Check for updates

OPEN ACCESS

EDITED BY Zhimin Tao, Jiangsu University, China

REVIEWED BY Dawei Cui, Zhejiang University School of Medicine, China Fei Xiao, Peking University, China

*CORRESPONDENCE Zujiang Yu johnyuem@zzu.edu.cn Zhigang Ren fccrenzg@zzu.edu.cn

[†]These authors have contributed equally to this work

SPECIALTY SECTION

This article was submitted to Infectious Diseases – Surveillance, Prevention and Treatment, a section of the journal Frontiers in Medicine

RECEIVED 16 May 2022 ACCEPTED 14 September 2022 PUBLISHED 12 October 2022

CITATION

Li Q, Liu X, Li L, Hu X, Cui G, Sun R, Zhang D, Li J, Li Y, Zhang Y, Shen S, He P, Li S, Liu Y, Yu Z and Ren Z (2022) Comparison of clinical characteristics between SARS-CoV-2 Omicron variant and Delta variant infections in China. *Front. Med.* 9:944909. doi: 10.3389/fmed.2022.944909

COPYRIGHT

© 2022 Li, Liu, Li, Hu, Cui, Sun, Zhang, Li, Li, Zhang, Shen, He, Li, Liu, Yu and Ren. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

Comparison of clinical characteristics between SARS-CoV-2 Omicron variant and Delta variant infections in China

Qinggang Li^{1,2†}, Xiaorui Liu^{1,2†}, Lei Li^{1,2†}, Xiaobo Hu^{1,2}, Guangying Cui^{1,2}, Ranran Sun^{1,2}, Donghua Zhang³, Juan Li³, Yonghong Li³, Yong Zhang³, Shen Shen^{1,2}, Ping He^{1,2}, Shasha Li^{1,2}, Yanmin Liu^{1,2}, Zujiang Yu^{1,2*} and Zhigang Ren^{1,2*}

¹Department of Infectious Diseases, The First Affiliated Hospital of Zhengzhou University, Zhengzhou, China, ²Precision Medicine Center, Gene Hospital of Henan Province, The First Affiliated Hospital of Zhengzhou University, Zhengzhou, China, ³Anyang City Fifth People's Hospital, Anyang, China

Background: The continued 'evolution' of Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) has led to the emergence of the Omicron variant after the Delta variant, resulting in a significant increase in the number of people with COVID-19. This increase in the number of cases continues to have a significant impact on lives. Therefore, a more detailed understanding of the clinical characteristics of Omicron infection is essential.

Methods: Using medical charts, we extracted clinical information for 384 patients infected with the Omicron variant in Anyang City, Henan Province, China. Epidemiology and clinical characteristics were compared with a cohort of people infected with the Delta variant in Zhengzhou in 2021.

Findings: Common initial symptoms at onset of illness were cough [240 (63%)], expectoration [112 (29%)], fever [96 (25%)], nasal congestion [96 (25%)] and myalgia or fatigue [30 (6%)]. In patients with the Omicron variant, levels of total cholesterol, low-density lipoprotein and creatinine increased in 52 (14%), 36 (9%) and 58 (15%) patients, respectively, compared with patients with the Delta variant [one (1%), one (1%) and two (2%)]. Levels of triglyceride and high-density lipoprotein also increased. In patients with the Omicron variant, the levels of specific gravity and the erythrocyte sedimentation rate were increased in 115 (30%) and 81 (21%) patients, and serum levels of complement 3 decreased in 93 (41%).

Results: Compared with patients infected with Delta, no major differences in initial clinical symptoms were identified in patients infected with Omicron.

However, dyslipidemia and kidney injury were much more severe in patients with the Omicron variant, and the erythrocyte sedimentation rate was increased. Due to decreased levels of complement 3, the immunity of patients with the Omicron variant was weak.

KEYWORDS

coronavirus disease 2019 (COVID-19), Omicron variant, Delta variant, severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), clinical characteristics

Introduction

Coronavirus disease 2019 (COVID-19) caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) was first announced as a pandemic on March 11, 2020 (1). The COVID-19 pandemic has resulted in significant morbidity and mortality in both developing and developed countries and has had a huge impact on the economies and livelihoods of global society (2-4). SARS-CoV-2 is a coronavirus with the natural capacity to undergo mutation and antigenic variation over time (5). For example, it has been reported that the increased transmissibility of the Delta variant is associated with higher viral loads, longer duration of infection, and high rates of reinfection because of its high immune escape ability (6-8). On November 26, 2021, a novel variant was reported after being named Omicron (B.1.1.529) and identified by the World Health Organization (WHO) as a variant of concern (VOC) (9). It was first identified in South Africa in November 2021, and early studies showed that infection with Omicron is associated with significantly reduced severity and mortality compared with COVID-19 caused by previous variants (10, 11). Serum from people vaccinated with two doses of mRNA or vector vaccine enabled neutralization of Omicron variant to a lesser extent than of the Delta variant (12). Overall, the Omicron variant is probably more infectious than the Delta variant (13). Compared with the original wild-type strain of SARS-CoV-2, Omicron carries up to 30 single point mutations, three deletion mutations, and one insertion mutation of the spike protein, which is the target of most therapeutic antibodies and COVID-19 vaccines (14). This high variability is reflected in diverse behaviors, with the Omicron variant showing antibody evasion, vaccine resistance, and enhanced transmission (15, 16). Furthermore, available evidence suggests that recovered patients remain susceptible to Omicron, with the occurrence of immune escape and breakthrough reinfection (17, 18). However, current studies have demonstrated that the viremic effect of Omicron is milder than previous variants or the original virus (19, 20).

