

Epidemiological feature, viral shedding, and antibody seroconversion among asymptomatic SARS-CoV-2 carriers and symptomatic/presymptomatic COVID-19 patients

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Research

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Abstract

Background

Novel coronavirus disease 2019 (COVID-19) caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) is pandemic. However, data concerning the epidemiological features, viral shedding, and antibody dynamics between asymptomatic SARS-CoV-2 carriers and COVID-19 patients remain controversial.

Methods

A total of 193 subjects in Ningbo and Zhoushan, Zhejiang, China, were enrolled in this study from January 21 to March 6, 2020. All subjects were tested positive for SARS-CoV-2 genomic RNA by quantitative reverse transcription PCR and then followed up to monitor the dynamics of serum antibody immunoglobulin M (IgM) and immunoglobulin G (IgG) against SARS-CoV-2 using enzyme-linked immunosorbent assays. Scatter diagram to demonstrate the distribution of IgM and IgG among asymptomatic carriers and COVID-19 patients were generated by R.

Results

Of the 193 subjects, 31 were asymptomatic SARS-CoV-2 carriers, 149 were symptomatic COVID-19 patients, and 14 were COVID-19 patients during the incubation. Compared to symptomatic COVID-19 patients, asymptomatic SARS-CoV-2 carriers were younger and had higher levels of white blood cell and lymphocyte, lower levels of C-reactive protein (CRP) and viral load, and shorter viral shedding time. Seroconversion of IgM against SARS-CoV-2 from positive to negative in asymptomatic carriers took 7.50 (IQR, 4.75–11.50) days, which was significantly shorter than 25.50 (IQR, 6.75–56.75) days in COVID-19 patients ($P=0.030$). The proportion of those persistently seropositive for IgG against SARS-CoV-2 was higher in COVID-19 patients than in asymptomatic carriers (66.1% vs. 33.3%, $P=0.037$). Viral load was higher in symptomatic than presymptomatic COVID-19 patients. Viral shedding was longer in presymptomatic COVID-19 patients than in asymptomatic carriers. In 4 familial clusters of SARS-CoV-2 infection, asymptomatic carriers were mainly children and young adults while severe COVID-19 was mainly found in family members older than 60 years with underlying diseases. Asymptomatic carriers acquired infection more from intra-familial transmission than did COVID-19 patients (89% vs. 61%, $P=0.028$).

Conclusion

Asymptomatic carriers might have a higher antiviral immunity to clear SARS-CoV-2 than symptomatic COVID-19 patients and this antiviral immunity might not be contributable to humoral immunity. The

severity of COVID-19 is associated with older age and underlying diseases in familial clustering cases.

Introduction

Novel coronavirus disease in 2019 (COVID-19) caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) has been pandemic since it was firstly recognized in China in late December 2019 [1]. The case number keeps increasing. Globally, as of 9:42am CET, December 11, 2020, there have been 68,845,368 confirmed cases of COVID-19, including 1,570,304 deaths, reported to World Health Organization [2]. Family clustering and hospital-based transmission were the two major epidemiological features of this outbreak at the early stage, and continued to be an important cause of community-based SARS-CoV-2 transmission in a low prevalence region [3–6]. COVID-19 patients and asymptomatic carriers are the main sources of SARS-CoV-2 infection but might have differences in some features [7]. However, data concerning SARS-CoV-2 transmission and viral shedding duration between COVID-19 patients and asymptomatic SARS-CoV-2 carriers remain controversial. It has been summarized from early studies that viral load of asymptomatic carriers is comparable to symptomatic patients [8]. In another study, it has been demonstrated that a considerably higher viral load is present in samples from fatal cases compared to asymptomatic carriers [9]. Difference in the dynamics of antibody against SARS-CoV-2 between asymptomatic carriers and COVID-19 patients remains unknown. To provide the information for recognizing differences between asymptomatic and symptomatic SARS-CoV-2 infected subjects, we conducted a study to investigate the epidemiological feature, laboratory findings, viral shedding, and antibody conversion of SARS-CoV-2 infected cases among asymptomatic SARS-CoV-2 carriers and symptomatic/presymptomatic COVID patients in two cities, a low prevalence region in Zhejiang, China.

