



# Treatment of Coronavirus Disease 2019 (COVID-19) Patients with Convalescent Plasma

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Coronavirus disease 2019 (COVID-19), caused by severe acute respiratory syndrome coronavirus 2, has spread globally, and no proven treatments are available. Convalescent plasma therapy has been used with varying degrees of success to treat severe microbial infections for >100 years. Patients ( $n = 25$ ) with severe and/or life-threatening COVID-19 disease were enrolled at the Houston Methodist hospitals from March 28, 2020, to April 14, 2020. Patients were transfused with convalescent plasma, obtained from donors with confirmed severe acute respiratory syndrome coronavirus 2 infection who had recovered. The primary study outcome was safety, and the secondary outcome was clinical status at day 14 after transfusion. Clinical improvement was assessed on the basis of a modified World Health Organization six-point ordinal scale and laboratory parameters. Viral genome sequencing was performed on donor and recipient strains. At day 7 after transfusion with convalescent plasma, nine patients had at least a one-point improvement in clinical scale, and seven of those were discharged. By day 14 after transfusion, 19 (76%) patients had at least a one-point improvement in clinical status, and 11 were discharged. No adverse events as a result of plasma transfusion were observed. Whole genome sequencing data did not identify a strain genotype-disease severity correlation. The data indicate that administration of convalescent plasma is a safe treatment option for those with severe COVID-19 disease. (*Am J Pathol* 2020, ■: 1–11; <https://doi.org/10.1016/j.ajpath.2020.05.014>)

Q8 The coronavirus disease 2019 (COVID-19) pandemic has spread globally and caused massive loss of life and economic hardship. As of May 2, 2020, there were 3,494,671 confirmed cases and 246,475 deaths worldwide, and in the United States, there were 1,154,340 confirmed cases and 67,447 deaths (Johns Hopkins University, <https://coronavirus.jhu.edu/map.html>, last accessed May 2, 2020). The disease is caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), a highly transmissible coronavirus first identified in Wuhan, China.<sup>1–3</sup> SARS-CoV-2 continues to

spread in many countries,<sup>4–8</sup> and despite aggressive research, no proven therapies have been described.

Treatment strategies for critically ill COVID-19 patients are lacking, with only limited evidence available for a

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battery of antiviral, antibiotic, and anti-inflammatory agents and aggressive supportive therapy. Multiple clinical trials are ongoing, including the repurposing of remdesivir, an antiviral investigated to treat Ebola, and hydroxychloroquine, an antimalarial chloroquine derivative used to treat lupus and rheumatoid arthritis. There are early anti-COVID-19 efficacy data with remdesivir,<sup>9</sup> and preliminary data supporting the use of hydroxychloroquine, alone or in combination with azithromycin,<sup>10</sup> have since been shown by larger controlled trials as misleading and potentially dangerous.<sup>11</sup> New therapies are needed to improve outcomes for critically ill COVID-19 patients.

In convalescent plasma therapy, blood plasma from a recovered patient is collected and transfused to a symptomatic patient. The transfer of convalescent plasma is an old concept, having been used since at least 1918 when it was employed to fight the Spanish flu pandemic.<sup>12</sup> More recently, convalescent plasma was used with some reported success during the 2003 SARS pandemic,<sup>13,14</sup> the 2009 influenza H1N1 pandemic,<sup>15</sup> and the 2015 Ebola outbreak in Africa.<sup>16</sup> Several small observational studies published during the COVID-19 pandemic suggest convalescent plasma is part of an effective treatment strategy for patients with severe disease.<sup>17–20</sup> The first report describing administration of convalescent plasma to five patients early in the COVID-19 outbreak in Wuhan was recently published.<sup>18</sup> Five critically ill patients received two, same-day infusions from five recovered healthy donors. In four of the five patients, inflammatory biomarkers decreased and A/a gradient improved, and all patients had improvement in pulmonary lesions on the basis of computed tomography scan.<sup>18</sup> A second study by Duan et al<sup>17</sup> reported improved clinical outcomes in 10 patients who received a single transfusion of convalescent plasma, with no adverse events reported. Two additional small case studies of five and six patients have since been published with similar findings.<sup>19,20</sup> A more recent study by Zeng et al<sup>21</sup> suggested that administration of convalescent plasma late in the disease course was ineffective for mortality reduction.

We performed the present study to provide additional data on these initial clinical observations of patients' clinical course and subsequent improvement after receiving convalescent plasma therapy for COVID-19. We transfused 25 COVID-19 patients with severe and/or life-threatening disease at the Houston Methodist hospitals, a large, quaternary-care hospital system that serves metropolitan Houston, TX (approximately 7 million people; United States Census Bureau, <https://www.census.gov/newsroom/press-kits/2020/pop-estimates-county-metro.html>, last accessed May 3, 2020). Patients were transfused once with 300 mL of convalescent plasma. The therapy was well tolerated, and no transfusion-related adverse events were observed. At day 7 after transfusion, 9 of 25 patients (36%) had improvement in the assessed clinical end points. By 14 days after transfusion, 19 patients (76%) had improved or been discharged. Although our study has limitations, the data

indicate that transfusion of convalescent plasma is a safe treatment option for those with severe COVID-19 disease.