As of May 9, 2022, more than 515 million individuals had laboratory-confirmed SARS-CoV-2 Omicron VOC infection globally; there have also been more than 6.25 million deaths, and these numbers continue to rise rapidly (21). Therefore, greater knowledge of SARS-CoV-2 Omicron infection is urgent.

On January 8, 2022, two patients were confirmed to be infected with SARS-CoV-2 in Anyang, Henan Province, China, and a new wave of COVID-19 infections ensued. Wholegenome sequencing of the virus showed that both local cases involved infection with the Omicron variant strain BA.1 branch. After epidemiological investigation and genetic sequencing, the first outbreak in Henan was found to be homologous to that in Tianjin. To thoroughly identify potential population infections through multiple nucleic acid tests and close contact tracing and isolation, a group of patients were diagnosed with Omicron variant infection by laboratory confirmation and sent to designated hospitals for centralized treatment. Hence, these patients provide an essential opportunity to understand the clinical characteristics, laboratory findings, treatment and prognosis of the Omicron variant.

We also collected clinical statistics for patients infected with the Delta variant in October 2021. In our study, patients infected with the Delta variant in 2021 and the Omicron variant of the present outbreak were compared to explore differences in clinical characteristics to better understand the characteristics of the SARS-CoV-2 Omicron variant.

Materials and methods

Data sources

On January 8, 2022, the first two cases of Omicron variant infection were reported in Henan Province, China, and given the strongly contagious and insidious nature of the variant, local nucleic acid tests were promptly carried out for the entire population. The Omicron variant had already spread insidiously for at least three generations by the time the first case was reported, causing a faster and more widespread spread. Case definitions of confirmed SARS-CoV-2 infection were in accordance with the guidelines of

Abbreviations: COVID-19, coronavirus disease 2019; SARS-CoV-2, severe acute respiratory syndrome coronavirus 2; HDL, high-density lipoprotein; CxLDL, low-density lipoprotein; OxLDL, oxidized LDL.

30 November 2021.

the National Health Commission of the People's Republic of China. A total of 384 individuals infected with the Omicron variant admitted to the Fifth People's Hospital of Anyang from January 8 to February 13 were enrolled as the Omicron variant cohort. We collected information on patients admitted to the designated treatment hospital with laboratory-confirmed SARS-CoV-2 Omicron variant infection, including dates of illness onset, close contact with confirmed or suspected Omicron variant cases, and hospital admissions. Epidemiological data were collected through brief interviews with each patient. Clinical data for 103 patients infected with the Delta variant were collected in Zhengzhou from 30 July 2021 to

This study was approved by the Institutional Review Board of the First Affiliated Hospital of Zhengzhou University (L2021-Y429-002). The study was implemented in accordance with the Helsinki Declaration and Rules of Good Clinical Practice. Due to the urgent need to collect data on this emerging pathogen, the requirement for informed consent was waived. By extracting the medical records of patients, we used a standard case report form to gather clinical data. If the details were unclear, the working group in Zhengzhou contacted the doctors responsible for patient treatment for clarification.

Laboratory confirmation

Cases of COVID-19 were diagnosed by detecting SARS-CoV-2 RNA in a combined nasopharyngeal and oropharyngeal swab or sputum by real-time reverse transcription-polymerase chain reaction (RT-PCR). The collection of nasopharyngeal swabs was performed by well-trained medical staff in the same hospital, and the standardized procedures were strictly followed. The samples were stored in virus medium. Viral RNA was extracted within 2 h using the Q1Aamp Viral RNA Mini Kit (Beijing Kinghawk Pharmaceutical Co., Ltd.) according to the manufacturer's instructions. RT-PCR was performed by using the RNA Detection Kit for SARS-CoV-2 (Beijing Kinghawk Pharmaceutical Co., Ltd.) subsequently. RT-PCR was conducted with specific (ORF1ab-F: TGGTACTGGTCAGGCAATAAC; primers ORF1ab-R: TGATCTATGTGGCAACGGC; N-F: GACCCCA AAATCAGCGAAATG; N-R: CCACTGCGTTCTCCATTCTG) and Taqmen probes (ORF1ab-P: CTTTGGTGGTGCAT CGTGTTGTCT; N-P: TGCCAGTTGAATCTGAGGGTCCAC) targeting at the N, ORF1a/b genes and a positive reference gene. Reaction system and amplification conditions were performed according to the manufacturer's specification (Beijing Kinghawk Pharmaceutical Co., Ltd.).

The detection limit of cycle threshold (Ct) was set to be 38.0 (500 copies/ml). Samples with Ct of less than 38.0 were considered positive. The cut-off Ct value of 38.0 was determined *via* the receiver operating characteristics (ROC) curve method.

All tests were performed under strict biosafety conditions and the standard operating procedures.