Methods

Study design and patients

It is an ambispective cohort study. The study enrolled all confirmed cases with SARS-CoV-2 infection on Ningbo city and a familial clustering infection with SARS-CoV-2 in Putuo district of Zhoushan city from January 21 to March 6, 2020. Epidemiological, clinical characteristics, pathogen and serological test results were collected by Ningbo Municipal Center for Disease Control and Prevention (Ningbo CDC) and CDC of Putuo district, Zhoushan (Putuo CDC). Some baseline information including demographic and pathogenic data of those patients were reported [10, 11]. After that, we continued to investigate viral load and viral shedding of SARS-CoV-2, laboratory tests, and the dynamics of serum immunoglobulin M (IgM) and immunoglobulin G (IgG) against SARS-CoV-2 between asymptomatic carriers and COVID-19 patients. All patients were followed up to monitor the dynamics of IgM and IgG against SARS-CoV-2. Those with antibody tests for two times or more within 160 days were included in antibody seroconversion analysis. Diagnoses and disease staging of COVID-19 were carried out according to the Protocol for the Diagnosis and Treatment of COVID-19 (Version 7th), National Health Commission of the People's Republic of China [12]. Specifically, COVID-19 was diagnosed if the patient was tested positive for SARS-CoV-2 genomic RNA and accompanied by clinical symptoms including fever and cough. Asymptomatic carrier was

identified in close contactors of COVID-19 patients if they did not have any symptom. We also classified as COVID-19 patients with onset of disease and those during the incubation period (presymptomatic). All diagnosed COVID-19 patients were classified as mild, common, severe, and extremely severe types. This study was approved by the Ethics Commissions of Ningbo CDC and Putuo CDC. Written informed consent was waived for emerging infectious diseases.

Epidemiological survey

A semi-structured questionnaire was applied to obtain demographic information, exposure information of the familiar clustering cases via face-to-face interview and telephone calls by well-trained professionals. The data regarding any travel history to high risk areas with COVID-19 epidemic, contact with confirmed cases or asymptomatic carriers tested positive for SARS-CoV-2 genomic RNA, contact with patients with some symptoms like fever, dry cough, and expectoration in the past 2–3 weeks before illness onset. Any chance and duration of attending any kinds of population gatherings were recorded as well.

Clinical information included the date of symptom onset and admission to designated hospitals, clinical manifestation, routine laboratory examinations, and radiographic examinations. The clinical manifestations, chest computed tomography images, and laboratory results of patients in Ningbo were collected from the electronic medical record systems in the two designated hospitals: Ningbo First Hospital and Huamei Hospital. The information of patients from Zhoushan was collected from Zhoushan Maternal and Child Health Care Hospital and Putuo Hospital. Two researchers (PL and YD) independently reviewed all of the data to doubly check the accuracy of data collected.

Examination of SARS-CoV-2 genomic RNA

Quantitative reverse transcription-PCR (qRT-PCR) assay was applied to detect SARS-CoV-2 genomic RNA in nasal and throat swabs, sputum, and feces of patients. Two targets, the open reading frame 1ab (ORF1ab) and nucleoprotein (N) genes of SARS-CoV-2 were detected using the testing kits. Patients in Ningbo were examined using the test kits manufactured by Shanghai BioGerm Medical Technology (Shanghai, China) and Daan Gene Co., Ltd (Guangzhou, China) [13]. Patients in Zhoushan were examined using the test kits manufactured by Shanghai GeneDx Biotech Company (Shanghai, China) [14]. Sample was positive if the cycling threshold (CT) values of reverse-transcription polymerase chain reaction (RT-PCR) for the ORF1ab and the N genes were less than 37. Sample was negative if no CT value, or CT value of greater than 40, or unrepeatable CT value in the range of 37–40.

Detection of antibodies against SARS-COV-2

IgM and IgG against SARS-CoV-2 in the frozen reserved fasting serum samples were detected using the diagnostic enzyme-linked immunosorbent assay (ELISA) kits. The diagnostic kits used for Ningbo patients and Zhoushan patients were manufactured by Innovita Biological Technology (Tangshan, China) and Nanjing Wending Biotech Company (Nanjing, China), respectively [15]. Briefly, testing results were determined by the color reaction. Dark band was considered positive. Light band indicated weak positive. Disappearance of the expected band was considered negative.

Statistical Analysis

Categorical variables were compared using the χ^2 test or the Fisher exact test. Continuous variables were described using median and interquartile range (IQR) values and then compared using Mann-Whitney U test or Kruskal Wallis test. These statistical analyses were two-sided and performed using R, version 3.6.2 (R Foundation for Statistical Computing, Canberra, Austria). Scatter diagram to demonstrate the distribution of IgM and IgG against SARS-CoV-2 among asymptomatic carriers and symptomatic and presymptomatic COVID-19 patients were generated by R software. A *P* value of < 0.05 was considered significant for two independent groups. An adjusted *P* value of < 0.017 was considered significant by Bonferroni-Dunn test for pairwise comparison among three groups.

