CASE REPORT Open Access

(2022) 11:28

Prolonged SARS-CoV-2 infection following rituximab treatment: clinical course and response to therapeutic interventions correlated with quantitative viral cultures and cycle threshold values

Christina S. Thornton¹, Kevin Huntley¹, Byron M. Berenger^{2,3}, Michael Bristow^{4,5}, David H. Evans⁶, Kevin Fonseca^{2,7}, Angela Franko³, Mark R. Gillrie^{1,7}, Yi-Chan Lin⁶, Marcus Povitz¹, Mona Shafey^{1,8}, John M. Conly^{1,7,10*†} and Alain Tremblay^{1,9*†}

Abstract

Background: Detection of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) RNA is completed through reverse transcriptase-PCR (RT-PCR) from either oropharyngeal or nasopharyngeal swabs, critically important for diagnostics but also from an infection control lens. Recent studies have suggested that COVID-19 patients can demonstrate prolonged viral shedding with immunosuppression as a key risk factor.

Case presentation: We present a case of an immunocompromised patient with SARS-CoV-2 infection demonstrating prolonged infectious viral shedding for 189 days with virus cultivability and clinical relapse with an identical strain based on whole genome sequencing, requiring a multi-modal therapeutic approach. We correlated clinical parameters, PCR cycle thresholds and viral culture until eventual resolution.

Conclusions: We successfully demonstrate resolution of viral shedding, administration of COVID-19 vaccination and maintenance of viral clearance. This case highlights implications in the immunosuppressed patient towards infection prevention and control that should consider those with prolonged viral shedding and may require ancillary testing to fully elucidate viral activity. Furthermore, this case raises several stimulating questions around complex COVID-19 patients around the role of steroids, effect of antiviral therapies in absence of B-cells, role for vaccination and the requirement of a multi-modal approach to eventually have successful clearance of the virus.

Keywords: SARS-CoV-2, Immunocompromised patient, Viral shedding, Therapeutics

Full list of author information is available at the end of the article

Background

The coronavirus disease 2019 (COVID-19) pandemic has had a global impact with significant morbidity and mortality. A positive upper respiratory or nasopharyngeal (NP) swab reverse-transcription polymerase chain reaction (RT-PCR) molecular amplification test is commonly used to confirm the presence of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) virus.



© The Author(s) 2022. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/by/4.0/. The Creative Commons Public Domain Dedication waiver (http://creativecommons.org/publicdomain/zero/1.0/) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

^{*}Correspondence: atrembla@ucalgary.ca; jconly@ucalgary.ca

[†]John M. Conly and Alain Tremblay: Senior corresponding authors contributed equally to this manuscript

¹ Department of Medicine, Cumming School of Medicine, University of Calgary, Calgary, AB, Canada

Risk factors for severe disease include older age, medical comorbidities, and immunosuppressed status [1].

Our understanding of COVID-19 has evolved as persistence of SARS-CoV-2 infectious viral shedding and recurrence of PCR positivity after one or more negative samples have been documented in both asymptomatic and symptomatic patients [2]. While RT-PCR is a rapid and easily accessible tool, limitations include prolonged positivity in the absence of infectious virus, thus confounding clinical management around possible reinfection or relapse. Understanding the natural progression of SARS-CoV-2 infectious viral shedding is critical from both an individual's management but also for public health and hospital infection control. Initial studies suggest infectivity declines within ten days in immunocompetent individuals [3]. Immunocompromised patients (ICPs) represent a unique population whereby prolonged infectious viral shedding and infectivity may occur with disease relapse. We present a complex ICP demonstrating protracted infectious viral shedding for 189 days with three distinct COVID-19 clinical relapses associated with waxing and waning of positive viral cultures and RT-PCR cycle threshold (Ct) values.

