

1 **Immune responses and residual SARS-CoV-2 in two critically ill**  
2 **COVID-19 patients before and after lung transplantation**

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33 To the Editor:

34 A small number of COVID-19 patients develop critical illness resulting in acute lung  
35 injury (ALI) and acute respiratory distress syndrome (ARDS). Although lung  
36 transplantation (LT) can be used to rescue patients from COVID-19-related ARDS<sup>1,2</sup>,  
37 current information concerning the immune statuses and pathogenic conditions of such  
38 transplant patients is lacking. In this study, we assessed the immune responses and the  
39 residual SARS-CoV-2 nucleic acids in patients before and after LT, including the first  
40 COVID-19 lung transplant patient in the world.

41 There were two critically ill COVID-19 patients in Wuxi of China, from January 25 to  
42 March 31, 2020, who eventually recovered after LT (Ethics No. 2020-014)<sup>2 3</sup>. For  
43 comparison purposes, we analyzed the whole blood lymphocytes, immunocyte  
44 subclasses (T, B and NK cells), blood cytokines and Ag-specific IgM and IgG of  
45 hospitalized COVID-19 patients in Wuxi (Ethics No. 2020-010-1). We focused on the  
46 analyses of the two critically ill COVID-19 patients (Patient 1: a 58-year-old male had  
47 COVID-19-associated ALI and ARDS; Patient 2: a 73-year-old male had COVID-19-  
48 associated multiple organ failure and ARDS).

49 In view of the limited number of critically ill patients, we merged severe and critical  
50 illness, designated as severe illness (Fig 1). Most of the severely ill patients had low  
51 levels of blood lymphocytes during hospitalization, and in particular, the blood  
52 lymphocytes in the two critically ill patients remained below the normal value before  
53 and after LT ( $<1.1 \times 10^9/L$ ) (Fig. 1A). There were statistically significant lower levels of

54 blood CD3<sup>+</sup> CD45<sup>+</sup> T (<60%) (Fig. 1B and 1C), CD4<sup>+</sup> T, CD8<sup>+</sup> T and NK cells in  
55 individuals with critical clinical manifestations ( $P < 0.05$ ), and a reduction of CD8<sup>+</sup> T  
56 cells was the most statistically significant in the severely ill patients ( $P < 0.01$ ) (Fig. 1B  
57 and 1D). Compared with untreated mildly ill patients, no significant increase of T and  
58 NK cells was observed in blood before and after LT, and only B cells increased slightly  
59 in the two critically ill patients, likely owing to mesenchymal stem cell infusion therapy  
60 (Fig. 1B and 1D). The two critically ill patients developed mildly positive SARS-CoV-  
61 2-specific IgM and IgG before LT, and such humoral immune responses became  
62 negative post LT (data not shown), likely due to blood transfusion therapy in the  
63 absence of new pathogen stimulation. It was reported that IL-6 and IL-10 play distinct  
64 roles in immune tolerance<sup>4,5</sup>. In our study of the two critically ill patients, long-lasting  
65 IL-6 and IL-10 levels in plasma exceeded the upper limits of normal values,  
66 accompanied by viral replication. The concentrations of proinflammatory cytokines  
67 (IL-6, IFN- $\gamma$ , and TNF- $\alpha$ ), anti-inflammatory cytokine IL-10 and B-/T-cell stimulating  
68 factor IL-4 in the severe period were significantly higher than those in the recovery  
69 period ( $p < 0.05$ ), especially for the critically ill patients post LT (Fig. 1E). The above  
70 findings together indicate that SARS-CoV-2 infection in critically ill patient results in  
71 lower levels of cellular and humoral immune responses.