## Materials and Methods

This study was conducted at the Houston Methodist hospitals from March 28, 2020, through April 28, 2020, with the approval of the Houston Methodist Research Institute ethics review board and with informed patient or legally authorized representative consent. Patients were treated under either emergency investigational new drug or investigational new drug applications, approved by the US Food and Drug Administration (<https://www.fda.gov/medical-devices/emergency-situations-medical-devices/emergency-use-authorizations>). Approval to treat the first patient by emergency investigational new drug was granted on March 28, 2020. The investigational new drug application was approved on April 3, 2020.

## Patients

COVID-19 patients in the Houston Methodist hospitals were considered for enrollment in this trial. SARS-CoV-2 infection was confirmed by RT-PCR. Patients were eligible if they had severe and/or life-threatening COVID-19 disease (US Food and Drug Administration, <https://www.fda.gov/vaccines-blood-biologics/investigational-new-drug-ind-or-device-exemption-ide-process-cber/recommendations-investigational-covid-19-convalescent-plasma/#Patient%20Eligibility2020>, last accessed May 3, 2020). Severe disease was defined as one or more of the following: shortness of breath (dyspnea), respiratory rate  $\geq$  30/min, blood oxygen saturation  $\leq$  93%, partial pressure of arterial oxygen to fraction of inspired oxygen ratio  $<$  300, and/or pulmonary infiltrates  $>$  50% within 24 to 48 hours. Life-threatening disease was defined as one or more of the following: respiratory failure, septic shock, and/or multiple organ dysfunction or failure. Clinical data for patients were obtained from the hospital electronic medical record.

## Definition of Clinical Disease Severity

Clinical severity for the purposes of outcome assessment was scored on the basis of a modified six-point clinical scale used by the World Health Organization R&D Blueprint group ([https://www.who.int/blueprint/priority-diseases/key-action/COVID-19\\_Treatment\\_Trial\\_Design\\_Master\\_Protocol\\_synopsis\\_Final\\_18022020.pdf](https://www.who.int/blueprint/priority-diseases/key-action/COVID-19_Treatment_Trial_Design_Master_Protocol_synopsis_Final_18022020.pdf), last accessed May 6, 2020). Patients were assigned a clinical status at baseline (day 0, date of transfusion) and evaluated at days 0, 7, and 14. The six-point scale is as follows: 1, discharged (alive); 2, hospitalized, not requiring supplemental oxygen but requiring ongoing medical care (for COVID-19 or otherwise); 3, hospitalized, requiring low-flow supplemental oxygen; 4, hospitalized, on noninvasive ventilation or

high-flow oxygen devices; 5, hospitalized and on invasive mechanical ventilation or extracorporeal membrane oxygenation (ECMO); and 6, death.

### Convalescent Plasma Donors

Convalescent plasma was obtained by apheresis using the Trima Accel automated blood collection system (Terumo BCT, Lakewood, CO). Plasma (600 mL) was collected from each donor and divided into two 300-mL units. Each donor had a documented history of laboratory-confirmed SARS-CoV-2 infection on the basis of a positive RT-PCR test result. All plasma was donated by recovered and healthy COVID-19 patients who had been asymptomatic for  $\geq 14$  days. Donors were between 23 and 67 years old. All donors provided written informed consent and tested negative for SARS-CoV-2 by RT-PCR. If eligible according to standard blood donor criteria, donors were enrolled in a frequent plasmapheresis program. Donors were negative for anti-human leukocyte antigen antibodies, hepatitis B virus, hepatitis C virus, HIV, human T-lymphotropic virus I/II, Chagas disease, West Nile virus, Zika virus, and syphilis, per standard blood banking practices.

### RT-PCR Testing for SARS-CoV-2 Infection

Symptomatic patients with a high degree of clinical suspicion for COVID-19 disease were tested in the Molecular Diagnostics Laboratory at Houston Methodist Hospital using a validated assay applied for under Emergency Use Authorization from the US Food and Drug Administration. The assay follows the protocol published by the World Health Organization<sup>22</sup> and uses a 7500 Fast Dx instrument (Applied Biosystems, Foster City, CA) and 7500 SDS software (Applied Biosystems). Testing was performed on nasopharyngeal or oropharyngeal swabs immersed in universal transport media, bronchoalveolar lavage fluid, or sputum treated with dithiothreitol.

### SARS-CoV-2 Spike Protein Expression: RBD and ECD Domains

The expression and purification of the receptor binding domain (RBD) and ectodomain (ECD) of the SARS-CoV-2 spike protein have been described previously.<sup>23</sup> Briefly, the RBD (residues 319 to 591) and ECD (residues 306 to 577) domains were cloned into the mammalian expression vector p $\alpha$ H (pNCOV-1), which contains an HRV3C cleavage site upstream of TwinStrep and 8xHis purification tags. The ColE1 vector was transformed and maintained in *Escherichia coli* DH10B at 37°C using ampicillin selection at 100  $\mu$ g/mL. Plasmids from single colonies were recovered using a mini-prep kit (Qiagen, Germantown, MD) after growing cells overnight in Superior broth supplemented with 100  $\mu$ g/mL ampicillin.