Methods

Nasopharyngeal (NP) swabs were collected at regular intervals beginning shortly after admission and tested by RT-PCR, quantitative culture [4] and whole genome sequencing (Additional file 1: Supplementary Material). The patient provided informed consent with approval of the University of Calgary's Health Research Ethics Board (no. 20-0444).

Case presentation

Preceding availability of COVID-19 vaccination, a 58-year-old Caucasian male presented to his primary care provider with one-week of cough, low-grade fever, and general malaise. He was diagnosed with follicular lymphoma 10 months prior and received 6 cycles of chemo-immunotherapy (bendamustine and rituximab), achieving clinical remission, and received an additional

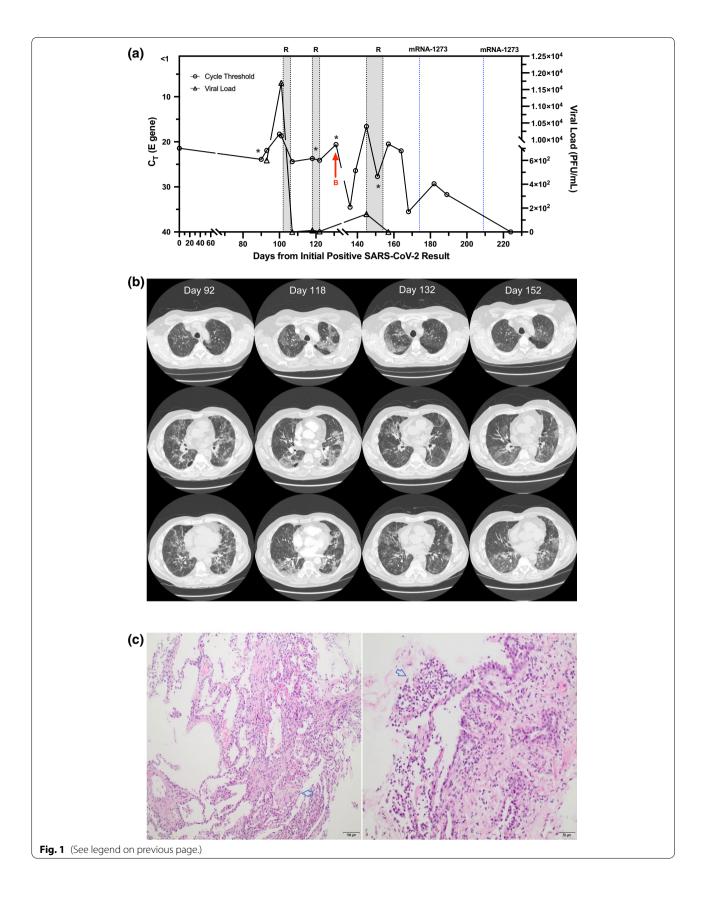
dose of maintenance rituximab one month prior. An oropharyngeal swab was positive for SARS-CoV-2, but he did not require hospitalization at that time.

Twenty-seven days after the initial positive swab, he developed hypoxemia prompting hospital admission. Initial NP swab was negative (Fig. 1A) and computed tomography (CT) imaging demonstrated recurrent ground glass opacities (Fig. 1B) with bilateral interstitial pneumonia, compatible with COVID-19. Blood tests revealed lymphopenia $(0.1\times10^9/L, \text{ normal: } 0.7-3.5\times10^9/L)$ with increased C-reactive protein (57.0 mg/L, normal: 0.0-8.0 mg/L) and ferritin (2047 µg/L, normal: 30-500ug/L). He improved with dexamethasone 6 mg daily for 10 days. However due to relapsing hypoxemia and radiographic progression, a diagnosis of organizing pneumonia as sequelae of COVID-19 was considered. Bronchoscopy was declined, and oral prednisone 50 mg daily was initiated with a planned dose taper.