Results

Epidemiological characteristics of asymptomatic SARS-CoV-2 carriers, symptomatic COVID-19 patients, and presymptomatic COVID-19 patients

A total of 193 SARS-CoV-2 infected subjects were enrolled in this study. Of those, 31 were asymptomatic carriers, 148 were symptomatic COVID-19 patients, and 14 were presymptomatic COVID-19 patients. Of the 187 patients from Ningbo, 3 family clusters were included. Those patients were close contacts of confirmed COVID-19 patients and then tested positive for SARS-CoV-2 genomic RNA, from January 21 to March 6, 2020. For a family cluster from Putuo, a 41-year-old man contacted with one relative, a confirmed COVID-19 patient from Xianning city, Hubei, China during January 26 and January 31, 2020. He then went to Putuo and lived with 6 family members from January 31 to February 3, 2020. All family members were tested positive for SARS-CoV-2 genomic RNA, with 1 asymptomatic SARS-CoV-2 carrier and 5 symptomatic COVID-19 cases. Our epidemiological survey indicated the family members did not have opportunity to get the infection from other sources.

Asymptomatic SARS-CoV-2 carriers were significantly younger than symptomatic COVID-19 patients and presymptomatic COVID-19 patients. Compared to symptomatic COVID-19 patients, asymptomatic SARS-CoV-2 carriers had higher levels of circulating white blood cell (WBC) and lymphocyte, lower levels of C-reactive protein (CRP) and viral load, and shorter viral shedding time. Interestingly, viral load was significantly lower in presymptomatic COVID-19 patients than in symptomatic COVID-19 patients. The viral shedding duration was significantly longer in presymptomatic COVID-19 patients than in asymptomatic carriers. The first-time serological tests showed that nearly one-third of asymptomatic carriers and symptomatic COVID-19 patients were seronegative for IgM against SARS-CoV-2 while the seronegative rates for IgG to SARS-CoV-2 were around 7% in the two populations, respectively (Table 1).

Table 1

Baseline information of COVID-19 patients and asymptomatic SARS-CoV-2 carriers in Ningbo and Zhoushan cities of Zhejiang province, China

	Asymptomatic carrier (N = 31)	Symptomatic COVID-19 patients (N = 148)	Presymptomatic COVID-19 patients (N = 14)	P*	P&	P#
Gender				0.453	1.000	0.750
Male	15 (48.4)	50 (33.8)	7 (50.0)			
Female	16 (51.6)	98 (66.2)	7 (50.0)			
Age, years	42.00(24.00–55.00)	53.00(38.00–62.75)	55.00(39.50–71.00)	< 0.001	0.002	1.000 [§]
< 30	9 (29.0)	17 (11.5)	1 (7.1)	0.076	0.345	1.000
30–59	18 (58.1)	89 (60.1)	8 (57.1)			
≥ 60	4 (12.9)	42 (28.4)	5 (35.7)			
Underlying diseases				1.000	1.000	1.000
No	24 (77.4)	104 (71.7)	9 (64.3)			
Yes	7 (22.6)	41 (28.3)	5 (35.7)			
WBC, 10 ⁹ /L	5.83(5.00–7.11)	4.63(3.80–5.69)	5.83(4.73–6.75)	< 0.001	1.000	0.075 [§]
Lymphocyte, 10 ⁹ /L	1.53(1.32–2.11)	1.22(0.86–1.60)	1.26(0.89–1.86)	0.003	0.420	1.000 [§]
CRP, mg/L	1.00(0.60–2.99)	6.90(2.04–18.20)	3.07(0.97–14.45)	< 0.001	0.317	0.885 [§]

Abbreviations: WBC, white blood cell; CRP, C-reactive protein; CT, cycling threshold; IQR, interquartile range.

P*, Asymptomatic carrier vs. COVID-19 patients.

P&, asymptomatic carrier vs. COVID-19 patients during the incubation.

P#, COVID-19 patients vs. COVID-19 patients during the incubation.

[§]Continuous variables are expressed as median (IQR). P value was calculated using Kruskal Wallis test.

Categorical variables are expressed as n (%). P values were calculated using χ^2 test and Fisher's exact test.