Expi293F cells (Thermo Fisher, Waltham, MA) were passaged twice and seeded to a density of  $7.5 \times 10^7$  cells in 25.5 mL Expi293 Expression Medium ( $2.9 \times 10^6$  cells/mL in a 125-mL flask). For each 30-mL transfection, plasmid DNA (30  $\mu$ g) was added to Opti-MEM I Reduced Serum Medium to a total volume of 1.5 mL and gently mixed. ExpiFectamine 293 Reagent (81  $\mu$ L) was diluted in Opti-MEM I medium to a total volume of 1.5 mL. After gently mixing, it was incubated for 5 minutes at room temperature. After incubation, the diluted DNA was added to the diluted ExpiFectamine 293 Reagent to obtain a total volume of 3 mL and gently mixed. The mixture was incubated for 20 minutes at room temperature to allow the DNA–ExpiFectamine 293 Reagent complexes to form and then added to the Expi293F cells. After incubating cells for 20 hours, 150  $\mu$ L of ExpiFectamine 293 Transfection Enhancer 1 and 1.5 mL of ExpiFectamine 293 Transfection Enhancer 2 were added to each flask. Cells were harvested at 7 days.

### Protein Purification

Immobilized metal affinity chromatography purification columns were used with 1 mL bed volume for each Ni-NTA column. Each prepared column was used to purify proteins from 200 to 250 mL of filtered tissue culture media. Following filtration, filtered tissue culture medium was applied to a previously prepared and equilibrated Ni-NTA column. Each column was washed with 20 mL equilibration buffer (50 mmol/L phosphate buffer, pH 7.5, 300 mmol/L NaCl, and 20 mmol/L imidazole). The target protein was eluted with 5 mL elution buffer (50 mmol/L phosphate buffer, pH 7.5, 300 mmol/L NaCl, and 250 mmol/L imidazole). The eluate was applied to a spin concentrator with 100 kDa molecular weight cutoff to concentrate target protein before fast protein liquid chromatography purification and for buffer exchange into cold 1 $\times$  phosphate-buffered saline (PBS). Spin concentrators were centrifuged at  $3000 \times g$ , at 4°C for 15 minutes. Following buffer exchange, the eluate was concentrated to approximately 600  $\mu$ L. The concentrated eluate was further purified using size-exclusion chromatography with a 24-mL Superose 6 10/300 GL column (GE Healthcare, Chicago, IL). The 0.5-mL sample loop was injected with 1 mL each of the following: 0.1 mol/L NaOH, RNase-free water, and 1 $\times$  PBS. The buffer-exchanged eluate was applied to the fast protein liquid chromatography sample loop and run with a flow rate 0.25 mL/minute. Fractions were collected after 0.2 CV, and fractionation volumes were collected at 0.33 mL.

### SARS-CoV-2 ELISA

Costar 96-well assay plates (Corning, Corning, NY) were coated with either SARS-CoV-2 spike (S protein) ECD or SARS-CoV-2 spike RBD (50  $\mu$ L at 2  $\mu$ g/mL in PBS)

overnight at 4°C. Plates were blocked with 2% milk in PBS at room temperature for 2 hours and washed 3× with PBS with 0.1% Tween 20. Plasma or monoclonal antibody was serially diluted in 50 µL/well across the entire 96-well plate. Negative plasma control was included on each antigen plate. Monoclonal antibody CR3022 was used as a positive control. CR3022 is a neutralizing antibody originally cloned from a convalescent SARS patient that targets the RBD of SARS-CoV<sup>24</sup> and binds to the RBD of SARS-CoV-2 with a binding affinity of 6.3 nmol/L.<sup>25</sup> Binding was performed at room temperature for 1 hour. Plates were washed, and anti-human IgG Fab HRP (Sigma A0293; 1:5000; Sigma-Aldrich, St. Louis, MO) was added to the plate (50 µL) and incubated at room temperature for 30 minutes. Plates were washed 3× with PBS with 0.1% Tween 20, enzyme-linked immunosorbent assay (ELISA) substrate (1-step Ultra TMB; Thermo Fisher) was added, plates were developed for 1 minute for RBD and 5 minutes for spike ECD, and the reaction was stopped with 50 µL of H<sub>2</sub>SO<sub>4</sub>. Plates were read at 450-nm absorbance. Threefold serial dilutions from 50 to 4050 were analyzed. Titer was defined as the last dilution showing an OD greater than a multiplate negative control average plus six SDs.

### SARS-CoV-2 Genome Sequencing and Analysis

Libraries for whole viral genome sequencing were prepared according to version 1 ARTIC nCoV-2019 sequencing protocol (<https://artic.network/ncov-2019>). Long reads were generated with the LSK-109 sequencing kit, 24 native barcodes (NBD104 and NBD114 kits), and a GridION instrument (Oxford Nanopore, Oxford, UK). Short reads were generated with the NexeraXT kit and a MiSeq or NextSeq 550 instrument (Illumina, San Diego, CA). Whole genome alignments of consensus viral genome sequence generated from the ARTIC nCoV-2019 bioinformatics pipeline were trimmed to the start of orf1ab and the end of orf10 and used to generate a phylogenetic tree using RAxML version 8.2 (<https://cme.h-its.org/exelixis/web/software/raxml/index.html>) to determine viral clade. Trees were visualized and annotated with CLC Genomics Workbench version 20 (Qiagen).

