

# Diffuse Alveolar Hemorrhage (DAH) Associated with Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2): A Case Report

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## Research Article

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# Abstract

Diffuse alveolar hemorrhage (DAH) is a rare syndrome resulting from the accumulation of intra-alveolar red blood cells (RBCs) originating most often from the alveolar capillaries and, less frequently, from precapillary arterioles or postcapillary venules. The causes of DAH can be divided into infectious and noninfectious. Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) is a novel coronavirus that has not been previously identified in humans, and it is responsible for the coronavirus disease-19 (COVID-19) infection. The long-term complications of COVID-19 are not entirely known and are still being investigated. Here, we present a case of DAH that is believed to be a consequence of COVID-19 infection.

## Introduction

Diffuse alveolar hemorrhage (DAH), commonly recognized by the signs of anemia, hemoptysis, diffuse radiographic pulmonary infiltrates, and hypoxemic respiratory failure, is a rare syndrome resulting from the accumulation of intra-alveolar red blood cells (RBCs) originating most often from the alveolar capillaries and, less frequently, from precapillary arterioles or postcapillary venules [1–3]. The prognosis of DAH is poor, with hospital mortality ranging from 20% to over 50% [4]. The causes of DAH can be divided into infectious and noninfectious. Pulmonary infections include those caused by viruses, bacteria, fungi, and parasites [3]. In immunocompromised patients, the common infectious diseases that may cause DAH can consist of cytomegalovirus (CMV), adenovirus, invasive aspergillosis, *Mycoplasma*, *Legionella*, and *Strongyloidiasis*. However, in immunocompetent patients, the infections that may cause DAH can include influenza A (H1N1), dengue, leptospirosis, and malaria [5–14].

Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) is a novel coronavirus that has not been previously identified in humans and is responsible for the coronavirus disease-19 (COVID-19) infection. The first case was identified in Wuhan, China, in December 2019, and it has spread to many countries around the world and was declared a global pandemic on March 11, 2020, by the World Health Organization (WHO). However, to our knowledge, the association between DAH and COVID-19 infection has not been reported. Here, we present a case of DAH that is believed to be a consequence of COVID-19 infection.

## Case Report

This report describes a 26-year-old Caucasian female with no known past medical history who presented to the Piedmont Columbus Regional Midtown emergency department with a chief complaint of worsening shortness of breath and dyspnea on exertion. The patient was found to be positive for COVID-19 around 25 days before this admission. The patient reported that she had isolated herself for ten days during that time and that her symptoms had improved. However, one week before the current admission, she started experiencing worsening shortness of breath and cough with hemoptysis. These symptoms prompted her visit to an emergency department, where she was subsequently diagnosed with atypical pneumonia and discharged on doxycycline [100 mg orally every 12 hours for five days].

The patient continued to worsen and was subsequently found to have generalized malaise, severe fatigue, non-productive cough, myalgias, fever, decreased appetite, and hemoptysis in the current admission. The patient had no family history of bleeding disorders. Her vital signs on admission were the following: temperature, 100.8°F (38.2°C); blood pressure, 128/72 mmHg; heart rate, 122 beats per minute; respiratory rate, 35 breaths per minute; and oxygen (O<sub>2</sub>) saturation, 90% on room air, for which she required a 100% nonrebreather for oxygen support. An arterial blood gas (ABG) panel was performed and revealed the following: pH of 7.43; pCO<sub>2</sub> of 32.9 mmHg; pO<sub>2</sub> of 73.6 mmHg; HCO<sub>3</sub> of 21.9 mmol/L. The patient had an elevated white blood cell (WBC) count of 11,200 cells/mcL [normal range, 4000–10500 cells/mcL], severe anemia with hemoglobin of 6.9 g/dL [normal range, 12.0–16 g/dL], hematocrit of 20.6% [normal range, 36.0–48.0%], and a platelet count of 432 10<sup>3</sup>/mcL [normal range, 130–400 10<sup>3</sup>/mcL]. The repeat COVID-19 test during this admission using the Xpert® Xpress SARS-CoV-2/Flu/RSV test (Cepheid®, Sunnyvale, California, United States) and the SARS-COV-2 Immunoglobulin G (IgG) antibody were positive, which indicated that she was in the late stage of the infection.