	Asymptomatic carrier (N = 31)	Symptomatic COVID-19 patients (N = 148)	Presymptomatic COVID-19 patients (N = 14)	P*	P&	P#
Viral shedding, day	24.00(21.00-30.80)	46.50(35.00-58.00)	48.00(23.75-51.25)	< 0.001	0.002	1.000 [§]
qRT-PCR-CT values	31.40(27.50-34.50)	29.00(24.25-32.00)	33.50(28.25-36.25)	0.004	0.410	0.003[§]
IgM				1.000	1.000	1.000
Negative	12 (38.7)	43 (31.9)	5 (50.0)			
Positive	19 (61.3)	91 (67.4)	5 (50.0)			
Weak Positive	0 (0.0)	1 (0.7)	0 (0.0)			
IgG				0.112	1.000	0.504
Negative	2 (6.5)	10 (7.4)	1 (10.0)			
Positive	25 (80.6)	122 (90.4)	8 (80.0)			
Weak Positive	4 (12.9)	3 (2.2)	1 (10.0)			
Abbreviations: WBC, white blood cell; CRP, C-reactive protein; CT, cycling threshold; IQR, interquartile range.						
P*, Asymptomatic carrier vs. COVID-19 patients.						
P&, asymptomatic carrier vs. COVID-19 patients during the incubation.						
P#, COVID-19 patients vs. COVID-19 patients during the incubation.						
[§] Continuous variables are expressed as median (IQR). P value was calculated using Kruskal Wallis test.						
Categorical variables are expressed as n (%). P values were calculated using χ^2 test and Fisher's exact test.						

Dynamics in seroconversion of IgM and IgG against SARS-CoV-2 between asymptomatic carriers and COVID-19 patients

Of the 193 study subjects, 74 (15 asymptomatic carriers and 59 COVID-19 patients) had two consecutive test results of IgM and IgG against SARS-CoV-2 within 160 days. SARS-CoV-2-specific IgM seroconversion from positive to negative or weak positive occurred in 9 (60.0%) asymptomatic carriers,

while this occurred in 28 (47.5%) COVID-19 patients ($P = 0.647$). However, the median time interval of IgM seroconversion from positive to negative was 7.50 (IQR, 4.75–11.50) days in asymptomatic carriers, which was significantly shorter than 25.50 (IQR, 6.75–56.75) days in COVID-19 patients ($P = 0.030$). SARS-CoV-2-specific IgG seroconversion from positive to negative or weak positive occurred in 8 (53.4%) asymptomatic carriers, while this occurred in 15 (25.5%) COVID-19 patients ($P = 0.059$). Importantly, 5 (33.3%) of asymptomatic carriers were consistently seropositive for IgG against SARS-CoV-2, however, this was 39 (66.1%) in COVID-19 patients ($P = 0.037$). Furthermore, there was no significant difference in the time interval of IgG seroconversion between the two groups (Table 2, Fig. 1).

Table 2
Seroconversion of IgG and/or IgM antibody against SARS-CoV-2 during follow-up time

	Overall	Asymptomatic carrier (N = 15)	COVID-19 patients (N = 59)	<i>P</i>
Seroconversion of IgM antibody				0.647
From positive to negative	28(37.8)	8(53.3)	20(33.9)	0.234
From positive to weak positive	9(12.2)	1(6.7)	8(13.6)	0.676
Consistent negative	28(37.8)	5(33.3)	23(39.0)	0.772
Consistent positive	9(12.2)	1(6.7)	8(13.6)	0.676
Time interval of IgM antibody seroconversion, days				
From positive to negative	13.00(6.00-33.75)	7.50(4.75-11.50)	25.50(6.75-56.75)	0.030
From positive to weak positive	16.00(3.00-28.00)	12.00(12.00-12.00)	19.00(3.00-30.00)	0.694
Consistent negative	11.50(7.00-25.50)	27.00(10.00-37.00)	11.00(7.00-22.50)	0.186
Consistent positive	9.00(8.00-13.00)	13.00(13.00-13.00)	8.50(7.50-14.50)	0.558
Seroconversion of IgG antibody				0.059
From positive to negative	10(13.5)	4(26.7)	6(10.2)	0.110
From positive to weak positive	13(17.6)	4(26.7)	9(15.3)	0.446
Consistent negative	7(9.5)	2(13.3)	5(8.5)	0.624
Consistent positive	44(59.5)	5(33.3)	39(66.1)	0.037
Time interval of IgG antibody seroconversion, days				
From positive to negative	24.00(10.00-48.00)	24.00(9.25-58.75)	27.50(10.00-48.00)	1.000
From positive to weak positive	7.00(4.00-22.00)	5.50(4.25-7.75)	11.00(4.00-30.00)	0.279

Data are median (IQR), n (%), P values compare using χ^2 test, Fisher's exact test, or Kruskal Wallis test. IQR, interquartile range.