## Results

### Overview of Patient Characteristics

Twenty-five patients with severe and/or life-threatening COVID-19 disease were enrolled in the study from March 28, 2020, to April 14, 2020. Patients ranged in age from 19 to 77 years [median, 51 years; interquartile range (IQR), 42.5 to 60 years], and 14 were female (Table 1). The median body mass index was 30.4 kg/m<sup>2</sup> (IQR, 26.5 to 37 kg/m<sup>2</sup>), and most (22/25, 88%) had no smoking history. Many patients (16/25, 64%) had one or more underlying chronic conditions, including diabetes mellitus (10 patients),

hypertension (9 patients), hyperlipidemia (5 patients), and gastrointestinal reflux disease (4 patients). Most patients (19/25, 76%) enrolled in the study had O-positive blood type. Bacterial or viral co-infections were identified in five patients (Table 1).

### Donor Characteristics

The characteristics of the donors of convalescent plasma are shown in Table 2. A total of nine donors provided plasma that was used to transfuse COVID-19 patients; two donors gave plasma on multiple occasions. The donors ranged in age from 23 to 67 years, and 56% (5/9) were males. On average, the donors gave plasma 26 days (range, 19 to 33 days) after their symptom start date and 21 days (range, 13 to 27 days) after their initial positive RT-PCR specimen collection date. Although all donors had been symptomatic, only one was ill enough to require hospitalization. To assess antibody titers, we used two ELISAs, one based on recombinant purified ECD of the spike protein and the second using recombinant RBD of the spike protein. The titers of the convalescent plasma used for transfusion ranged from 0 to 1350 for the RBD and ECD domains (Figure 1 and Supplemental Table S1).

### Transfusion of Severe COVID-19 Patients with Convalescent-Phase Donor Plasma

The median time from symptom onset to hospitalization was 6 days (IQR, 4 to 8 days) (Table 3). Most patients received concomitant anti-inflammatory treatments within 5 days of the plasma transfusion, including tocilizumab and steroids. Most received other investigational treatments, including courses of hydroxychloroquine and azithromycin, ribavirin, and/or lopinavir/ritonavir; and two patients received remdesivir (Table 3). All patients required oxygen support before transfusion (Figure 1), including 12 patients on mechanical ventilation, 10 on low-flow oxygen, and 3 on high-flow oxygen. One patient (Patient 9) was placed on ECMO on the day of transfusion before transfusion. More than half (13/25, 52%) had acute respiratory distress syndrome<sup>26</sup> at the time of transfusion (Table 3). The median time from symptom onset to transfusion was 10 days (IQR, 7.5 to 12.5 days), and the median time from hospitalization to transfusion was 2 days (IQR, 2 to 4 days) (Table 3). All patients received one 300-mL dose of convalescent-phase plasma, and one patient received a second transfusion 6 days after the initial transfusion. Clinical outcomes and laboratory parameters were assessed at days 0, 7, and 14 after transfusion.

### Outcomes

The primary clinical end point of the study was safety. No adverse events attributed to plasma transfusion occurred within 24 hours after transfusion. One patient developed a

**Table 1** Demographics and Clinical Characteristics of Patients with COVID-19 Disease Who Received Convalescent Plasma

Patient	Sex	Age, years	Weight, kg	BMI, kg/m <sup>2</sup>	Smoking history	Blood type	Co-infections	Co-existing chronic diseases
1	F	39	90	34	Never	O pos	None	DM2
2	F	63	104	38	Never	O pos	None	DM2, HTN, HLP, GERD
3	F	48	63	23	Never	O pos	None	None
4	M	57	96	29	Never	O pos	None	None
5	F	38	99	35	Never	O pos	Influenza B	DM2, HTN, GERD
6	M	46	133	32	Former	O pos	MSSA PNA	DM2
7	M	51	94	32	Former	A pos	None	DM2
8	M	74	84	27	Never	A pos	VAP: MSSA and GAS	DM2, HTN, CKD
9	F	55	73	26	Never	O pos	None	None
10	F	19	113	49	Never	O pos	<i>Enterococcus</i> BSI	None
11	F	22	91	40	Never	O pos	None	Asthma
12	F	46	65.8	24.9	Never	O pos	None	None
13	M	61	88	30	Unknown	O pos	None	None
14	F	49	101	31.9	Never	O pos	None	GERD, HTN
15	M	29	126	44	Never	O pos	None	None
16	F	30	94.7	38.2	Never	O pos	None	Post-partum, hypothyroidism
17	F	54	79	30	Never	O pos	None	HTN
18	M	56	102	40	Never	O pos	None	HTN, HLP
19	M	60	81.6	32	Never	O pos	None	DM2, HLD
20	F	77	95	36	Never	O pos	None	HTN, DM2
21	F	60	65	23	Never	O neg	None	None
22	F	77	86.5	29.8	Never	A pos	GAS	Atrial fibrillation, DM2, HLD
23	M	60	85	30.4	Never	O pos	None	DM2, HLD, HTN
24	M	54	72	25	Never	B pos	None	HLD
25	M	50	58	22.6	Never	B pos	None	None

F, female; M, male; BMI, body mass index; BSI, bloodstream infection; CKD, chronic kidney disease; COVID-19, coronavirus disease 2019; DM2, diabetes mellitus type 2; GAS, group A *Streptococcus*; GERD, gastrointestinal reflux disease; HLD, hyperlipidemia; HTN, hypertension; MSSA, methicillin-susceptible *Staphylococcus aureus*; neg, negative; None, no infection identified; PNA, pneumonia; pos, positive; VAP, ventilator-associated pneumonia.