On day 90, 73 days post admission, following ongoing reduction of prednisone (10 mg), the patient presented with new onset of pyrexia and recurrence of hypoxemia. RT-PCR testing was again positive by NP swab (Fig. 1A, Ct 18.3). The patient was re-initiated on dexamethasone with bronchoscopy performed. Bronchoalveolar lavage was both SARS-CoV-2 RT-PCR (Ct 18.7) and culture positive $(1.17 \times 10^4 \text{ pfu/mL}, \text{ wild-type G clade})$ and negative for bacterial, mycobacterial, other viral and fungal growth. Transbronchial biopsies (Fig. 1C) demonstrated active pneumonitis with occasional foci of organizing pneumonia, and patchy interstitial fibrosis. Symptoms were consistent with COVID-19 relapse and remdesivir was given (200 mg intravenous loading dose then 100 mg daily) for 5 days concurrently with dexamethasone (Fig. 1A). On day 107, there was clinical improvement with reduction in oxygen requirements to room air and RT-PCR Ct of 24.4 Subsequent investigations revealed hypogammaglobinemia (IgA 0.48 g/L, IgG 3.29 g/L, IgM 0.05 g/L (normal: 0.6–4.2 g/L, 6.8–18.0 g/L, 0.4-3.00 g/L, respectively), likely secondary to prior rituximab administration. He was initiated on monthly intravenous immunoglobulin. He clinically improved and

(See figure on next page.)

Fig. 1 Depiction of prolonged COVID-19 viral shedding in an immunosuppressed patient. A Timeline of patient's clinical course from time of initial COVID-19 positive swab (day 0). Cycle threshold values and culture viral load are represented. Treatment modalities throughout the clinical course are depicted. R: Remdesivir; B: bamlanivimab; *indicates date of corresponding CT scans depicted in B. Cycle threshold values greater than 40 were considered negative. B Serial CT scans show multifocal, patchy areas of predominantly ground-glass opacities. The first CT scan shown (day 92) was obtained at the time of the second relapse. The subsequent studies (days 118, 132 and 152) were obtained during the third relapse. On each of the serial scans, some areas of involvement improve while there is worsening in other areas, consistent with migratory opacities. There are features of organizing pneumonia, which is typical later in the course of COVID-19 pneumonia. C Transbronchial biopsy. Left image: Low power view of transbronchial biopsy shows alveolated lung parenchyma with expansive interstitial cellular infiltrates composed of lymphocytes and neutrophils. There are subtle occasional foci of organizing pneumonia (blue arrow highlights an intra-alveolar fibroblastic plug of organizing pneumonia) and patchy interstitial fibrosis (hematoxylin–eosin, at original magnification × 10). Right image: High power view with focus of intra-alveolar neutrophils (highlighted with blue arrow) with associated reactive type II pneumocytes lining alveolar spaces (hematoxylin–eosin, at original magnification × 20)



was discharged home on day 107 with an oral dexamethasone (4 mg) taper. Viral culture load at time of discharge was not detectable.

On day 118 the patient re-presented to hospital with respiratory failure. A repeat SARS-CoV-2 RT-PCR swab was positive with Ct of 23.7and a viral culture load of 1.25×10^{1} pfu/mL, suggestive of ongoing viral persistence and a third COVID-19 relapse (Fig. 1A). SARS-CoV-2 testing failed to demonstrate remdesivir resistance (supplementary material). He was re-treated with another course of remdesivir for 5 days with modest clinical improvement. Repeat CT scans demonstrated persistent, migratory bilateral ground glass opacities (Fig. 1B). Subsequent medical therapy included bamlanivimab 700 mg (day 133; one dose), IVIG 75 mg (days 137, 165 and 193; one dose each) and a third course of remdesivir (day 146; ten-day course). The patient demonstrated a gradual clinical recovery with progressive reduction in oxygen therapy and eventual negative swab by RT-PCR on day 189. On days 174 and 202, the mRNA-1273 COVID-19 vaccine was administered. On day 202 after initial presentation, he was ultimately discharged home with repeat COVID-19 swabs (days 209 and 224) negative. He has continued to improve with gradually increasing exercise tolerance.