	Overall	Asymptomatic carrier (N = 15)	COVID-19 patients (N = 59)	<i>P</i>
Consistent negative	10.00(7.50–18.50)	9.00(8.50–9.50)	12.00(7.00–25.00)	0.699
Consistent positive	13.50(6.75–24.00)	13.00(12.00–14.00)	14.00(6.00–25.00)	0.970
Data are median (IQR), n (%), <i>P</i> values compare using χ^2 test, Fisher's exact test, or Kruskal Wallis test. IQR, interquartile range.				

Intrafamilial transmission of SARS-CoV-2

Four family clusters were recorded in the study, 3 from Ningbo and 1 from Zhoushan. A total of 15 subjects (7 asymptomatic carriers and 8 COVID-19 patients) were confirmed to be infected by SARS-CoV-2 in the 4 familial clusters. Of the 7 asymptomatic carriers, 3 were children at the age of 12 years or younger, 3 adults aged from 18 to 60 years, and a 75-year-old woman. Of the 8 COVID-19 cases, 4 were older than 60 years and diagnosed as severe cases, 3 of the 4 severe cases had underlying diseases. The remaining 4 were mild patients at the age between 18 and 60 years. Only one of the 4 mild cases had an underlying disease. Although intra-familial transmission was the major cause of acquiring SARS-CoV-2 infection, the proportion of those who acquired SARS-CoV-2 infection via intra-familial transmission was significantly higher in asymptomatic carriers than in COVID-19 patients (89% vs. 61%, $P = 0.028$) (Fig. 2).

Discussion

To characterize the epidemiological features including immunological response, viral transmission, and antibody seroconversion in asymptomatic SARS-CoV-2 carriers, we made a comprehensive comparison between asymptomatic carriers and COVID-19 patients in this study. Compared to symptomatic COVID-19 patients, asymptomatic SARS-CoV-2 carriers were younger and had higher levels of circulating WBC and lymphocyte and a lower level of CRP. These data indicate that asymptomatic carriers have a stronger antiviral immunity and a lower level of systemic inflammation. It has been proven that innate and adaptive lymphocytes and inflammatory factors were closely related to disease progression of COVID-19, from mild to severe [16, 17]. In a previous prospective study, we have demonstrated that lower circulating counts of T lymphocytes, CD4⁺ T cells, and CD8⁺ T cells as well as higher circulating levels of neutrophil proportion, neutrophil/lymphocyte ratio, interleukin-6, CRP, and procalcitonin facilitate the progression of COVID-19. Of those, CD8⁺ T cell exhaustion plays an important role in the pathogenesis of COVID-19 [18]. Other studies have also shown that disease severity is negatively associated with NK cells and CD3⁺, CD4⁺, and CD8⁺ T lymphocyte levels, while intensive expansion of highly cytotoxic effector T cell subsets, such as CD4⁺ effector-granulysin, CD8⁺ effector-granulysin, and NKT CD160, is associated with convalescence of COVID-19 patients [19–21]. These evidences strongly indicate that damage of innate immunity and T cell-mediated immunity, which might be caused by proinflammatory factor-induced inflammation, play key roles in the development of COVID-19.

In this study, we also found that around 7% of asymptomatic carriers and COVID-19 patients during or after the incubation were seronegative for IgG against SARS-CoV-2, indicating SARS-CoV-2 infection might not induce sufficient humoral immunity against SARS-CoV-2. In the follow-up study, IgM seroconversion from positive to negative was much faster in asymptomatic carriers than in COVID-19 patients ($P= 0.030$). The overall rate of IgG seroconversion from positive to negative or weak positive was around 30% within 160 days after the diagnosis (Table 2), indicating that IgG against SARS-CoV-2 is not stable. Importantly, seroconversion of IgG against SARS-CoV-2 from positive to negative or weak positive occurred 53.4% in asymptomatic carriers and 25.5% in COVID-19 patients ($P= 0.059$), while consistently seropositive rate of IgG against SARS-CoV-2 was significantly higher in COVID-19 patients than in asymptomatic carriers ($P= 0.037$). The similar observations concerning rapid seroconversion of the antibody against SARS-CoV-2 or short-lived immune response after mild infection were also reported in the frontline health care personnel in the US and active workers in France [22, 23]. These data indicate that humoral immunity against SARS-CoV-2 was not efficiently aroused by a relative short exposure of SARS-CoV-2 in asymptomatic carriers or in those with a stronger innate and cell immunity. Long-term seropositive rate of antibody against SARS-CoV-2 in COVID-19 patients, which has been previously reported [24, 25], indicates the feasibility of antibody-generating vaccination in the worldwide prophylactic action. However, it might be possible that the antibody against SARS-CoV-2 take part in the pathogenic process of COVID-19.