morbilloform rash 1 day after transfusion that lasted for several days. Punch biopsy findings were compatible with an exanthematous drug eruption, and classic histologic findings of serum sickness (leukocytoclastic vasculitis) were not seen. Two patients developed deep vein thrombosis 4 and 8 days after transfusion, and one patient developed a deep vein thrombosis and a pulmonary embolism 4 days after transfusion. The observed thrombotic complications are consistent with findings reported for COVID-19 patients.<sup>27</sup> The secondary end point was an improvement in the modified six-point World Health Organization ordinal scale at day 14 after transfusion, including discharge from the hospital (Supplemental Table S2). At day 7 after transfusion, 9 patients (36%) improved from baseline, 13 (52%) had no change, and 3 deteriorated (Figure 2). Seven of the nine improved patients (28%) had been discharged. By day 14 after transfusion, 19 patients (76%) improved from baseline: an additional four patients were discharged, eight patients improved from baseline, three patients remained unchanged, three had deteriorated, and one patient died from a condition not caused by plasma transfusion (Figure 2 and Supplemental Table S2). The average overall length of hospital stay was 14.3 days (range, 2 to 25 days). The average post-transfusion length of hospital stay was 11 days (range, 1 to 21 days) (Table 3).

## Laboratory Results

Laboratory results were assessed for parameters associated with inflammation and liver function. The median value for C-reactive protein decreased in our cohort from 14.66 mg/dL at day 0 to 2.9 and 0.45 mg/dL at days 7 and 14 after transfusion, respectively (Table 4). There was a trend toward increasing ferritin by day 3, which tended to decrease by day 7. No significant increases in liver enzymes were noted (Table 4 and Supplemental Table S3).

## Viral Genome Sequencing of SARS-CoV-2 Strains from Recipients and Donors

A recent analysis of the genomic heterogeneity of the SARS-CoV-2 virus strains circulating in Houston early in the pandemic showed that the predominant clades isolated were A2a, B, and B1.<sup>28</sup> Amino acid polymorphisms, especially in the spike protein, can potentially alter the character of the antibody response and virulence profile of the virus.<sup>23,29–31</sup> Therefore, we sequenced the genomes of the SARS-CoV-2 virus strains infecting donors and recipients to assess the magnitude of nucleotide and amino acid mismatch between the viral genotype of donors and plasma recipients. Of the 34 patients and donors, we were able to

**Table 2** Characteristics of Convalescent Plasma Donors

Donor	Age, years	Sex	Blood type	Symptom start date	Positive test date	Hospitalized	Symptoms resolved	Plasma collected date(s)	Symptom resolution to first donation
1	44	M	O pos	3/7/20	3/14/20	No	3/10/20	3/27/20, 3/31/20, 4/3/20, 4/7/20	17
2	36	M	O pos	3/6/20	3/12/20	No	3/13/20	3/31/20, 4/3/20, 4/8/20	19
3	67	F	A pos	3/6/20	3/17/20	No	3/17/20	4/3/20	17
4	23	F	O pos	3/11/20	3/18/20	No	3/24/20	4/9/20	16
5	50	M	O pos	3/13/20	3/14/20	No	3/27/20	4/10/20	14
6	41	F	O pos	3/21/20	3/23/20	No	3/24/20	4/9/20	16
7	54	F	A pos	3/18/20	3/20/20	No	3/19/20	4/7/20	19
8	61	M	A pos	3/8/20	3/16/20	Yes	3/22/20	4/10/20	19
9	23	M	B pos	3/13/20	3/17/20	No	3/25/20	4/13/20	19

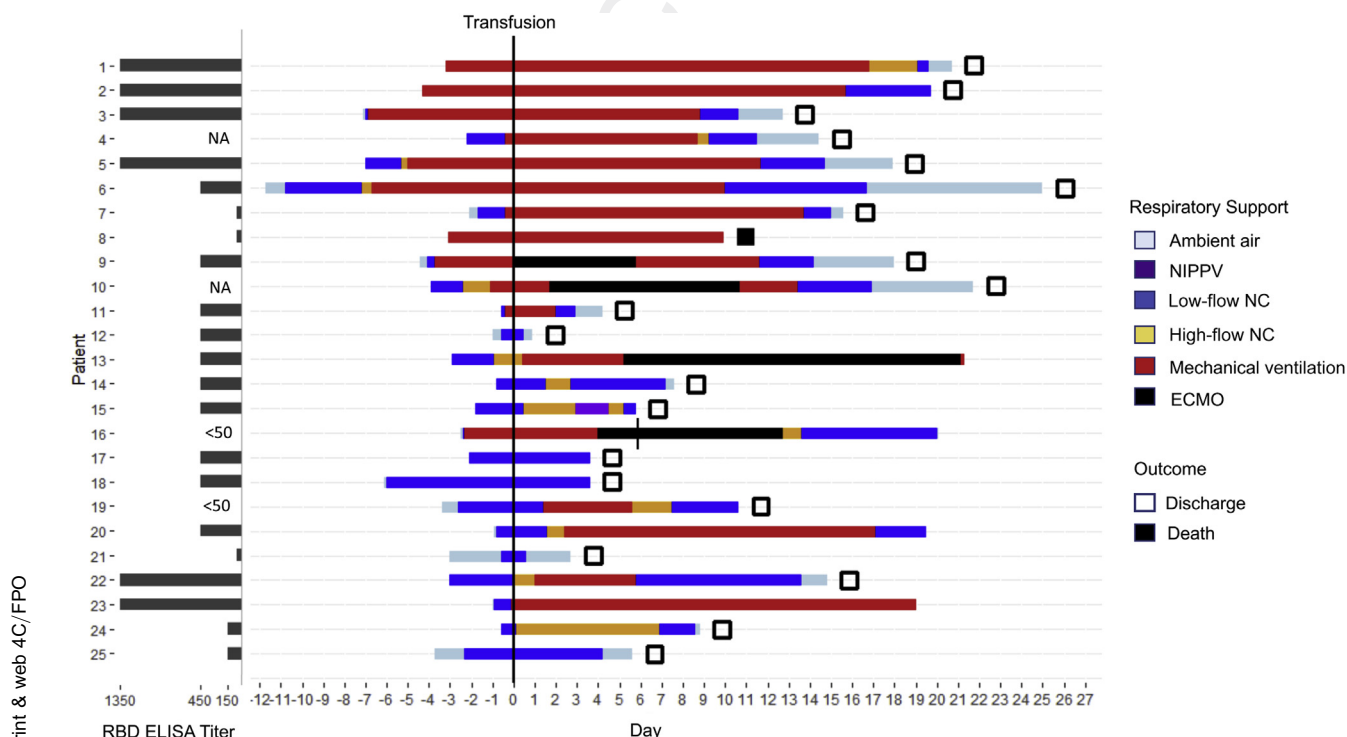
F, female; M, male; pos, positive.