The chest computed tomography (CT) scan was negative for pulmonary embolism (PE). However, it revealed extensive bilateral ground-glass opacities with confluent densities suggestive of multifocal pneumonia. Consequently, pulmonology and infectious diseases were consulted for evaluation and management. The patient was started on community-acquired pneumonia (CAP) treatment with ceftriaxone [1 g intravenously (IV) every 24 hours for five days] and azithromycin [500 mg IV every 24 hours for three days]. Additionally, there was strong suspicion for diffuse alveolar hemorrhage (DAH), most likely due to a COVID-19-associated vasculitis process, for which methylprednisolone [250 mg IV four times per day] was started.

A series of serologic tests were ordered to investigate the cause of DAH. The patient was found to be positive for perinuclear anti-neutrophil cytoplasmic antibodies (P-ANCA) [1:40 Titer; normal range, < 1:20 Titer]. She was also found to be negative for antinuclear antibody (ANA), anti-double-stranded deoxyribonucleic acid (anti-dsDNA), and cyclic citrullinated peptide (CCP) antibodies. Additionally, her total complement (CH50) was 49 U/mL [normal range, 31–60 U/mL], her C3 complement was 141 mg/dL [normal range, 90–180 mg/dL], and her C4 complement was 16 mg/dL [normal range, 10–40 mg/dL]. Based on these serologic findings, the patient was thought to have microscopic polyangiitis.

At the time of discharge, the patient had an oxygen saturation of 96% on room air and a respiratory rate of 18 breaths per minute with no evidence of ongoing hypoxia. The patient was prescribed a tapering course of prednisone [40 mg orally twice daily for five days, then 30 mg orally twice daily for five days, then 20 mg orally twice daily for five days, then 10 mg orally twice daily for five days, then 10 mg orally once daily for five days] and advised to follow up with both pulmonology and her primary care physician.

Three days after discharge, the anti-glomerular basement membrane (anti-GBM) antibodies were found to be positive [4.5 AI; normal range, < 1.0 AI]. In addition, the urinalysis (UA) obtained during admission confirmed the presence of blood and protein. The patient was informed about the positive results and advised to schedule an appointment with a nephrologist for further assessment.

A renal biopsy was obtained during her visit with the nephrologist. The renal parenchyma available for light microscopic examination was represented by approximately 90% cortex and 10% medulla. Forty-eight glomeruli were present, none of which were globally sclerotic. Two glomeruli showed segmental rupture of the capillary loops associated with fibrinoid necrosis of the glomerular tuft and cellular crescent formation. Twenty-six additional glomeruli showed areas of segmental glomerulosclerosis, eleven of which were associated with fibro-cellular crescents and fifteen with fibrous crescents. The twenty uninvolved glomeruli showed no mesangial or endocapillary hypercellularity. The findings of the light microscopy can be seen in Fig. 1a and Fig. 1b.

The renal parenchyma available for immunofluorescence studies was entirely represented by cortex and contained nine glomeruli, none of which were globally sclerotic. The sections were stained for IgG, IgM, IgA, C3, complement 1q (C1q), albumin, fibrinogen, kappa, and lambda light chains. In addition, there was focal intense fibrinogen staining within the glomerular tuft. Additionally, there was mild mesangial finely granular staining for IgA (trace), IgG (2+), kappa (2+), and lambda (1+). All other stains were negative within the glomeruli. The findings of the immunofluorescence studies can be seen in Fig. 2a and Fig. 2b.

Based on the serology results mentioned above and the renal biopsy findings, the patient was determined to be having ANCA-associated vasculitis with glomerulonephritis and was started on methylprednisolone [500 mg IV daily for three days], followed by rituximab [375 mg/m<sup>2</sup> IV once weekly for four weeks].

## Discussion

A variety of diseases and conditions can be associated with the development of diffuse alveolar hemorrhage. One of the histopathologic patterns of DAH is pulmonary, or alveolar, capillaritis, which is characterized by the neutrophilic infiltration of the alveolar septa or the lung interstitium [4, 15]. Although bronchoscopy with bronchoalveolar lavage (BAL) is indicated in patients suspected of having DAH to confirm the diagnosis [16], this procedure was not performed in our case due to the risk of aerosolization and spread of COVID-19 [17].