Interestingly, compared to COVID-19 patients, asymptomatic carriers had a lower level of viral load and shorter viral shedding time (Table 1). Our finding is different from a study carried out in Chongqing that asymptomatic carriers had a significantly longer duration of viral shedding than the symptomatic patients [26], possibly because of small sample size. Lower viral load and shorter viral shedding duration in asymptomatic carriers should be unlikely caused by the neutralizing antibody, because the antibody, either IgM or IgG, was declining more rapidly in asymptomatic carriers than in COVID-19 patients. Innate immunity and cell-mediated immunity should play key roles in repressing viral replication in asymptomatic carriers [27]. Lower viral load and shorter viral shedding time should be due to a relative stronger antiviral immunity, as a high viral load often predisposes adverse outcomes of COVID-19 [9, 28]. To develop effective vaccine against SARS-CoV-2, it is important to arouse the specific cell immunity, instead of focusing on humoral immunity.

In this study, we found that viral load increased from COVID-19 patients during the incubation to symptomatic COVID-19 patients (Table 1), indicating that infectivity is the highest at the stage of disease onset. Asymptomatic carriers had a lower level of viral load and shorter viral shedding duration, indicating that the transmissibility of asymptomatic carriers was relative weaker. In the 4 familial clusters, we found that asymptomatic carriers were mostly children and young adults, mild patients were young and middle-aged adults between 18 and 60 years, and severe cases were older than 60 years with underlying diseases. Family members were exposed to the same source of infection. However, they had diverse clinical manifestations: children were often asymptomatic whereas old members were very sick. This observation is quite in consistent with the findings of large epidemiological studies that children acquire SARS-CoV-2 infection mostly have mild respiratory symptoms or are asymptomatic, whereas

elderly patients with COVID-19, especially male patients, are more likely to progress into severe-type and even die of this disease [29–33]. Thus, the host immunity and underlying inflammation, which is often affected by ageing, underlying diseases, and dysregulated macrophage response [34], should be the major determinants of disease severity of COVID-19. A family cluster occurred in Zhoushan is a suitable example to address this issue. Although asymptomatic carriers often acquire the infection from family members, they can transmit SARS-CoV-2 into family members and hospital centers, and eventually kill aged members. As a considerable percentage of SARS-CoV-2 infections may be asymptomatic or presymptomatic, enhanced testing approaches may be needed to detect those who transmit the virus.

Our study has some limitations. First, follow-up should be extended to observe the duration of SARS-CoV-2-specific antibodies. Second, sample size of asymptomatic carriers with SARS-CoV-2 infected was relatively small.

Conclusions

Asymptomatic carriers have a higher level of antiviral immunity and lower level of inflammation to clear SARS-CoV-2, resulting in a lower capacity of SARS-CoV-2 transmission, than do symptomatic COVID-19 patients. This antiviral immunity should not be contributable to humoral immunity because both IgM and IgG against SARS-CoV-2 are declining more rapidly in asymptomatic carriers than in COVID-19 patients. The severity of COVID-19 is associated with older age and underlying diseases in familial clustering cases. This study is not only important in elucidating the mechanisms by which SARS-CoV-2 interacts with host immunity in determining the outcome of SARS-CoV-2 infection, but also help in optimizing the strategy for the worldwide prophylactic action to develop SARS-CoV-2 vaccine.

Abbreviations

COVID-19: coronavirus disease 2019; SARS-CoV-2: severe acute respiratory syndrome coronavirus 2; qRT-PCR: Quantitative reverse transcription-PCR assay; IgM: Immunoglobulin M; IgG: immunoglobulin G.

Declarations

Acknowledgments

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Ethics approval and consent to participate

This study was approved by the Ethics Commission of Ningbo CDC and Putuo CDC.

Availability of data and materials

The datasets used and analyzed during the current study are available from the corresponding author on reasonable request.

Competing interests

The authors declare that they have no competing interests

Consent for publication

Written informed consent was waived for emerging infectious diseases.

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Authors' contributions

Miao Liu and Yi Chen: investigation, field survey, data organization. Ping Li and Yi-bo Ding: quality control, data analysis, and composition of figures and table. Lei-jie Liu and Bo Yi: investigation, etiological identification, antibody test. Hong-jun Dong, Xu-ying Lao and Ting Wu: project management. Dong-liang Zhang and Xiao-Jie Tan: data entry and analysis. Ke-qing Ding, Hai-bo Wang and Zhong-fa Wang: laboratory test and etiological identification. Guo-zhang Xu and Guang-wen Cao: conceptualization, investigation, supervision, and funding acquisition. Guang-wen Cao wrote this manuscript.