analyze all plasma recipient genotypes and four donor genotypes. Overall, there were few polymorphisms in the sequenced viruses, and there was no correlation between infecting strains and disease severity (Supplemental Figure S1). Analysis of the first four donors found that, in general, donor and recipient S proteins matched when their SARS-CoV-2 isolates were from the same clade (Supplemental Figure S1). This is primarily a result of a D614G amino acid change in S protein that defines the clade A2a.<sup>28,32</sup> However, there are at least three instances of an

additional amino acid change in the S2 domain of the S protein,<sup>23,30,31</sup> one in a donor (M731I) and two in recipients (S967R and L1203F) (Supplemental Figure S1).

## Discussion

Our study was performed to evaluate the safety and potential benefit of transfusing convalescent plasma to patients with severe COVID-19 disease. To date, this is the largest



**Figure 1** Respiratory support status, clinical score, patient outcomes (discharge/death), and receptor binding domain (RBD) titer of transfused plasma in 25-patient cohort. Respiratory support requirements for the duration of hospitalization are color coded per the key. Discharge or death is indicated by open or closed squares, respectively. Patient 16 was given a second transfusion on day 6, indicated by a vertical line. The convalescent plasma titers for the RBD domain of the severe acute respiratory syndrome coronavirus 2 spike protein are indicated to the left. ECMO, extracorporeal membrane oxygenation; ELISA, enzyme-linked immunosorbent assay; NC, nasal cannula; NIPPV, noninvasive, positive-pressure ventilation.

**Table 3** Disease Course and Additional Treatments of Patients Receiving Convalescent Plasma

Patient No.	Symptom onset to admission, days	Symptom onset to positive SARS test, days	Admission to transfusion, days	Complications before transfusion	Anti-inflammatory treatments	Antiviral treatments	Length of hospital stay, days	Post-transfusion length of hospital stay, days
1	7	3	1	ARDS	Tocilizumab	HCQ, RBV	24	21
2	7	9	4	ARDS, CRRT	Interferon, steroids	HCQ, AZM, RBV	24	20
3	8	3	6	ARDS	Tocilizumab, steroids	HCQ, RBV, LPVr	20	13
4	8	9	2	ARDS	Tocilizumab, steroids	HCQ, AZM, RBV	17	15
5	3	4	7	ARDS	None	HCQ, AZM, RBV	25	18
6	3	4	13	ARDS	Tocilizumab	HCQ, AZM, RBV	37	NA
7	3	3	2	ARDS	Tocilizumab	HCQ, LPVr	20	16
8	4	5	3	ARDS	Steroids	HCQ, RBV, LPVr	13	10
9	4	4	4	ARDS, CRRT, ECMO (VV)	Tocilizumab, steroids	HCQ, RBV	22	18
10	6	10	5	ARDS	Tocilizumab	HCQ, AZM, RBV, LPVr, remdesivir	28	22
11	5	3	1	ARDS	Steroids	HCQ, AZM, RBV	5	4
12	10	6	2	None	None	HCQ, AZM	2	1
13	5	6	3	None	Tocilizumab	HCQ, AZM, RBV	NA	NA
14	12	6	1	None	Tocilizumab, steroids	HCQ, AZM, RBV	9	8
15	7	8	2	None	None	HCQ, AZM, RBV	8	6
16	8	3	2	ARDS	Tocilizumab, steroids	HCQ, AZM, RBV	NA	NA
17	4	4	2	None	None	HCQ, AZM	6	4
18	8	8	6	None	None	HCQ, AZM	10	4
19	6	6	3	None	Tocilizumab	HCQ, AZM	14	11
20	3	4	1	None	None	HCQ, RBV, AZM, remdesivir	NA	NA
21	8	8	3	None	None	HCQ, RBV	6	3
22	4	4	2	None	Steroids	HCQ, AZM, RBV	18	15
23	14	1	2	ARDS	Tocilizumab, steroids	HCQ, AZM	NA	NA
24	9	6	2	None	Tocilizumab	HCQ, AZM	10	9
25	11	11	3	None	Tocilizumab	HCQ, AZM	9	6