Notably, whole genome sequencing of the patient's viral samples over the duration of the time-period demonstrated an identical sequence belonging to SARS-CoV-2 clade D614G equivalent to the PANGO B.1 lineage viral strain with a specific sublineage of B.1.160. A review of the local epidemiology during this period noted B.1.160 as the 12th highest lineage, accounting for only 1.4% of the those sequenced from the general population. Based on the sequencing results, relapse of infection by an identical virus strain was demonstrated rather than re-infection with different strains. Finally, serology testing completed was negative to the nucleocapsid antibody, however positive for the spike antibody (value of 1515.3AU/mL; positive threshold of > 50AU/mL).

Discussion and conclusions

Cases of prolonged infectious viral shedding are emerging in the literature, including B-cell deficiency where shedding of greater than 100 days has been reported [5, 6]. The impact of COVID-19 on immune-deficient individuals and optimal treatment is unknown. Reports of patients post B-cell depletion therapy with spontaneous clearance of SARS-CoV-2 are known; however, others undergoing rituximab therapy had severe and prolonged COVID-19 infections with persistent detectable virus [7].

This case adds to emerging literature on COVID-19 relapse and prolonged SARS-CoV-2 infectious virus shedding in the ICP. Such individuals may pose important

infection prevention concerns for similar patient populations as they are often cared in clinical settings with other ICPs and may serve as unrecognized viral reservoirs for superspreading local transmission. Current CDC guidelines recommend duration of isolation for severely immunocompromised COVID-19 patients to be up to 20 days [1]. However, emerging reports have clearly demonstrated prolonged infectious viral shedding far exceeding this timeframe. In the hospital setting, isolation guidelines are of critical importance as prolonged infectious viral shedding poses risk to both patients and healthcare workers.

Emerging data has highlighted the difficulty in SARS-CoV-2 eradication in those with blunted immune responses, prolonged viral shedding, and protracted clinical courses. Rituximab is a chimeric monoclonal antibody that binds to the CD20 molecule on the surface of B-cells leading to B-cell depletion lasting 6-9 months. Theoretically, the CD20 surface molecule is lost during B-cell differentiation into plasma cells, the major source of circulating immunoglobulins, and effects on immunoglobulin production after short courses of rituximab should be a rare event but real-world experience has demonstrated reduced response to vaccines and reactivation of viral infections, such as COVID-19. A recent retrospective study revealed up to a 45% incidence of hypogammaglobinemia in rituximab-treated individuals, persisting up to 42 months, and 4% of patients failing to recover CD20 cellular function [8]. Naïve B-cells, disproportionally depleted by rituximab, are critical to produce neutralizing antibodies in COVID-19 [9]. Similarly, rituximab-treated patients often fail to respond to SARS-CoV2 vaccines, suggesting that B-cells are required not only for viral clearance during natural infection but also for adequate responses to vaccination [9].

Treatments including remdesivir and dexamethasone have shown modest benefit in the general population, but efficacy is unclear in ICPs. The use of COVID-19 convalescent plasma in severely ICPs due to prior treatment with anti-CD20 monoclonal antibodies had early promise with favourable safety efficacy early in disease course, but a recent randomized control trial failed to demonstrate benefit [10]. Similarly, while no strong evidence exists for use of anti-SARS-CoV-2 monoclonal antibody treatment in ICPs, a rational argument for their use in such situations may be made, and in our case resulted in a significant but transient improvement in Ct values. Our study was unique in utilizing a multimodal therapeutic targeted approach and successfully allowing vaccination against COVID-19, while highlighting the profound effect of rituximab on the B cell lineage.

