Clinical and immunological characteristics in COVID-19 convalescent patients

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Abstract
The humoral and cellular immunity of convalescent COVID-19 patients is involved in pathogenesis and vaccine immunity. In this study, through CoV-psV neutralization assay and IFN-γ ELISpot testing in 30 cases of COVID-19 patients after 9 months post-SARS-CoV-2 infection, it found that the ratio of memory/naive CD4+ T lymphocytes cells and levels of anti-SARS-CoV-2-IgM and RBD-IgM were slightly but significantly higher in COVID-19 severe convalescent patients than that in non-severe patients. The specific cellular and humoral immunity against SARS-CoV-2 were detectable, regardless of the severity of the disease in the acute phase. This information may help understanding the immune status after SARS-CoV-2 infection.

Keywords SARS-CoV-2 · Virus-specific T cells · Neutralizing antibodies · Humoral immunity

Introduction
COVID-19 is currently a major public health issue of international concern. In the context of the widespread global vaccination of COVID-19 to establish herd immunity, recent studies have been focused on whether a single dose of COVID-19 vaccine can establish an adequate protective barrier for previously SARS-CoV-2 infected persons [1–5]. Therefore, it is of great significance to investigate the specific cellular immunity and humoral immunity during the convalescence period after SARS-CoV-2 infection.

Most severe COVID-19 patients developed lymphopenia as well as pneumonia with higher plasma levels of pro-inflammatory cytokines [6–8], suggesting that the host immune system is involved in pathogenesis. Studies reported by Zhou et al. [9] and Krammer et al. [10] found that sera
of COVID-19 patients could inhibit SARS-CoV-2 entry in target cells, indicating the involvement of humoral immunity. A study from Finland found a low level of neutralizing antibody titer in COVID-19 patient [11]. SARS-CoV-2-specific antibody responses have been identified in several studies [12–14]. However, to what extent adaptive immune responses to SARS-CoV-2 infection associate with the clinical course of COVID-19 remains unclarified. How long the neutralizing antibodies can sustain after SARS-CoV-2 infection and their relationships with the severity of the disease also needs to be understood.

In this study, we followed up with COVID-19 patients who were discharged about 9 months post-SARS-CoV-2 infection. The relation of SARS-CoV-2-specific antibody and T cell responses with the severity of disease was reported.

Methods

Study design and participants

All patients with COVID-19 enrolled in this study were diagnosed according to World Health Organization interim guidance [15]. They were initially admitted to Zhongnan Hospital of Wuhan University from 21 January to 3 March 2020. In severity assessment on admission, severe COVID-19 was defined according to a previous study [16]. They were treated at our hospital throughout the whole course of the disease and completed the follow-up test within the specified time. In another 16 cases, sex- and age-matched healthy blood donors were included as healthy control. The source of study subjects in this study was shown in Fig. 1.

Flow cytometry and detection of SARS-CoV-2-specific IgG and IgM

The CD3+/CD4+/CD8+ T-cell, CD19+ B-cell, and CD16+/CD56+ NK-cell counts testing procedure were performed according to a previously study [17]. The detection of SARS-CoV-2-specific IgG and IgM was performed according to the manufacturer’s instructions. An antibody concentration of ≥ 10.0 AU/mL was considered reactive and of < 10.0 AU/mL non-reactive.

Coronavirus spike protein pseudotyped virus (CoV-psV) neutralization assay

Coronavirus spike protein pseudotyped virus (CoV-psV) were packaged following a previously described protocol using a replicate-deficient VSV-based rhabdoviral pseudotyping system (VSV-dG) [18]. Serial dilutions of nanobodies were mixed with pseudoviruses, incubated for 30 min at room temperature, and then added to Caco-2 cells in a 96-well plate. Sixteen- to twenty-four hours later, cells were lysed by 1 × passive lysis buffer (Promega, United States) at room temperature for 15 min. Luciferase activity in the cell lysate was determined by a Bright-Glo luciferase assay kit (Promega, United States) and measured through a GloMax 20/20 Luminometer (Promega, United States).

Interferon gamma (IFN-γ) ELISpot

ELISpot assay procedures were referred to as described according to the previous report [19]. The spot increment ≥ 6 was considered positive. However, when the numbers of

Fig. 1 Flowchart of the study subjects
negative (unstimulated) spot forming units (SFU) ≥ 6, stimulated spot numbers ≥ twofold higher than negative (unstimulated) controls to any of the antigens were considered positive.

**Statistical analysis**

Categorical variables were described as frequency rates and percentages, and continuous variables were described using mean, median, and interquartile range (IQR) values. \( \chi^2 \) analysis or Fisher exact test was conducted to examine the categorical variables. Means for continuous variables in this study were compared using independent group t tests when the data were normally distributed; otherwise, the Mann–Whitney test was used. Two-sided \( p \) values < 0.05 were considered statistically significant.