ARDS, acute respiratory distress syndrome; AZM, azithromycin; CRRT, cardiac rapid response team; ECMO (VV), extracorporeal mechanical oxygenation (venovenous); HCQ, hydroxychloroquine; LPVr, lopinavir/ritonavir; NA, still hospitalized at day 14 after transfusion (study end point); RBV, ribavirin; SARS, severe acute respiratory syndrome.

cohort assessed for outcomes pertaining to convalescent plasma transfusion for COVID-19. Of our 25 patients, 9 had improvement by day 7, and an additional 12 patients (for a total of 19) had improvement by day 14, as assessed by discharge or at least a one-point improvement on a modified clinical scale. Several case studies investigating the use of convalescent plasma to treat severe COVID-19 have recently been published,<sup>17–21</sup> and the overall findings presented herein are consistent with these reports.

Convalescent plasma therapy has been administered on the frontlines during emergencies, and we and others recognize the need for controlled clinical trials to determine its therapeutic efficacy.<sup>13,14,18,33,34</sup> The timing of the transfusion after symptom onset, the number of transfusions, the volume and its adjustment on the basis of body mass index, donor antibody titers, and other parameters need to be evaluated to optimize this therapy. For example, some studies have observed that the sooner after the onset of symptoms that the transfusion was administered, the better

the outcomes.<sup>13,14,34</sup> Variability existed among our cohort with respect to symptom onset and severity of illness.

The anti-SARS-CoV-2 anti-spike protein IgG titers varied significantly among individual donors, as assessed by ELISA (Supplemental Table S1). Early in the study period, ELISA titers were not available, and thus, transfusions were given solely on the basis of ABO compatibility. Among the five patients who received plasma from a donor with an anti-RBD IgG titer of  $\leq 50$ , one is deceased, and one was placed on ECMO. The patient placed on ECMO received a second dose of convalescent plasma confirmed to have high IgG titer before transfusion. The patient was eventually extubated and weaned off ECMO. Regardless, at this time, no clear correlation between ELISA IgG titer and patient outcomes can be established in this small patient cohort. In addition, more studies are needed to better understand why donors present with a range of anti-spike antibody titers, and whether there is a correlation between donor disease presentation and antibody titers. Additional studies are

Baseline Oxygen Support, n (%)	Day 7						Day 14					
	Death	Inv.	High-flow	Low-flow	Room air	Discharged	Death	Inv.	High-flow	Low-flow	Room air	Discharged
Invasive n = 13	0	12 (92%)	0	0	0	1 (8%)	1 (8%)	3 (23%)	0	6 (46%)	1 (8%)	2 (15%)
High-flow n = 3	0	1 (33%)	0	2 (66%)	0	0	0	1 (33%)	0	0	1 (33%)	1 (33%)
Low-flow n = 9	0	1 (11%)	1 (11%)	1 (11%)	0	6 (67%)	0	1 (11%)	0	0	0	8 (89%)
Room air n = 0	0	0	0	0	0	0	0	0	0	0	0	0

Worse from baseline
  No change from baseline
  Improved from baseline

**Figure 2** Clinical outcomes at days 7 and 14 after transfusion. Distribution of patients on low-flow, high-flow, invasive, or no oxygen support at days 0 (day of transfusion), 7, and 14. By day 7 after transfusion, 36% (9/25) of patients had improved from baseline; 76% (19/25) of patients improved by day 14 after transfusion. Inv., Invasive.

underway to better understand the correlation between anti-SARS-CoV-2 antibody titers and virus neutralization.

The results from our study support the existing data from the COVID-19 literature that point to underlying medical conditions, such as obesity, type 2 diabetes, and hypertension, playing a large role in patients' COVID-19 disease course and outcomes.<sup>35-37</sup> Of transfused patients in this study, 68% (17/25) had a body mass index in the obese category and 84% were considered overweight.

A confounding variable in many convalescent plasma studies is the addition of other treatment regimens, such as antivirals and anti-inflammatory compounds. Adjunct therapies hinder the ability to draw definitive conclusions regarding the contribution of the convalescent plasma. In our study, all 25 patients received hydroxychloroquine and azithromycin, as these were reported to have beneficial effects early in the pandemic.<sup>10</sup> Subsequent larger and more controlled studies determined that this combination has no benefits to patients and, in fact, could be harmful.<sup>11</sup> Many (68%) of our patients were also administered oral ribavirin. Despite inconclusive data on ribavirin's efficacy in the treatment of SARS during the 2003 epidemic,<sup>38</sup> proven safety and ready availability supported its use in the treatment of our COVID-19 patients. Two patients received remdesivir, which was recently shown to modestly reduce recovery time (NIH, <https://www.niaid.nih.gov/news-events/nih-clinical-trial-shows-remdesivir-accelerates-recovery-advanced-covid-19>, last accessed May 5, 2020).<sup>9</sup> Anti-inflammatory compounds, such as the IL-6 inhibitor tocilizumab and methylprednisolone, were administered per institutional protocols within 5 days of the plasma transfusion to 72% of our cohort. Tocilizumab was recently shown to reduce mortality in a retrospective analysis of 20

severe COVID-19 patients.<sup>39</sup> Because convalescent plasma therapy is typically performed in emergency situations for the ill, it is difficult to assess its benefits as a stand-alone treatment. A blinded, randomized controlled trial is currently being considered.