According to sequencing viral RNA gene and comparing the results with known viral reference sequences, it was found that all patients were infected with Omicron variant. Laboratory tests, together with a whole blood count and serum biochemistry, were conducted on admission to the hospital.

Definitions

The date of diagnosis was defined as the date of first positive nasopharyngeal swab collection. The incubation period was defined as the date from exposure to the onset of illness, which was estimated for patients who could provide the exact date of close contact with individuals with confirmed or suspected SARS-CoV-2 Omicron variant infection. Patients were discharged when their respiratory symptoms and chest CT images improved and when nasopharyngeal and oropharyngeal swab specimens collected at least 24 h apart were negative by two consecutive real-time reverse transcription-polymerase chain reaction tests.

Statistical analysis

Categorical variables are described as percentages. For continuous variables, we calculated either means and standard deviations or medians with interquartile ranges. Means of continuous variables were compared using independent Group t tests when the data were normally distributed; otherwise, the Mann–Whitney test was used. Proportions for categorical variables were compared using the chi-square test; Fisher's exact test was used when the data were limited. Differences were statistically significant when the 2-sided P was less than 0.05. SPSS software for Windows, version 26 was used for statistical analysis.

Results

Epidemiological characteristics

Clinical data for 384 patients in Henan Province with laboratory-confirmed SARS-CoV-2 Omicron variant infection were gathered up to February 13, 2022. The mean age among these patients was 26.4 years (SD \pm 0.9 years; **Table 1**); it was 41.2 years (\pm 1.9) in patients infected with the Delta variant. Compared with patients infected with the Delta variant, the majority of patients in the current cohort were adolescents, accounting for 235 (61%) patients. Less than half of the 384 patients (162, 42%) were men. One hundred and sixty-six (46%) cases were associated with family clusters, which was notably

	Omicron variant (n = 384)	Delta variant (<i>n</i> = 103)	Р
Age (Mean \pm SD)	26.4 ± 0.9	41.2 ± 1.9	0
Age groups (years)			0
≤ 18	235 (61%)	11 (11%)	
	72 (19%)	40 (39%)	
41-65	58 (15%)	39 (38%)	
≥ 66	19 (5%)	13 (13%)	
— Male sex	162 (42%)	45 (44%)	0.784
Familial cluster	166 (46%)	28 (27%)	0.00
Number of vaccination doses			0
0	2 (1%)	40 (39%)	
1	11 (3%)	7 (7%)	
2	346 (90%)	53 (51%)	
3	21 (5%)	1 (1%)	
Coexisting conditions			
Any	40 (10%)	24 (23%)	0.00
Hypertension	23 (6%)	12 (12%)	0.04
Diabetes	10 (3%)	14 (14%)	0
Cardiovascular disease	6 (1.6%)	8 (8%)	0.00
Pregnancy	1 (0.3%)	1 (1%)	0.37
Malignancy	1 (0.3%)	0	1
Renal diseases	1 (0.3%)	1 (1%)	0.37
Liver disease	6 (1.6%)	3 (3%)	0.40
Fever	96 (25%)	36 (35%)	0.03
Highest temperature, °C			0.08
< 37.3	283 (74%)	67 (65%)	
37.3-38	48 (12%)	17 (17%)	
> 38	53 (14%)	19 (18%)	
Cough	240 (63%)	26 (25%)	0
Expectoration	112 (29%)	16 (16%)	0.00
Myalgia or fatigue	30 (6%)	9 (9%)	0.002
Nasal congestion	96 (25%)	7 (7%)	0
Loss of smell and taste	4 (1%)	2 (2%)	0.81
Headache	7 (2%)	3 (3%)	0.76
Diarrhea	6 (1.6%)	4 (4%)	0.27
Dyspnea	3 (0.8%)	0	1
Incubation period (days)	2(1-2)(n=97)	3 (2–7) (<i>n</i> = 81)	0
Time from illness onset to first hospital admission (days)	2 (1–3) (<i>n</i> = 362)	2.5 (0-5.0)	0.840
Systolic pressure, mm Hg	120 (110–125)	122 (113–130)	0.002
Heart rate	80 (75-82)	78.0 (75.0-81.3)	0.42
Respiratory rate	19 (18–20)	19 (18–20)	0.27
Oxygen saturation	99 (98–99)	97.0 (96.3-98.0)	0

TABLE 1 Personal and clinical baseline characteristics of patients infected with the Omicron variant or Delta variant.

Percentages do not total 100% owing to missing data.

Data are Mean \pm SD, median (IQR), n (%), or n/N (%), where N is the total number of patients with available data.

P comparing Omicron variant and Delta variant are from χ^2 , Fisher's exact test, or Mann–Whitney U test.

different from the Delta variant cases [28 (27%)]. It is significant that 367 (95%) patients infected with the Omicron variant received at least two doses of a COVID-19 vaccine, compared with 101 (97%) patients with the Delta variant. Moreover, 97 (25%) patients could provide the exact date of close contact

with someone who was confirmed or suspected to have a SARS-CoV-2 Omicron variant infection.