The patient's clinical course may also have been impacted by corticosteroid treatment. While

dexamethasone can improve mortality in severe COVID-19 illness, evidence in immunosuppressed individuals is lacking. The development of organizing pneumonia following COVID-19 illness and associated use of prolonged courses of corticosteroids have been described [11], as highlighted in our case with hypoxemia improving with escalating steroid doses. However, despite initial improvement, additional immunosuppression may have contributed to prolonged viral replication with eventual viral clearance coinciding with tapering of corticosteroid treatment to physiologic doses. Regardless of the mechanisms of viral clearance, all anti-viral therapies attempted provided modest improvement in Ct values and cultivatable virus, although often with transient effect, highlighting the underlying impact of host immune defects.

In conclusion, our case describes a prolonged clinical course with multiple COVID-19 relapses in an immunosuppressed lymphoma patient caused by the same strain of SARS-CoV-2. Our findings have larger implications towards infection prevention and control guidelines that should consider prolonged isolation and rigorous followup. Use of quantitative PCR that are well validated and correlated with viral culture titres in parallel with Ct values may serve as a powerful clinical tool to elucidate infectious viral shedding, guide treatments, and assess outcomes. Response to specific anti-viral treatments may be seen, but repeated interventions are likely needed due to relapse while waiting for effects of immunosuppressive agents such as rituximab to diminish. The use of corticosteroids for organizing pneumonia needs to be balanced carefully with the risk of compounding pre-existing immunosuppression.

Abbreviations

SARS-CoV-2: Severe acute respiratory syndrome coronavirus 2; RNA: Ribonucleic acid; RT-PCR: Reverse transcriptase-polymerase chain reaction; COVID-19: Coronavirus disease 2019; ICP: Immunocompromised patients; CT: Computed tomography; IVIG: Intravenous immune globulin; Ct: Cycle threshold; CDC: Centers for disease control and prevention.

Supplementary Information

The online version contains supplementary material available at https://doi.org/10.1186/s13756-022-01067-1.

Additional file 1. Viral culture load at time of discharge was not detectable (Fig. 1A).

Acknowledgements

We thank the patient in this manuscript for granting permission to share his story.

Authors' contributions

CST, KH, MG, MP, MS, JMC and AT were involved in the clinical care of the patient described in this case. BMB, DHE, KF, and YL provided the laboratory results with viral culture titres and cycle thresholds. MB provided radiographic

analysis. AF provided pathological analysis. CST created the first draft of the manuscript and all authors assisted in the creation of the final manuscript. All authors read and approved the final manuscript.

Funding

Not applicable.

Availability of data and materials

Not applicable.

Declarations

Ethics approval and consent to participate

Not applicable.

Consent for publication

Written informed consent was obtained from the patient for publication of this case report and any accompanying images. A copy of the written consent is available for review by the Editor-in-Chief of this journal.

Competing interests

The authors have none to declare relevant to this manuscript.

Author details

¹Department of Medicine, Cumming School of Medicine, University of Calgary, Calgary, AB, Canada. ²Alberta Precision Laboratories, Alberta Public Health Laboratory, Calgary, AB, Canada. ³Department of Pathology and Laboratory Medicine, University of Calgary, Calgary, AB, Canada. ⁴Department of Radiology, University of Calgary, Calgary, AB, Canada. ⁵Department of Cardiac Sciences, University of Calgary, Calgary, AB, Canada. ⁶Department of Medical Microbiology and Immunology and Li Ka Shing Institute of Virology, University of Alberta, Calgary, AB, Canada. ⁷Department of Microbiology, Immunology and Infectious Diseases, University of Calgary, Calgary, AB, Canada. ⁸Tom Baker Cancer Centre, University of Calgary, Calgary, AB, Canada. ⁹Foothills Medical Centre, 3300 Hospital Drive NW, Calgary, AB T2N4N1, Canada. ¹⁰Foothills Medical Centre, SSB, 1403 29 St NW, Calgary, AB T2N2T9, Canada.