The patient outcomes in our study are similar to those recently published describing treating COVID-19 patients with remdesivir on a compassionate-use basis.<sup>9</sup> In that review, patients were prescribed a 10-day course of remdesivir, with follow-up for 28 days or until discharge or death. Both study cohorts included patients who required invasive ventilation, including 35 of 53 (66%) of remdesivir patients compared with 17 of 25 (68%) of the patients in our study. Clinical improvement was less frequent among patients who received invasive ventilation at any time or were aged  $\geq 70$  years. In the remdesivir study, 36 of 53 patients (68%) showed clinical improvement at follow-up (median time to follow-up, 18 days), whereas 19 of 25 patients (76%) receiving convalescent plasma improved by day 14 after transfusion. These data suggest that treatment with convalescent plasma and remdesivir resulted in similar outcomes among patients on the basis of oxygenation requirements and age. The mortality difference between the cohorts cannot be compared as the remdesivir cohort represented an older population (median age, 64 years, versus 51 years in our study), where the risk of death was greater at baseline. Delays in obtaining remdesivir on a compassionate-use basis (12 days from symptom onset) may have artificially extended the cohort's opportunity to demonstrate clinical improvement and does not reflect the eligibility criteria for any ongoing clinical trials (World Health Organization, <https://www.who.int/emergencies/diseases/novel-coronavirus-2019/global-research-on-novel-coronavirus-2019-ncov/>



**Table 4** Median Laboratory Values of Plasma Recipients at Days 0, 7, and 14 after Transfusion

Laboratory test (normal range)	Median values		
	Day 0	Day 7	Day 14
CRP (0–0.5 mg/dL)	14.66	2.9	0.45
WBC count (4.5–11 k/ $\mu$ L)	10.9	11.3	13.1
LDH (87–225 U/L)	380	394	305
ALT (5–50 U/L)	38	60.5	47
AST (10–35 U/L)	51	41	32
Ferritin (13–150 ng/mL)	878	1633.5	718
Total bilirubin (0–1.2 mg/dL)	0.4	0.75	0.9

ALT, alanine aminotransferase; AST, aspartate aminotransferase; CRP, C-reactive protein; LDH, lactate dehydrogenase; WBC, white blood cell.

*solidarity-clinical-trial-for-covid-19-treatments*, last accessed May 5, 2020; NIH, <https://clinicaltrials.gov/ct2/show/NCT04292899>, last accessed May 5, 2020; NIH, <https://clinicaltrials.gov/ct2/show/NCT04280705>, last accessed May 5, 2020). Clinical outcomes data to inform timing of therapeutic interventions, like remdesivir or convalescent plasma, are lacking.

We analyzed the genomes of the infecting SARS-CoV-2 strain from both the donors and recipients. One could conceive of a situation in which the donor genotype of the SARS-CoV-2 infecting strain was matched with the genotype of the patient's strain to maximize potential immune benefit. We found few differences in the inferred amino acid sequences of the plasma donor and recipient strains and no association between disease severity and infecting strain genotype.

Most of the donors and plasma recipients in our study had type O blood (25/34, 74%). Our initial donors, who donated repeatedly, were blood type O. Because ABO compatibility was a requirement for recipient selection early in the study, many of our early recipients were also type O. Zhao et al<sup>40</sup> have reported that of the 2173 patients analyzed in their study of COVID-19 patients in China, most had type A blood. More studies are needed to determine if this association holds true in geographically distinct areas of infection. Regardless, our data do not reflect a higher rate of blood type A in COVID-19 patients.

## Limitations

As with the great majority of the studies using convalescent plasma to treat severe infections, our study has several important limitations. First, the study was a small case series and no control group was included. Thus, it is not clear if the 25 patients given convalescent plasma would have improved without this treatment. Second, all patients were treated with multiple other medications, including antiviral and anti-inflammatory agents. Thus, we cannot conclude that the patient outcomes were due solely to administration of convalescent plasma. Third, 24 of the 25 patients

received only one transfusion of plasma. Whether treatment with multiple transfusions on  $\geq 1$  day would be a more effective regimen is not clear. An expanded donor pool providing higher-titer convalescent plasma would allow for dose escalation studies. Fourth, many patients had severe COVID-19 disease. It is possible that transfusion of convalescent plasma earlier in the course of disease or in patients with less severe symptoms would be a better approach. Fifth, our plasma donors had a range of anti-S protein IgG titers. Several patients were transfused with plasma with low titer of anti-S protein antibody. Sixth, the small number of patients treated, coupled with the experimental design, did not permit us to determine if this therapy significantly reduces mortality or other measures of disease outcome. Finally, although this study assessed outcomes at days 7 and 14 after transfusion, at the time of this writing, all but two of the surviving patients who were intubated had been extubated. Similarly, all patients that were on ECMO had been weaned, and 20 of the 25 patients had been discharged.

## Conclusion

Outcomes from this case series of 25 patients indicate that administration of convalescent plasma is a safe treatment option for those with severe COVID-19 disease.

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## Supplemental Data

Supplemental material for this article can be found at <http://doi.org/10.1016/j.ajpath.2020.05.014>.

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