Clinical features

Forty of the 384 (10%) patients had one or more coexisting medical conditions: 23 (6%) had hypertension, ten (3%) had diabetes, six (1.6%) had cardiovascular disease and six (1.6%) had liver disease; pregnancy, malignancy and renal disease occurred in one patient (0.3%) each (Table 1). In patients infected with the Delta variant, the most common coexisting conditions were diabetes [14 (14%)], hypertension [12 (12%)] and cardiovascular disease [eight (8%)]. The most common initial symptoms at onset of illness were cough [240 (63%)], expectoration [112 (29%)], fever [96 (25%)], nasal congestion [96 (25%)] and myalgia or fatigue [30 (6%)]. Less common symptoms were headache, loss of smell and taste and diarrhea. Only three (0.8%) patients developed dyspnea (Table 1). No major variations in initial clinical symptoms were identified among Omicron patients.

Among 97 patients who could provide the exact date of close contact with someone with confirmed or suspected Omicron variant infection, the median incubation period from exposure to symptoms was 2 days (IQR 1–2 days). According to our data, the incubation period in patients infected with the Omicron variant was significantly shorter than that in those infected with the Delta variant. The median time from illness onset to first hospital admission was also 2 days (1–3).

Vital signs and laboratory parameters

Heart rate and respiratory rate did not differ between patients infected with the Omicron variant and with the Delta variant. Compared with patients infected with Delta, systolic pressure in patients with Omicron was reduced, though oxygen saturation was increased, with both within the normal range (Table 1). These measures were recorded for all patients on the day of hospital admission. The blood counts showed leukopenia (white blood cell count less than 4 \times 10⁹/L; 87 [23%] of 384 patients; Table 2). Compared with Deltainfected patients, the lymphocyte count, platelet count and hemoglobin were increased in Omicron-infected patients, and the neutrophil count was decreased. The proportion of patients with lymphopenia (lymphocyte count $< 1.0 \times 10^9$ /L; 20 [5%] patients) was lower in the 384 patients with Omicron than in the 103 patients with Delta (15 [15%] patients). The activated partial thromboplastin time and D-dimer level were lower in Omicron cases [median activated partial thromboplastin time 30.9 s (IQR 28.7-32.9); median D-dimer level 0.04 mg/L (0.02-0.09)] than in Delta cases [33.2 s (28.9-37.3); 0.33 mg/L (0.19-0.60)]. Compared with Delta cases [12 (12%)], levels

	Omicron variant (<i>n</i> = 384)	Delta variant (<i>n</i> = 103)	Р
White blood cell count, × 10 ⁹ per L	4.91 (4.06-6.22)	4.98 (4.00-6.21)	0.935
< 4	87 (23%)	25 (24%)	0.729
4-10	293 (76%)	76 (74%)	
> 10	4 (1%)	2 (2%)	
Neutrophil count, $ imes 10^9$ per L	2.43 (1.82-4.33)	2.72 (2.05-3.91)	0.011
Lymphocyte count, $\times 10^9$ per L	1.89 (1.49–2.41)	1.54 (1.23–2.10)	0
< 1.0	20 (5%)	15 (15%)	0.001
Eosinophil count, × 10 ⁹ per L	0.08 (0.04-0.15)	0.08 (0.03-0.14)	0.244
Platelet count, $\times 10^9$ per L	205 (173-241.5)	164 (119–204)	0
> 300	18 (5%)	4 (4%)	0.935
Hemoglobin, g/L	143 (131–159)	128 (119–146)	0
Prothrombin time, s	11.7 (10.9–12.5)	11.8 (11.4–12.4)	0.071
Activated partial thromboplastin time, s	30.9 (28.7-32.9)	33.2 (28.9–37.3)	0
> 37	15 (4%)	22 (21%)	0
D-dimer, mg/L	0.04 (0.02-0.09)	0.33 (0.19-0.60)	0
≥ 1.0	2 (0.5%)	10 (10%)	0
Alanine aminotransferase, U/L	17 (12–26)	17 (12–29)	0.699
> 40	39 (10%)	26 (25%)	0.472
Aspartate aminotransferase, U/L	19 (15–24)	23 (17–30)	0
> 40	17 (4%)	12 (12%)	0.006
Albumin, g/L	43 (41.1–45.1)	42.68 (40.16-45.60)	0.203
A/G	1.6 (1.4–1.9)	1.7 (1.4–1.9)	0.186
< 1.5	123 (32%)	26 (25%)	0.184
Γotal Bilirubin, μmol/L	11.2 (8.48–15.03)	8.5 (6.0–12.4)	0
Total cholesterol, mmol/L	4.58 (3.99-5.21)	3.86 (3.45-4.25)	0
≥ 5.72	52 (14%)	1 (1%)	0
Triglyceride, mmol/L	0.83 (0.64–1.19)	1.14 (0.91–1.60)	0
> 2.26	11 (3%)	4 (4%)	0.833
High-density lipoprotein, mmol/L	1.48 (1.26–1.72)	1.05 (0.91–1.23)	0
< 0.91	4 (1%)	23 (22%)	0
Low-density lipoprotein, mmol/L	2.64 (2.31–3.11)	1.93 (1.64–2.29)	0
≥ 3.64	36 (9%)	1 (1%)	0.004
Creatinine, µmol/L	69 (62–78)	59.5 (49–71.5)	0
≥ 133	58 (15%)	2 (2%)	0.044
Creatinine kinase, U/L	86 (63–116)	61.5 (38.7–91.7)	0
Blood urea nitrogen, μ mol/L	3.4 (3.2-4.9)	4.44 (3.82–5.21)	0.012
Lactate dehydrogenase, U/L	158 (140–180)	200 (173-243)	0
≥ 250	8 (2%)	25 (24%)	0

TABLE 2 Laboratory findings of patients infected with the Omicron variant or Delta variant on admission to the hospital.