Anti-neutrophil cytoplasmic antibody (ANCA)-associated vasculitis (AAV) is a group of autoimmune diseases including granulomatosis with polyangiitis (GPA) (formerly Wegener's granulomatosis), eosinophilic granulomatosis with polyangiitis (formerly Churg-Strauss syndrome), and microscopic polyangiitis (MPA). These diseases are characterized by inflammatory cell infiltration causing necrosis of small blood vessels [18–20]. Our patient was thought to have diffuse alveolar hemorrhage caused by ANCA-associated vasculitis, but with the specific disease not wholly identified.

A positive ANCA test can indicate if DAH is due to GPA or microscopic polyangiitis. Cytoplasmic ANCA (C-ANCA), as determined by anti-proteinase 3 antibodies (anti-PR3), is most consistent with GPA. At the same time, perinuclear ANCA (P-ANCA) with positive anti-myeloperoxidase (anti-MPO) favors the diagnosis of microscopic polyangiitis or eosinophilic granulomatosis with polyangiitis syndrome [21]. Anti-GBM antibodies are more specific to an anti-GBM disease or Goodpasture's syndrome, an

autoimmune disorder characterized by autoantibodies directed against the glomerular basement membrane. However, 60 to 80% of the patients with anti-GBM disease have clinical manifestations of pulmonary and renal disease [22]. In addition, patients with acute glomerulonephritis with or without pulmonary hemorrhage also may have GPA or MPA. Thus, the serum should be tested for ANCA as well as anti-GBM antibodies. Anti-GBM disease and systemic vasculitis not only have similar clinical manifestations, but between 10% and 50% of patients with the anti-GBM disease also test positively for ANCA (usually anti-MPO-ANCA) at the time of diagnosis and may have signs of a systemic vasculitis or a marked systemic inflammatory response. Based on our patient's serologic findings throughout her admission, she was diagnosed initially with microscopic polyangiitis. However, the discovery of the positive anti-GBM antibodies raised the suspicion for possible kidney involvement, which was subsequently confirmed by renal biopsy, even though the patient did not have any signs of kidney injury during her admission.

COVID-19 infection is classified into three stages: 1) early infection; 2) pulmonary involvement with or without hypoxia; 3) and the severe systemic hyper inflammation stage. The late stage of COVID-19 is characterized by the elevation of systemic inflammation markers [23]. It has been reported that inflammatory cytokines and biomarkers such as Interleukins (e.g., IL-2, IL-6, and IL-7), granulocyte colony-stimulating factor, macrophage inflammatory protein 1-a, tumor necrosis factor (TNF)-a, C-reactive protein (CRP), ferritin, and D-dimer are significantly elevated during this stage [24]. In addition, some of these inflammatory markers, including IL-6, IL-8, and TNF-a, are elevated as well with ANCA-associated vasculitis [21]. Since our patient was in the late stage of her COVID-19 infection, which was supported by the time since the onset of her COVID-19 symptoms, her positive SARS-COV-2 IgG antibodies, and her inflammatory markers, it was possible that the elevation of inflammatory markers at the time of admission could have contributed to the incidence of ANCA-associated vasculitis and, subsequently, the diffuse alveolar hemorrhage.

## Conclusion

We presented a rare case of diffuse alveolar hemorrhage in a patient following infection with SARS-COV-2. The underlying cause is thought to be ANCA-associated vasculitis with glomerulonephritis. The association between COVID-19 infection and DAH is not fully known. However, the inflammatory process of COVID-19 infection may have a role in vasculitis, leading to DAH. Nevertheless, this case cannot establish a causal relationship between the two conditions, but it describes a clinical observation that might need to be considered when evaluating COVID-19 patients.

## Declarations

### Funding

The authors received no funding for this publication.

## Conflicts of Interest/Competing Interests

All the authors report no potential conflicts of interest.

## Availability of Data and Material

Not applicable.

## Code Availability

Not applicable.

## Authors' Contributions

The first draft of the manuscript was written by Haytham Wali, and all authors commented on previous versions of the manuscript. All authors read and approved the final manuscript.

## Ethics Approval

The Columbus Regional Research Integrity Panel has confirmed that no ethical approval is required.

## Consent to Publish

The patient has consented to the submission of the case report to the journal.

