infect Dis 2021; 73:e1762–5.
- Gandhi S, Klein J, Robertson AJ, et al. De novo emergence of a remdesivir resistance mutation during treatment of persistent SARS-CoV-2 infection in an immunocompromised patient: a case report. Nat Commun 2022; 13:1547.
- Szemiel AM, Merits A, Orton RJ, et al. In vitro selection of remdesivir resistance suggests evolutionary predictability of SARS-CoV-2. Plos Pathog 2021; 17: e1009929.
- Verderese JP, Stepanova M, Lam B, et al. Neutralizing monoclonal antibody treatment reduces hospitalization for mild and moderate coronavirus disease 2019 (COVID-19): a real-world experience. Clin Infect Dis 2021; 74:1063–9.