(Continued)

TABLE 2 (Continued)
-----------	------------

	Omicron variant (<i>n</i> = 384)	Delta variant (<i>n</i> = 103)	Р
Specific gravity			
≥ 1.030	115 (30%)	NA	
Erythrocyte sedimentation rate, mm/h	13 (8–20)		
> 20	81 (21%)	NA	
Complement 3	0.73 (0.61 - 0.88) (n = 225)	NA	
< 0.7	93 (41%)	NA	
Complement 4	0.31 (0.25 - 0.44) (<i>n</i> = 225)	NA	
C-reactive protein, mg/L	1.57 (0.14-4.20)	3.9 (1.5-18.5)	0
> 10	28 (7%)	36 (35%)	0
Procalcitonin, ng/mL			
< 0.1	314 (82%)	95 (92%)	0.01
Pneumonia	116 (30%)	63 (61%)	0
Bilateral involvement of chest radiographs	51 (13%)	66 (50%)	0

Percentages do not total 100% owing to missing data.

Data are median (IQR), n (%), or n/N (%), where N is the total number of patients with available data.

P comparing Omicron variant and Delta variant are from $\chi^2,$ Fisher's exact test, or Mann–Whitney U test.

of aspartate aminotransferase were increased in 17 (4%) of the 384 Omicron cases. Moreover, some of the patients with the Omicron variant developed dyslipidemia. Levels of total cholesterol, low-density lipoprotein and creatinine in patients with the Omicron variant were increased in 52 (14%), 36 (9%) and 58 (15%) cases, respectively, and in one [1%], one [1%] and two [2%] Delta variant cases, respectively (**Figures 1A,C**). Compared with patients infected with Delta, levels of triglyceride and high-density lipoprotein were increased in those infected with Omicron, and lactate dehydrogenase was increased in eight (2%) (**Figures 1B,C**). In patients infected with the Omicron variant, levels of specific gravity and erythrocyte sedimentation rate were increased in 115 (30%) and 81 (21%) cases, respectively.

On admission, serum levels of complement 3 were decreased in 93 (41%) of 225 Omicron patients. Most patients [314 (82%)] had normal serum levels of procalcitonin, and a few [28 (7%)] had abnormal levels of C-reactive protein. On admission, 116 (30%) of the 384 enrolled patients infected with Omicron developed pneumonia, and only 51 (13%) showed bilateral involvement of chest CT scans (Figure 2).

Clinical treatment and outcomes

The complications, treatment and outcomes of the 384 patients infected with Omicron are shown in Table 3.



Lipid laboratory tests between patients infected with the Omicron variant or Delta variant on admission to the hospital. (A) Comparison of total cholesterol in patients infected with the Omicron or Delta variant. (B) Comparison of triglyceride in patients infected with the Omicron or Delta variant. (C) Comparison of high-density lipoprotein in patients infected with the Omicron or Delta variant. (D) Comparison of low-density lipoprotein in patients infected with the Omicron or Delta variant. The asterisk indicates a statistically significant difference compared with delta patients. *** means P < 0.001.

None of these patients was transferred to an intensive care unit. Complications among the 384 patients included acute kidney injury [1 (0.3%)], anemia [1 (0.3%)] and secondary infection [1 (0.3%)]. Almost all patients [381 (99%)] received traditional Chinese medicine treatment, and 143 (37%) received immunotherapy. Only a few patients received heparin [33 (9%)], interferon inhalation [22 (6%)] or antiviral treatment [8 (2%)], with a lower proportion of use than in patients infected with the Delta variant. Furthermore, nasal cannula was required in only four (1%) patients.

As of April 10, 2022, all patients had been discharged from the hospital, with no deaths. In patients infected with the Omicron variant, the median time from first hospital admission to discharge was 19 days (interquartile range 15–25 days), which was shorter than that in patients with the Delta variant [26 (18–46) days]. Disease duration, the time from illness onset to discharge, was lengthened [22 (17–28) days] in patients with Omicron compared with those with Delta [28 (21–47) days].

Discussion

As of 9 May 2022, over 220,000 laboratory-confirmed cases of infection with SARS-CoV-2 have been reported in China (22).



Representative chest CT images. (A) Transverse chest CT images for a 16-year-old female infected with Omicron only showing ground-glass opacity on day four after symptom onset. (B) Transverse chest CT images for a 35-year-old female infected with Omicron showing signs of bilateral pneumonia on day four after symptom onset. (C) Transverse chest CT images for a 51-year-old male infected with Omicron showing signs of bilateral pneumonia on day three after symptom onset. (D) Transverse chest CT images for a 73-year-old male infected with Omicron showing multiple bilateral pneumonia on day six after symptom onset.