**Results**

**Patient characteristics**

Of the 30 COVID-19 convalescent patients, the median age was 56 ± 12 years (range, 33–76 years) and 15 were male. Nine (30.0%) had 1 or more comorbidity: hypertension (2 (6.67%), diabetes (1 (3.33%), cardiovascular disease (2 (6.67%), chronic kidney disease (1 (3.33%), hypertension and diabetes coexisting conditions (2 (6.67%)), and hypertension and cardiovascular disease coexisting conditions (2 (6.67%)). The age of the 16 healthy blood donors was comparable to those COVID-19 convalescent patients and 8 were male. Five (31.25%) had comorbidity: hypertension (2 (12.50%), diabetes (2 (12.50%)), and hyperthyroidism (1 (6.25%). SARS-CoV-2-specific IgG and IgM were negative for all blood donors.
**Clinical laboratory findings**

A complete blood count, liver and renal function, inflammatory biomarkers, and coagulation function were measured in COVID-19 convalescent individuals to monitor the degree of recovery from SARS-CoV-2 infection. The results showed that the levels of white blood cells, lymphocyte counts, PLT, BUN, ALT, Alb, GGT, LDH, IL-6, CRP, PT, and D-Dimer were significantly improved and returned to normal levels in the convalescent phase. The clinical results of these above indicators were significantly different from those in the acute phase during hospitalization.

**Fig. 3** SARS-CoV-2-specific antibodies response in COVID-19 convalescent patients by the severity of the disease. **a** The levels of anti-SARS-CoV-2-IgM ($P = 0.014$), anti-S1-IgM ($P = 0.0004$), and anti-RBD-IgM ($P = 0.0002$) in severe convalescent patients were higher than that in non-severe convalescent patients, except for the anti-N-IgM ($P = 0.917$); **b** all the levels of IgG antibodies, including anti-SARS-CoV-2-IgG, anti-S1-IgG, anti-RBD-IgG, and anti-N-IgG, were similar in convalescent patients, regardless of the disease severity in the acute phase.
Lymphocyte subtypes

Compared with the lymphocyte subtypes in the acute phase during hospitalization, the levels of CD4+ T lymphocyte cells (Fig. 2a), CD8+ T lymphocyte cells (Fig. 2b), CD19+ B cells (Fig. 2c), and CD16+CD56+ NK cells (Fig. 2d) significantly increased and returned to normal levels in convalescent phase, except for the ratio of memory/naive CD4+ T lymphocytes cells (Fig. 2e, \( P = 0.0135 \)). Moreover, the ratio of memory/naive CD4+ T lymphocytes cells in COVID-19 severe convalescent patients was higher than that in non-severe patients (Fig. 2f, \( P = 0.0431 \)).

SARS-CoV-2-specific antibodies

The levels of anti-SARS-CoV-2-IgG, anti-SARS-CoV-2-N-IgG, S1-IgG, and RBD-IgG in severe COVID-19 patients were similar to those in non-severe COVID-19 patients (Fig. 3a). Interestingly, except for N-IgM, the levels of anti-SARS-CoV-2-IgM, S1-IgM, and RBD-IgM in severe COVID-19 patients were significantly greater than those in non-severe COVID-19 patients (Fig. 3b).

Moreover, all the 13 non-severe patients were negative for anti-SARS-CoV-2-IgM, whereas 29.4% (5/17) of the severe patients were positive for anti-SARS-CoV-2-IgM \( (\chi^2 = 4.588, P = 0.032) \) (Table 1).

Serum neutralization capabilities in COVID-19 convalescent patients

When the serums of COVID-19 convalescent patients were diluted as 1:125, the inhibition rate against SARS-CoV-2 infection was as high as 60%, whereas the inhibition rate was less than 30% when the serums of healthy blood donors were diluted as 1:5. However, there was no significant difference in terms of the inhibition rate against SARS-CoV-2 infection among severe and non-severe COVID-19 patients (Fig. 4).

| Table 1 Follow-up results of anti-SARS-CoV-2 IgG and IgM between severe and non-severe patients at 283 days after the diagnosis of SARS-CoV-2 infection |
|---------------------------------------------------|-----------------|----------------|-----------------|
| Severe cases (n = 17) | Non-severe cases (n = 13) | Test | P value |
| IgG | | | |
| Total IgG (+) (n, %) | 17 (100%) | 12 (92.3%) | 1.353 | 0.245 |
| N-IgG (+) (n, %) | 16 (94.1%) | 10 (76.9%) | 1.885 | 0.170 |
| S1-IgG (+) (n, %) | 17 (100%) | 13 (100%) | - | - |
| RBD-IgG (+) (n, %) | 17 (100%) | 12 (92.3%) | 1.353 | 0.245 |
| IgM | | | |
| Total IgM (+) (n, %) | 5 (29.4%) | 0 (0%) | 4.588 | 0.032 |
| N-IgM (+) (n, %) | 1 (5.9%) | 1 (7.7%) | 0.039 | 0.844 |
| S1-IgM (+) (n, %) | 4 (23.5%) | 0 (0%) | 3.529 | 0.060 |
| RBD-IgM (+) (n, %) | 3 (17.6%) | 0 (0%) | 2.549 | 0.110 |