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Figures

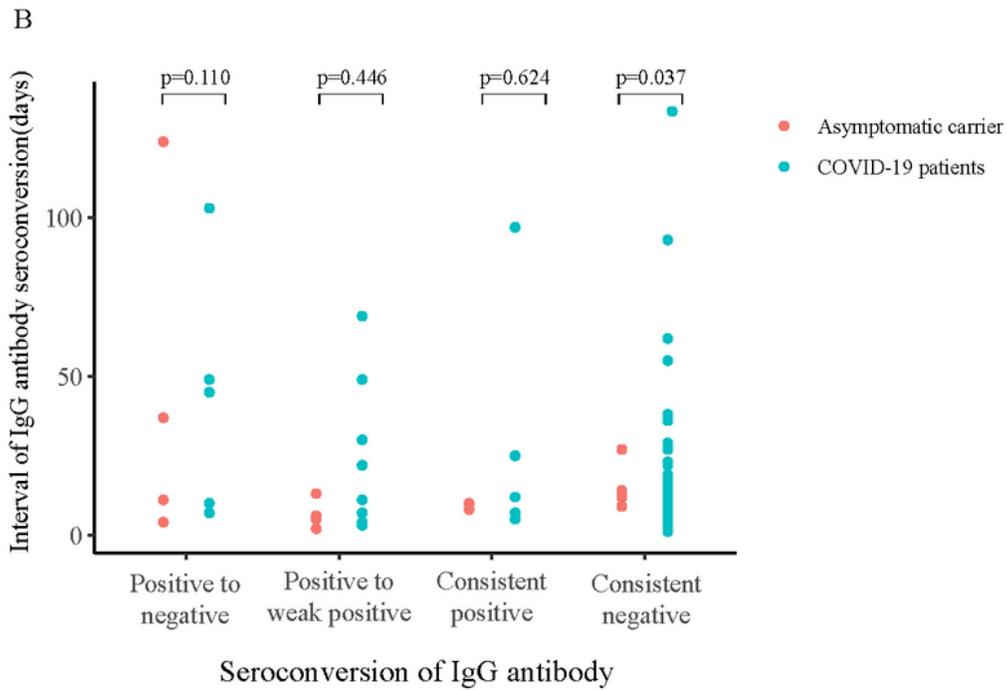
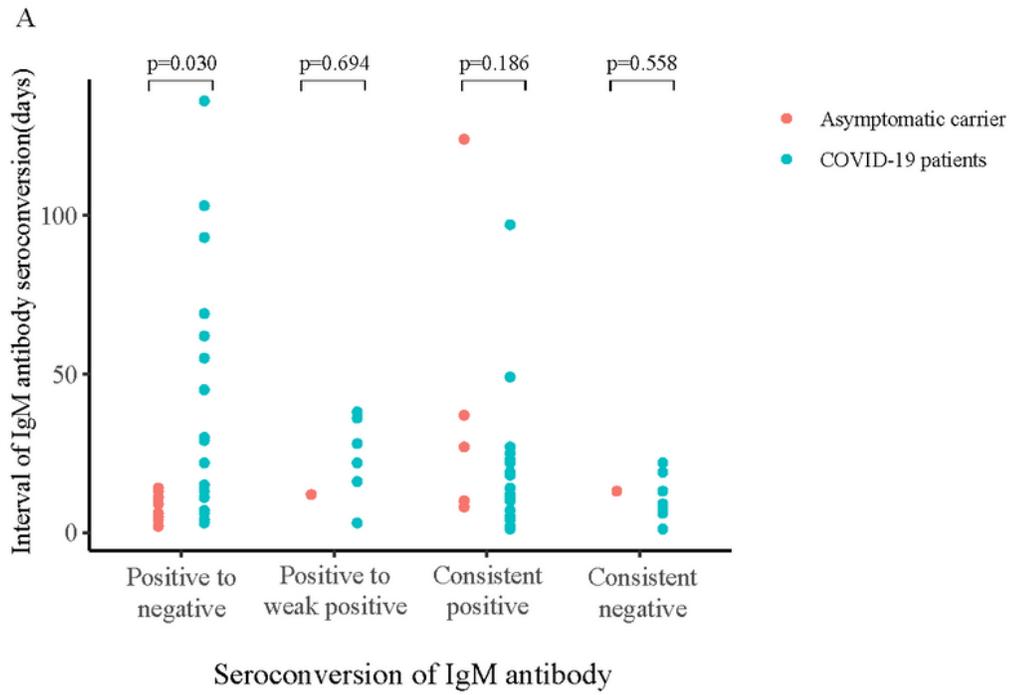