According to a report from the WHO, the current global epidemiology of SARS-CoV-2 is characterized by continued rapid global spread of the Omicron variant. In previous studies, patients infected with the Omicron variant were less likely to be hospitalized and develop severe disease than those infected with previous variants or the original virus causing COVID-19 (23). Here, we report a cohort of 384 patients with laboratoryconfirmed SARS-CoV-2 Omicron variant infection. Based on epidemiological investigation, 378 (98%) of the patients in our study had received at least two doses of COVID-19 vaccination, indicating immune escape by the Omicron variant. Common initial symptoms at onset of illness were cough, expectoration, fever, nasal congestion, myalgia and fatigue, consistent with the original wild-type strain of SARS-CoV-2 and other variant infections (19, 24). Most of the patients in Henan Province had mild symptoms, and only three patients developed dyspnea. No

	Omicron variant (n = 384)	Delta variant (<i>n</i> = 103)	Р
Admission to intensive care unit	0	10 (10%)	0
Acute kidney injury	1 (0.3%)	0	1
Anemia	1 (0.3%)	0	1
Secondary infection	1 (0.3%)	0	1
Shock	0	1 (1%)	0.211
Treatment			
TCM treatment	381 (99%)	93 (90%)	0
Immunotherapy	143 (37%)	35 (34%)	0.542
Antiviral treatment	8 (2%)	16 (16%)	0
Antibiotics	11 (3%)	16 (16%)	0
Interferon inhalation	22 (6%)	13 (13%)	0.016
Heparin	33 (9%)	26 (25%)	0
Chloroquine phosphate	0	3 (3%)	0.008
Glucocorticoid treatment	0	0	
Continuous renal-replacement therapy	0	0	
Oxygen support			
Nasal cannula	4 (1%)	17 (17%)	0
Non-invasive ventilation or high-flow nasal cannula	0	2 (2%)	0.044
Invasive mechanical ventilation	0	2 (2%)	0.044
Invasive mechanical ventilation and ECMO	0	2 (2%)	0.044
Prognosis			
Hospitalization	0	0	
Discharge	384 (100%)	103 (100%)	
Death	0	0	
Hospital stay, days	19 (15–25)	26 (18-46)	0
Disease duration, days	22 (17–28) (<i>n</i> = 362)	28 (21–47) (<i>n</i> = 101)	0

TABLE 3 Treatments and outcomes of patients infected with the Omicron variant or Delta variant.

Percentages do not total 100% owing to missing data.

Data are median (IQR), n (%), or n/N (%), where N is the total number of patients with available data.

P comparing Omicron variant and Delta variant are from χ^2 , Fisher's exact test, or Mann–Whitney U test.

patients were admitted to the intensive care unit. Compared with patients infected with the Delta variant, laboratory findings showed that the patients with the Omicron variant experienced mild illness. Our results showed that most Omicron patients had levels of white blood cell counts, neutrophil counts, and lymphocyte counts in the normal range. This suggests an active interaction between immunity and the Omicron variant (19). Compared with those infected with Delta, levels of activated partial thromboplastin time were increased in patients infected with Omicron, which may be related to the increased activity of coagulation factors. There were a few patients with abnormal alanine aminotransferase, aspartate aminotransferase, lactate dehydrogenase, C-reactive protein or procalcitonin, which suggests little hepatic injury, little myocardial injury and a weak inflammatory response. Compared with patients infected with the Delta variant, the incidence of pneumonia and bilateral involvement of chest radiographs were reduced in patients infected with the Omicron variant, resulting in little pulmonary injury. Omicron infections generally cause less severe disease than infections with prior variants (25). Coexisting conditions including hypertension, diabetes and cardiovascular diseases demonstrated lower frequencies in patients with Omicron than those with Delta, which might be also associated with their younger ages. Furthermore, the significant age difference between the two cohorts may have affected the laboratory results, leading to fewer patients with mild infections than among patients infected with Delta.

Compared with those infected with Delta, levels of high-density lipoprotein were increased in patients infected with Omicron. It has been reported that low high-density lipoprotein and high triglyceride concentrations before or during hospitalization are strong predictors of COVID-19 severity (26). A higher proportion of patients with the Omicron variant had abnormally elevated total cholesterol and lowdensity lipoprotein than in patients with Delta. Dyslipidemia is a potential risk factor for COVID-19 (27). Studies have shown that Dyslipidaemia was associated with the severity and mortality of COVID-19 (28). A sustained inflammatory response driven by a 'cytokine storm', including the release of pro-inflammatory cytokines (TNF-α, IL-6, IL-8 and IL-10) and lymphopenia, is thought to be a major cause of life-threatening complications in SARS-CoV-2 patients. In the context of acute inflammation, reduced plasma levels of lecithin cholesterol acyltransferase (LCAT) may also alter HDL function and further worsen the inflammatory response (29). At the same time, LDL and its major apolipoprotein, apolipoprotein B (apoB), are oxidized (oxLDL) (30). On admission, levels of creatinine and specific gravity were increased in patients with Omicron, which indicates that Omicron variant infection might be associated with kidney injury. Similarly, a high erythrocyte sedimentation rate indicates the active stage of the disease. In addition, low serum levels of complement 3 in Omicron patients indicated reduced immunity.