Fig. 4 Serum neutralization capabilities in COVID-19 convalescent patients (The inhibition rate against SARS-CoV-2 infection in COVID-19 convalescent patients was as high as 60%, whereas the inhibition rate was less than 30% when the serums of healthy blood donors were diluted as 1:5.)
Proliferative capabilities of SARS-CoV-2-specific T cells in convalescent COVID-19

In severe and non-severe COVID-19 convalescent patients, the proportion of SARS-CoV-2-specific T cells response was 82.4% (14/17) and 84.6% (11/13), respectively. Moreover, the spot forming units by the IFN-γ ELISpot test were 23 ± 4 and 30 ± 7 in the 14 severe and 11 non-severe COVID-19 convalescent patients, respectively. The spot forming units by the IFN-γ ELISpot test was similar in severe and non-severe COVID-19 convalescent patients (Fig. 5).

Discussion

In this follow-up study, as expected, we found complete blood cell count, liver and kidney function, coagulation function, and inflammatory cytokine levels of the patients all returned to normal levels. Interestingly, the ratio of memory/naive CD4+ T lymphocytes cells and levels of anti-SARS-CoV-2-IgM and RBD-IgM in severe COVID-19 convalescent patients were slightly but significantly higher than that in non-severe cases. Importantly, it found that in the convalescence phase, SARS-CoV-2 infection patients had specific cellular immunity, and the specific antibodies had the capacity to neutralize SARS-CoV-2.

In this study, SARS-CoV-2 IgM was positive in 29.4% (5/17) severe COVID-19 convalescent individuals, whereas none was positive in non-severe patients. Further follow-up is needed to determine how long SARS-CoV-2 IgM lasts in severely COVID-19 patients, but a conclusion we can draw now was that the late disappearance of SARS-CoV-2 IgM may indicate a more serious condition.

As previously reported [8], SARS-CoV-2-specific humoral immunity was detected in newly discharged patients. In this study, we demonstrated that immune-mediated protection to viral infection can be detected after 9 months post-SARS-CoV-2 infection. Recently, several studies characterizing adaptive immune responses to SARS-CoV-2 infection have reported that most COVID-19 convalescent individuals have detectable neutralizing antibodies, which correlate with the numbers of virus-specific T cells [8, 9, 20–23]. This study further found that virus-stimulated neutralizing antibodies can sustain for several months, and the capacity to neutralize SARS-CoV-2 in severe convalescent patients was close to that in non-severe patients.

Antibody responses in patients previously infected with SARS-CoV tended to be short-lived [24], whereas SARS-CoV-specific memory T cells were found to persist for 6 years after infection in SARS survivors [25–27]. Memory CD4+ T cells have multiple roles in initiating and propagating the immune response [28]. SARS-CoV-2-specific T cells in the acute phase and the convalescent phase were proved to be different, displaying as a highly activated cytotoxic phenotype that correlated with various clinical markers of disease severity in the acute phase and a stem-like memory phenotype in the convalescent phase [29]. However, little is known about whether the proportion of these cells changed after SARS-CoV-2 infection. In this study, the ratio of memory/naive CD4+ T lymphocytes cells in severe COVID-19 convalescent patients was found to be greater than that in non-severe COVID-19 convalescent patients, which indicated that specific cellular immunity plays an important role in pathogenesis after SARS-CoV-2 infection. This potential effect can persist about 9 months post-SARS-CoV-2 infection. Whether this memory T response has a protective effect on SARS-CoV-2 re-infection is still worth further study.

We recognize that our study has limitations. Given the single-center study and the small number of samples, the representativeness of the outcome may be limited to some extent. Also, although we confirmed that humoral immunity responses and specific cellular immunity in COVID-19 convalescent patients can be detectable after 9 months of SARS-CoV-2 infection, whether the neutralizing antibodies and specific cellular immunity have a protective effect on
Delta or Lambda variants is still worth exploring. Our team will try our best to make up for the defects in future studies.

Conclusion

The specific cellular and humoral immunity against SARS-CoV-2 were detectable after 9 months post-SARS-CoV-2 infection, regardless of the severity of the disease in the acute phase. This information may help understanding the immune status after SARS-CoV-2 infection.

Author contribution

Y.X. and X.W. conceptualized the study design; Z.S. and W.H. recruited the patients, collected specimens, collected demographic, and clinical data; Q.L., Z.S., Y.Z., Z.Z., and W.H. did the laboratory tests; T.C., R.Y., Y.Z., and Y.C. interpreted the results; R.Y. wrote the initial drafts of the manuscript; Y.C., R.Y., Y.X., and X.W. revised the manuscript. All authors read and approved the final manuscript.

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