Figure 1

Proportion of patients owing to intra-familial transmission among asymptomatic carriers with SARS-CoV-2 infection and COVID-19 patients

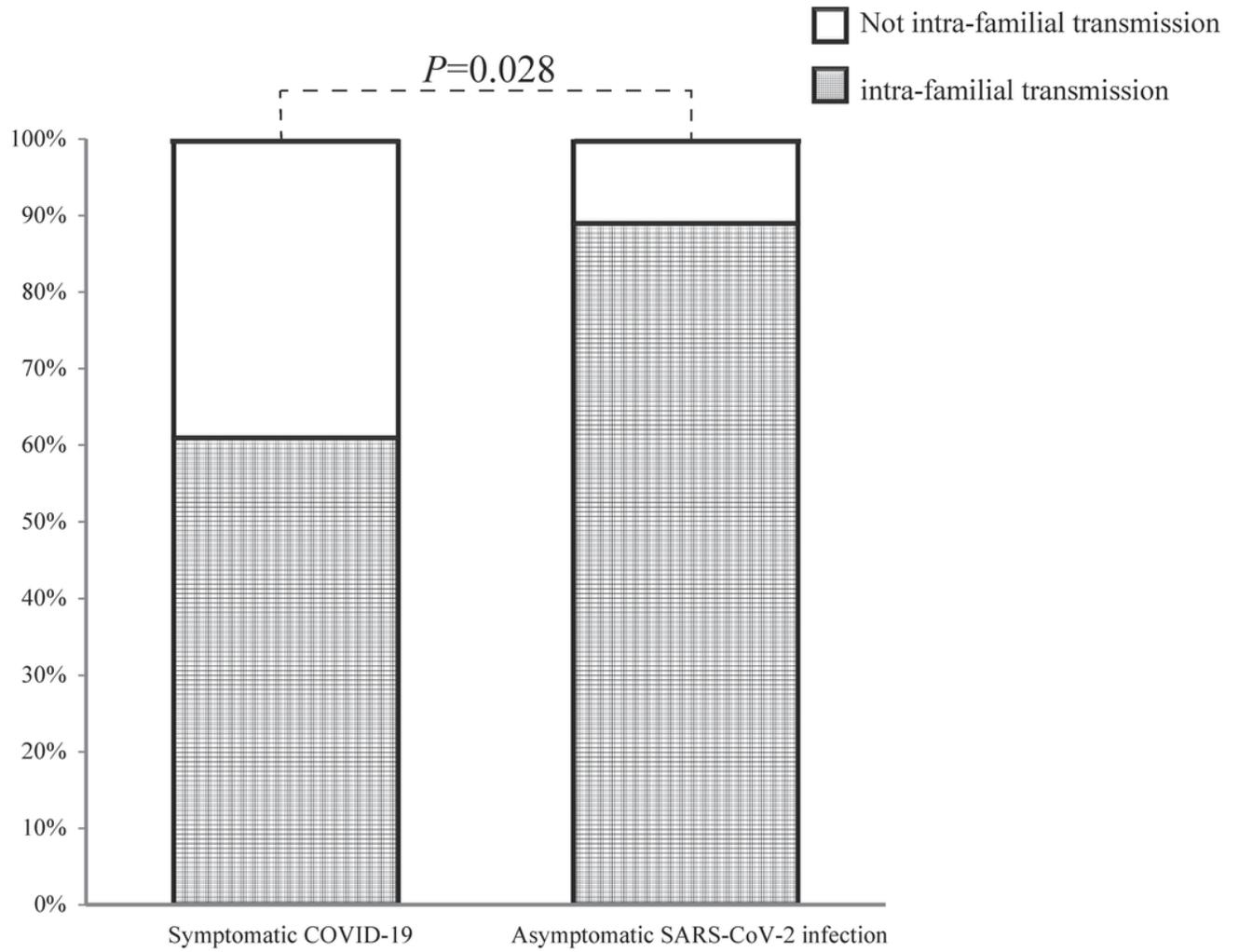


Figure 2

The distribution of time intervals of IgM and IgG antibody seroconversion among asymptomatic carriers with SARS-CoV-2 infection and COVID-19 patients during follow-up time