Almost all patients in the two cohorts received traditional Chinese medicine treatment, but the types of drugs used varied. Because of the reduced immunity in patients with the Omicron variant, 143 (37%) received immunotherapy. In addition, the use of antiviral drugs, antibiotics and heparin was decreased in patients with the Omicron variant compared with those infected with the Delta variant. Both disease duration and hospital stay were reduced in Omicron patients. These results indicate that SARS-CoV-2 Omicron variant infection might be milder than Delta variant infection.

Given the large number of patients infected with Omicron in this study, our findings provide valuable information for understanding the clinical characteristics of Omicron variant infection. Due to the increased transmissibility and risk of reinfection of Omicron variant, it is still important to remain vigilant and protect ourselves against COVID-19.

There are several limitations in our study. First, as the patients were only from Henan Province, it is possible that more clinical features associated with Omicron variant infection were identified. Second, because of the strict control measures associated with COVID-19 in China, including epidemiological history investigation and follow-up, isolation of close contacts and early treatment of patients, patients with mostly mild disease were enrolled in our study.

Conclusion

No major differences were identified between initial clinical symptoms in patients infected with Omicron versus those with Delta. Compared with patients infected with the Delta variant, dyslipidemia and kidney injury were much more severe in patients infected with the Omicron variant, and the erythrocyte sedimentation rate was increased. Due to decreased levels of complement 3, the immunity of patients with the Omicron variant was reduced.

Data availability statement

The datasets presented in this article are not readily available because COVID-19 patient information is private. Requests to access the datasets should be directed to ZR, fccrenzg@zzu.edu.cn.

Author contributions

ZY and ZR designed the study. ZY, ZR, QL, XL, XH, GC, RS, DZ, JL, YHL, YZ, SS, PH, SL, and YML treated the patients

References

1. Acuti Martellucci C, Flacco ME, Cappadona R, Bravi F, Mantovani L, Manzoli L. SARS-CoV-2 pandemic: an overview. *Adv Biol Regul.* (2020) 77:100736. doi: 10.1016/j.jbior.2020.100736

2. Singh AK, Misra A. Impact of COVID-19 and comorbidities on health and economics: focus on developing countries and India. *Diabetes Metab Syndr.* (2020) 14:1625–30. doi: 10.1016/j.dsx.2020.08.032

3. Karadag E. Increase in COVID-19 cases and case-fatality and case-recovery rates in Europe: a cross-temporal meta-analysis. *J Med Virol.* (2020) 92:1511–7. doi: 10.1002/jmv.26035

4. Matta S, Chopra KK, Arora VK. Morbidity and mortality trends of Covid 19 in top 10 countries. *Indian J Tuberc*. (2020) 67:S167–72. doi: 10.1016/j.ijtb.2020.09.031

5. Awadasseid A, Wu Y, Tanaka Y, Zhang W. SARS-CoV-2 variants evolved during the early stage of the pandemic and effects of mutations on adaptation in Wuhan populations. *Int J Biol Sci.* (2021) 17:97–106. doi: 10.7150/ijbs.47827

and collected the clinical data. ZR and LL analyzed the clinical data. LL and ZR wrote the manuscript. All authors reviewed and approved the manuscript.

Funding

This study was sponsored by grants from Corona Virus Disease 2019 Chinese Medicine Emergency Special (2022ZYLCYJ01-3), National Key R&D Program of China (2018YFC2000501), and National Natural Science Foundation of China (U2004121, 82070643, and U1904164).

Acknowledgments

We thank all the generous volunteer subjects who participated in the study. We also thank all the doctors, nurses, and medical technicians who bravely fight the COVID-19 pandemic.

Conflict of interest

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

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

6. Wang Y, Chen R, Hu F, Lan Y, Yang Z, Zhan C, et al. Transmission, viral kinetics and clinical characteristics of the emergent SARS-CoV-2 Delta VOC in Guangzhou, China. *EClinicalMedicine*. (2021) 40:101129. doi: 10.1016/j.eclinm. 2021.101129

7. Liu C, Ginn HM, Dejnirattisai W, Supasa P, Wang B, Tuekprakhon A, et al. Reduced neutralization of SARS-CoV-2 B.1.617 by vaccine and convalescent serum. *Cell.* (2021) 184:4220–4236e13. doi: 10.1016/j.cell.2021.06.020

8. Hu Z, Huang X, Zhang J, Fu S, Ding D, Tao Z. Differences in clinical characteristics between delta variant and wild-type SARS-CoV-2 infected patients. *Front Med (Lausanne)*. (2021) 8:792135. doi: 10.3389/fmed.2021.792135

9. WHO. The Name of SARS-Cov-2 Variants and Currently Designated Variants of Concern. Geneva: WHO (2022).

10. Maslo C, Friedland R, Toubkin M, Laubscher A, Akaloo T, Kama B. Characteristics and outcomes of hospitalized patients in South Africa during the

COVID-19 omicron wave compared with previous waves. JAMA. (2022) 327:583-4. doi: 10.1001/jama.2021.24868