72 Pathological analyses were performed by immunostaining for CD3<sup>+</sup> T, IgA<sup>+</sup> and SARS-  
73 CoV-2 S protein<sup>+</sup> cells in the diseased lungs. In immunohistochemistry, critically ill  
74 patient' lungs (Patient 2) showed obscure mature CD3<sup>+</sup> cells in tissues, and extensively  
75 fibrosis (Fig. 1F), interstitial hemorrhage (Fig. 1G) and mucous exudative necrosis in

76 the bronchioles (Fig. 1H), as well as alveolar epithelial atrophy, hyperplasia and  
77 shedding in the alveolar cavity (Fig. 1I). The number of IgA<sup>+</sup> cells from alveoli  
78 epithelial cells decreased in both the right and the left pulmonary lobes (Fig. 1J-1M).  
79 Residual SARS-CoV-2 in the lungs has been suggested to be the main reason for viral  
80 positivity of discharged COVID-19 patients <sup>6, 7</sup>. We observed a direct evidence of  
81 residual SARS-CoV-2 in excised lungs (Fig. 1N-1Q), suggesting that antiviral therapy  
82 may not completely eliminate the virus in the dysfunctional lungs. For the two critically  
83 ill COVID-19 patients under therapies, convalescent plasma and mesenchymal stem  
84 cell infusions appeared unable to restore a systemic immunity, including cellular and  
85 mucosal (IgA) immune responses in the diseased lungs. A previous pathologic study  
86 showed that SARS-CoV-2 was highly destructive to the immune system, resulting in  
87 reduced splenic T and B cell compositions due to necrosis and apoptosis <sup>8</sup>. This may  
88 account for the long-term low systemic immunity of COVID-19 transplant patients.

89 SARS-CoV-2 RNA can be detected in the gastrointestinal tract using swabs and stool  
90 sampling <sup>9</sup>, and in particular, SARS-CoV-2 particles can be found in the gut  
91 endothelium <sup>10</sup>, suggesting the potential significance of the gut in viral transmission and  
92 pathogenesis. Prior to LT, the two critically ill patients in the current study were under  
93 treatments with convalescent plasma infusion, mesenchymal stem cell infusion and  
94 antiviral agents until SARS-CoV-2 nucleic acid turned into negative in blood and  
95 nasopharyngeal and anal swabs. Post LT, residual SARS-CoV-2 in nasopharyngeal and  
96 anal swabs was also examined. In anal swabs, SARS-CoV-2 was mildly positive at day  
97 26, 30, and 43 post LT in Patient 1 (Fig. 2A), and mildly positive at day 28 post

98 hospitalization and day 17 post LT in Patient 2 (Fig. 2B). Of note, in the absence of  
99 antiviral treatment and under immune suppression therapy (doses of drugs were only  
100 1/6 of those for ordinary transplant patients) post LT, there was no indication of SARS-  
101 CoV-2 infection in the new (donor) lungs according to the chest radiographs and axial  
102 pulmonary CT graphs in the negative-pressure ward up to 45 days (Fig. 2A and 2B),  
103 while viral nucleic acid remained negative after the patients were transferred to the  
104 general ward. There was no medical staff infected by SARS-CoV-2 during medical care  
105 in general ward. The above results indicated that the detection of viral positive nucleic  
106 acids by anal swabbing does not necessarily reflect a contagious SARS-CoV-2 in the  
107 gut.

108 In conclusion, following LT, the two critically ill COVID-19 patients in the absence of  
109 antiviral treatment have not had a second SARS-CoV-2 infection in the new lungs. For  
110 the first time, our study provides information relating to the immune status and SARS-  
111 CoV-2 positivity of lung transplant patients.

112 **References**

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136 **Declaration of Competing Interest**

137 The authors declare that the research was conducted in the absence of any commercial  
138 or financial relationships that could be construed as a potential conflict of interest.

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149 **Figure legends**

150 **Figure 1. Immune responses of critically ill COVID-19 patients and**  
151 **immunohistochemistry of the diseased lungs.**

152 **A.** Blood T lymphocyte counts. Blood T lymphocyte counts were detected by clinical  
153 blood cell analyzer. The gray area is the location of the dangerous values ( $<1.1 \times 10^9/L$ ).