Received: 7 October 2021 Accepted: 21 January 2022 Published online: 05 February 2022

References

- Huang C, Wang Y, Li X, et al. Clinical features of patients infected with 2019 novel coronavirus in Wuhan, China. Lancet. 2020;395(10223):497–506.
- Oran DP, Topol EJ. The proportion of SARS-CoV-2 infections that are asymptomatic: a systematic review. Ann Intern Med. 2021;174(5):655–62.
- Van Kampen JJA, Van de Vijver D, Fraaij PLA, et al. Duration and key determinants of infectious virus shedding in hospitalized patients with coronavirus disease-2019 (COVID-19). Nat Commun. 2021;12(1):267.
- Rajakumar I, Isaac DL, Fine NM, et al. Extensive environmental contamination and prolonged severe acute respiratory coronavirus-2 (SARS CoV-2) viability in immunosuppressed recent heart transplant recipients with clinical and virologic benefit with remdesivir. Infect Control Hosp Epidemiol. 2021: 1–3.
- Baang JH, Smith C, Mirabelli C, et al. Prolonged severe acute respiratory syndrome coronavirus 2 replication in an immunocompromised patient. J Infect Dis. 2021;223(1):23–7.
- 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(23):2291–3.
- Yasuda H, Tsukune Y, Watanabe N, et al. Persistent COVID-19 pneumonia and failure to develop anti-SARS-CoV-2 antibodies during rituximab maintenance therapy for follicular lymphoma. Clin Lymphoma Myeloma Leuk. 2020;20(11):774–6.
- Roberts DM, Jones RB, Smith RM, et al. Rituximab-associated hypogammaglobulinemia: incidence, predictors and outcomes in patients with multi-system autoimmune disease. J Autoimmun. 2015;57:60–5.

- Sette A, Crotty S. Adaptive immunity to SARS-CoV-2 and COVID-19. Cell. 2021;184(4):861–80.
- Simonovich VA, Burgos Pratx LD, Scibona P, et al. A randomized trial of convalescent plasma in Covid-19 severe pneumonia. N Engl J Med. 2021;384(7):619–29.
- Kory P, Kanne JP. SARS-CoV-2 organising pneumonia: "Has there been a widespread failure to identify and treat this prevalent condition in COVID-19?" BMJ Open Respir Res. 2020;7(1):e000724.
- Berenger BM, Conly JM, Fonseca K, et al. Saliva collected in universal transport media is an effective, simple and high-volume amenable method to detect SARS-CoV-2. Clin Microbiol Infect. 2021;27(4):656–7.
- Freed NE, Vlkova M, Faisal MB, Silander OK. Rapid and inexpensive whole-genome sequencing of SARS-CoV-2 using 1200 bp tiled amplicons and Oxford Nanopore Rapid Barcoding. Biol Methods Protoc. 2020;5(1):bpaa014.
- Resende PC, Motta FC, Roy S, Appolinario L, Fabri A, Xavier J, Harris K, Matos AR, Caetano B, Orgeswalska M, Miranda M, Garcia C, Abreu A, Williams R, Breuer J, Siqueira MM. 2020. SARS-CoV-2 genomes recovered by long amplicon tiling multiplex approach using nanopore sequencing and applicable to other sequencing platforms. bioRxiv. https://doi.org/10. 1101/2020.04.30.069039.
- Zelyas N, Pabbaraju K, Croxen MA, Lynch T, Buss E, Murphy SA, et al. Precision response to the rise of the SARS-CoV-2 B.1.1.7 variant of concern by combining novel PCR assays and genome sequencing for rapid variant detection and surveillance. Microbiol Spectr. 2021;9(1):e0031521.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Ready to submit your research? Choose BMC and benefit from:

- fast, convenient online submission
- thorough peer review by experienced researchers in your field
- rapid publication on acceptance
- support for research data, including large and complex data types
- gold Open Access which fosters wider collaboration and increased citations
- maximum visibility for your research: over 100M website views per year

At BMC, research is always in progress.

Learn more biomedcentral.com/submissions