11. Wolter N, Jassat W, Walaza S, Welch R, Moultrie H, Groome M, et al. Early assessment of the clinical severity of the SARS-CoV-2 omicron variant in South Africa: a data linkage study. *Lancet.* (2022) 399:437–46. doi: 10.1016/s0140-6736(22)00017-4

12. Rössler ARL, Bante D, von Laer D, Kimpel J. SARS-CoV-2 omicron variant neutralization in serum from vaccinated and convalescent persons. *N Eng J Med.* (2022) 386:698–700. doi: 10.1056/NEJMc2119236

13. He X, Hong W, Pan X, Lu G, Wei X. SARS-CoV-2 Omicron variant: characteristics and prevention. *MedComm.* (2021) 2:838–45. doi: 10.1002/mco2. 110

14. GISAID. Tracking of Variants. Munich: GISAID (2022).

15. Yin WXY, Xu P, Cao X, Wu C, Gu C, He X, et al. Structures of the omicron spike trimer with ACE2 and an anti-omicron antibody. *Science*. (2022) 375:1048–53.

16. Liu L, Iketani S, Guo Y, Chan JF, Wang M, Liu L, et al. Striking antibody evasion manifested by the omicron variant of SARS-CoV-2. *Nature.* (2022) 602:676–81. doi: 10.1038/s41586-021-04388-0

17. Zhang X, Wu S, Wu B, Yang Q, Chen A, Li Y, et al. SARS-CoV-2 Omicron strain exhibits potent capabilities for immune evasion and viral entrance. *Signal Transduct Target Ther.* (2021) 6:430. doi: 10.1038/s41392-021-00852-5

18. Wu L, Zhou L, Mo M, Liu T, Wu C, Gong C, et al. SARS-CoV-2 Omicron RBD shows weaker binding affinity than the currently dominant Delta variant to human ACE2. *Signal Transduct Target Ther*. (2022) 7:8. doi: 10.1038/s41392-021-00863-2

19. Zhang J, Chen N, Zhao D, Zhang J, Hu Z, Tao Z. Clinical characteristics of COVID-19 Patients Infected by the omicron variant of SARS-CoV-2. *Front Med (Lausanne)*. (2022) 9:912367. doi: 10.3389/fmed.2022.912367

20. Kim MK, Lee B, Choi YY, Um J, Lee KS, Sung HK, et al. Clinical characteristics of 40 patients infected with the SARS-CoV-2 omicron variant in Korea. *J Korean Med Sci.* (2022) 37:e31. doi: 10.3346/jkms.2022.37.e31

21. WHO. Coronavirus (COVID-19) Dashboard. Geneva: WHO (2022).

22. Chinese Center for Disease Control and Prevention. *Update on the Novel Coronavirus Pneumonia Outbreak as at 24:00 on 9 May.* Beijing: Chinese Center for Disease Control and Prevention (2020).

23. Wang D, Hu B, Hu C, Zhu F, Liu X, Zhang J, et al. Clinical characteristics of 138 hospitalized patients with 2019 novel coronavirus-infected pneumonia in Wuhan, China. *JAMA*. (2020) 323:1061–9. doi: 10.1001/jama.2020.1585

24. Xu XW, Wu XX, Jiang XG, Xu KJ, Ying LJ, Ma CL, et al. Clinical findings in a group of patients infected with the 2019 novel coronavirus (SARS-Cov-2) outside of Wuhan, China: retrospective case series. *BMJ*. (2020) 368:m606. doi: 10.1136/bmj.m606

25. Huang C, Wang Y, Li X, Ren L, Zhao J, Hu Y, et al. Clinical features of patients infected with 2019 novel coronavirus in Wuhan, China. *Lancet.* (2020) 395:497–506. doi: 10.1016/s0140-6736(20)30183-5

26. Masana L, Correig E, Ibarretxe D, Anoro E, Arroyo JA, Jerico C, et al. Low HDL and high triglycerides predict COVID-19 severity. *Sci Rep.* (2021) 11:7217. doi: 10.1038/s41598-021-86747-5

27. Ribeiro HG, Dantas-Komatsu RCS, Medeiros JFP, Carvalho M, Soares VL, Reis BZ, et al. Previous vitamin D status and total cholesterol are associated with SARS-CoV-2 infection. *Clin Chim Acta*. (2021) 522:8–13. doi: 10.1016/j.cca.2021. 08.003

28. Liu Y, Pan Y, Yin Y, Chen W, Li X. Association of dyslipidemia with the severity and mortality of coronavirus disease 2019 (COVID-19): a meta-analysis. *Virol J.* (2021) 18:157. doi: 10.1186/s12985-021-01604-1

29. Khovidhunkit W, Shigenaga JK, Moser AH, Feingold KR, Grunfeld C. Cholesterol efflux by acute-phase high density lipoprotein: role of lecithin:cholesterol acyltransferase. *J Lipid Res.* (2001) 42:967–75. doi: 10.1016/s0022-2275(20)31621-7

30. Sorokin AV, Karathanasis SK, Yang ZH, Freeman L, Kotani K, Remaley AT. COVID-19-Associated dyslipidemia: implications for mechanism of impaired resolution and novel therapeutic approaches. *FASEB J.* (2020) 34:9843–53. doi: 10.1096/fj.202001451