154 **B.** Gating strategies for blood immunocytes.  $CD3^+$  and  $CD45^+$  T lymphocytes were  
155 gated for  $CD4^+$  and  $CD8^+$  T cells.  $CD3^-$  and  $CD45^+$  immunocytes were gated for B  
156 ( $CD19^+$ ) and NK ( $CD16/56^+$ ) cells. **C.** Percentage comparison of blood  $CD3^+$  and

157  $CD45^+$  lymphocyte levels. **D.** Absolute percentage comparison of blood immunocytes.

158 **E.** Plasma cytokine levels. Plasma samples from COVID-19 patients (n=6) were  
159 collected during the early and recovery periods (mildly and severely ill patients were at  
160 around day 30; transplant patients were before being transferred to the general ward).

161 IL-6, IL-10, IFN- $\gamma$ , IL-17A, TNF- $\alpha$ , and IL-4 beads were used for staining the cytokines  
162 in plasma samples. Quantitative detection and comparison of inflammatory cytokine  
163 expression levels in different periods were conducted by flow cytometry. Mildly ill

164 patients (Mild): individuals who had mild manifestation and no apparent or long-term  
165 decrease in lymphocyte levels. Severely ill patients (Severe): individuals who had  
166 lasting lower levels of lymphocytes and needed supplemental oxygen and intensive care.

167 Critically ill patients: individuals who had failure of respiratory organs leading to  
168 dependency on ventilators. The red solid symbol ( $\blacktriangle$  or  $\bullet$ ) stands for critically ill  
169 patients. The results are expressed as the mean  $\pm$  SEM; NS, not statistically significant,

170  $P < 0.05$  was considered statistically significant. **F-I.** Hominine CD3 immunostaining  
171 of diseased lung tissues. **J-M.** Hominine IgA immunostaining. **N-Q.** SARS-CoV-2 S  
172 protein immunostaining. Post LT, approximately 2-cm segments from excised human  
173 lungs were collected and fixed in formalin for 24 h. After fixation, tissue was embedded  
174 and sectioned. For the detection of CD3-, IgA- and SARS-CoV-2 S protein-positive  
175 cells in lungs, the slides were stained with CD3 $\epsilon$  rabbit monoclonal antibody (Cell  
176 Signaling Technology, U.S.A, 1: 1000, E4T1B), human IgA heavy chain rabbit  
177 polyclonal Ab (Proteintech, U.S.A., 1: 400), and SARS-CoV-2 S protein rabbit  
178 polyclonal Ab (Sino Biological Inc., China, 1: 50), respectively, overnight at 4°C. After  
179 labeling with a goat anti-rabbit secondary IgG Ab, a color reaction was developed with  
180 the addition of 3, 3'-diaminobenzidine free base (DBA), followed by counterstaining  
181 with hematoxylin. Data shown are representative immunohistochemistry results. S  
182 protein: spike protein; R, right upper lobe; L, left lower lobe; The red arrow shows the  
183 positive cells.

184 **Figure 2. Chest imaging and viral nucleic acid detection before and after LT.**

185 **A.** Chest radiographic images and SARS-CoV-2 nucleic acid results of patient 1. Chest  
186 radiographic images were obtained before and post LT. Axial CT images on day 45 post  
187 LT, and SARS-CoV-2 nucleic acid results. **B.** Axial chest CT images before and after  
188 LT and SARS-CoV-2 nucleic acid results of patient 2. Nasopharyngeal and anal swabs  
189 were placed in viral transport medium at a low temperature. Total RNA extraction and  
190 SARS-CoV-2 detection were performed using the commodity reagents of Da An Gene

191 (Da An Gene, China) according to the manufacturer's instructions. Briefly, SARS-CoV-  
192 2 RNA was detected by real-time RT-PCR. Target genes (ORF1ab + N) were set as  
193 described in the reagent instructions. A cycle threshold value (Ct value) less than 36  
194 was defined as a positive (+), and Ct value between 36 and 37 was defined as mildly  
195 positive ( $\pm$ ). 1, 2 and 3 on CT images mean the different axial images within the lungs.  
196 ph: post-hospitalization; plt: post LT.