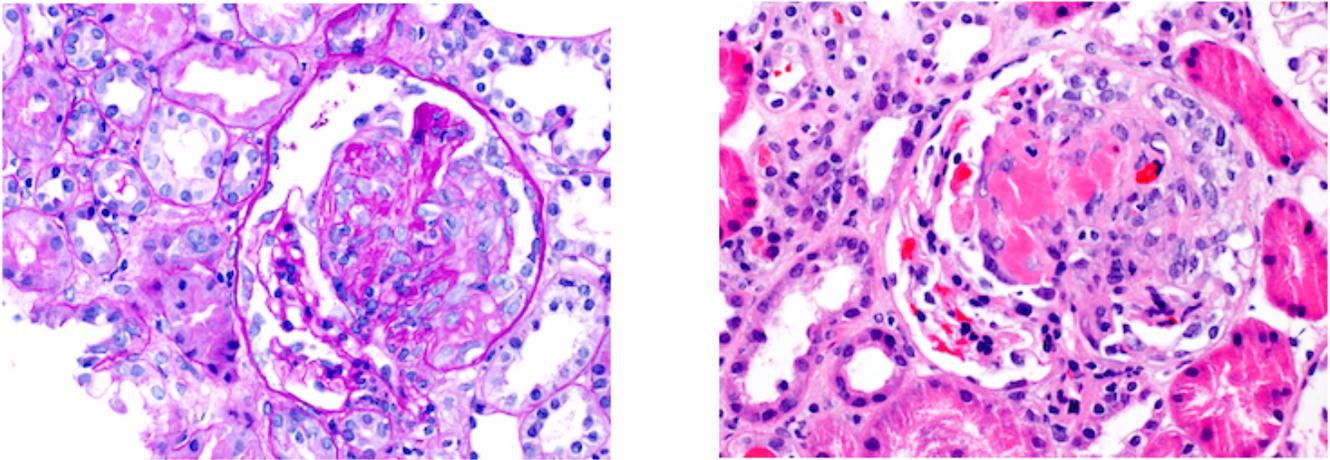
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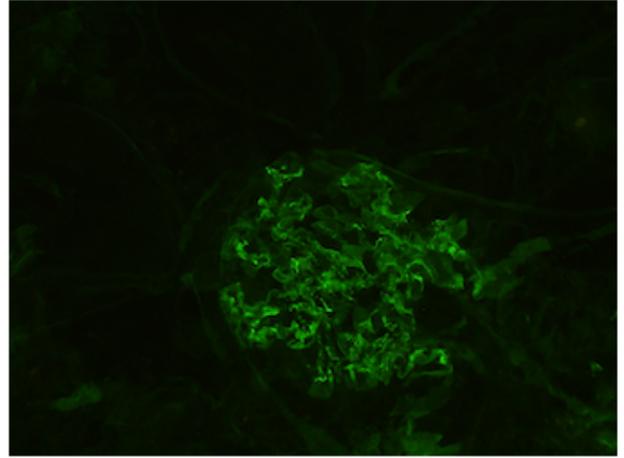
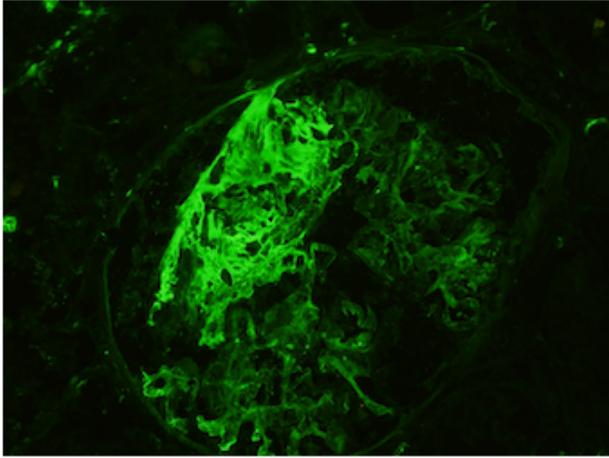
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## Figures



**Figure 1**

1a Light microscopy of the renal biopsy showing fibrocellular crescent. 1b Light microscopy of the renal biopsy showing fibrinoid necrosis.



**Figure 2**

2a The immunofluorescence stain of the fibrinogen in the renal biopsy. 2b The immunofluorescence stain of the immunoglobulin G (IgG) in the renal biopsy.