

Disease profile and plasma neutralizing activity of post-vaccination Omicron BA.1 infection in Tianjin, China: a retrospective study

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Supplementary Information, Materials and Methods

Data Sources

This study analyzed clinical and demographic data on the 430 Omicron patients reported by Tianjin Municipal Health Commission between January 8 and February 7, 2022, including COVID-19 vaccination history, laboratory tests, clinical symptoms, ICU admission and death. These data were extracted from the medical records obtained from Tianjin Haihe Hospital and Tianjin First Central Hospital where the patients received treatment after diagnosis.

Laboratory Confirmation

In this study, laboratory diagnosis was done in two designated COVID-19 hospitals: Tianjin Haihe Hospital and Tianjin First Central Hospital. Nucleic acid was extracted from respiratory samples using commercial kits (Zybio, 5203050). Reverse Transcription-Polymerase Chain Reaction (RT-PCR) was performed following the WHO protocol to detect two target genes, the open reading frame of 1ab (ORF1ab) and the nucleocapsid protein (N).⁵ ORF1ab forward primer (F): CCCTGTGGGTTTTACTTAA, reverse primer (R): ACGATTGTGCATCAGCTGA, probe (P): 5'-FAM-CCGTCTGCGGTATGTGGAAAGGTTATGG-BHQ1-3'. N forward primer: GGGGAAGTTCTCCTGCTAGAAT, reverse primer: CAGACATTTTGCTCTCAAGCTG, probe: 5'-FAM-TTGCTGCTGCTTGACAGATT-TAMRA-3'. 40 cycles of amplification (50°C for 10 min, 90°C for 5 min, 95°C for 10 s; 55°C for 40 s) were performed. A cycle threshold value (Ct-value) less than 37 with an S-shape amplification curve was defined as positive, and a Ct-value no less than 40 without a S-shape amplification curve was defined as negative. In the case of $37 \leq Ct < 40$, retesting would be recommended. If both target genes (ORF1ab and N) are tested positive on real-time RT-PCR for the same sample, the result would be regarded as positive; if only one of the two target genes is tested positive, resampling and retesting would be required. If one of the two target genes is tested positive for two samples, the result would be positive.

Plasma Isolation

Peripheral blood was collected from 135 Omicron BA.1 convalescent patients 28 days after discharge and 114 healthy recipients of 3 doses of inactivated vaccines (CoronaVac) around 90 days after the booster shot. Whole blood samples were then diluted with PBS+2%FBS at 1:1 for Ficoll gradient centrifugation (Cytiva, 17-1440-03). Plasma was collected from the supernatants and restored at -80°C.

Authentic neutralization assay

A neutralization assay of authentic SARS-CoV-2 was performed using a cytopathic effect (CPE) assay. Briefly, various concentrations (2-fold serial dilution using DMEM) of plasma were mixed with the same volume of 100 PFU of authentic SARS-CoV-2 and incubated at 37°C for 1 h. The mixture was added to a monolayer of Vero-E6 cells (5x10³ cells per well) in a 96-well plate and incubated for 1 h at 37°C. The supernatant was removed, and 200 µL of DMEM supplied with 2% (v/v) FBS was added to the infected cells. After incubation at 37°C supplied with 5% CO₂ for 5 days, all wells were examined for the CPE effect. All experiments were performed in a Biosafety Level 3 facility. Neutralization titers were calculated by the Spearman-Kärber method.

Outcomes

The primary endpoints were COVID-19 severity and neutralizing antibody titers. COVID-19 disease severity was defined as asymptomatic, mild, moderate, severe and critical according to WHO living guidance for clinical management of COVID-19.

Secondary endpoints were intensive care unit (ICU) admission, re-positive results on nucleic acid tests during convalescence phase, and duration of hospitalization and recovery. PCR re-positive during convalescence phase was defined as PCR Ct value < 40 after two independent PCR-negative results at an interval of more than 24 hours. Duration of hospitalization and recovery was defined as the days spent in Tianjin Haihe Hospital and Tianjin First Central Hospital, respectively. All patients, including asymptomatic and mild cases, were hospitalized in Tianjin Haihe Hospital upon positive PCR results. Patients were discharged from Tianjin Haihe Hospital if the following criteria were met: 1) body temperature restored and stayed normal for over 3 days; 2) respiratory symptoms significantly relieved; 3) acute exudation substantially resolved on imaging study of the lungs; 4) negative on two consecutive PCR tests (at an interval of at least 24 hours) of samples collected from the respiratory tract. For patients whose PCR assays remained positive for over 4 weeks after criteria 1), 2) and 3) had been met, antibody assay and virus culture were applied to assess the risk of transmission before deciding whether these patients could be discharged.

Discharged patients from Tianjin Haihe Hospital were then transferred to Tianjin First Central Hospital for at least 14 days under medical observation. Patients received PCR assays on the 1st, 7th and 14th days after being transferred to Tianjin First Central Hospital. After 14 days of observation, patients with negative results on PCR and without other conditions in need of

hospitalization were discharged. Re-positive cases during convalescence phase were required to yield negative on consecutive PCR assays at an interval of at least 24 hours.

Study Oversight

This study was approved by the Tianjin Municipal Health Commission and the Ethics Committee of Tianjin First Central Hospital (Ethics committee archiving No. 2022N045KY). All patients/participants provided their written informed consent to have their clinical information collected for study purposes and the data generated from the study published. All the authors contributed to data collection and analysis, discussion and interpretation of the results. All the authors read and approved the final manuscript.

Statistical Analysis

Continuous variables were shown in medians and interquartile ranges (IQR), and the Mann-Whitney U test was used to analyze the differences between the two groups. Categorical variables were summarized as counts and percentages, and analyzed by Pearson's χ^2 test. Ordered multi-class logistic regression model was used to analyze the relations between age/gender/receipt of inactivated vaccine and COVID-19 severity. No imputation was made for missing data. All the analyses were conducted using the SPSS software, version 22.0. Two-sided $P < 0.05$ was used in tests of significance.

Role of the funding source

The funders of the study had no role in study design, data collection, data analysis, data interpretation, or writing of the report.

Data sharing

Although all data used in this analysis were anonymized, the individual level nature of the data used risks individuals being identified, or being able to self-identify, if the data are released publicly. Extracted data are available upon request.

Ethical statement

This study was performed in accordance with Declaration of Helsinki after approved by the Tianjin Municipal Health Commission, and the Ethics Committee of Tianjin First Central Hospital (Ethics committee archiving No. 2022N045KY). Informed consent was obtained from all human research participants.

References

- 1 Tianjin Municipal Health Commission. http://wsjk.tj.gov.cn/ZTZL1/ZTZL750/YQFKZL9424/FKDT1207/202201/t20220109_5774785.html 2022.
- 2 Tianjin Municipal Health Commission. Vaccination of COVID-19 in Tianjin. http://wsjktjgovcn/ZTZL1/ZTZL750/YQFKZL9424/wjwymjzqk/202201/t20220109_5774994html 2022.
- 3 Fu L, Wang B, Yuan T *et al.* Clinical characteristics of coronavirus disease 2019 (COVID-19) in China: A systematic review and meta-analysis. *Journal of Infection* 2020; **80**:656-665.
- 4 Tan L, Wang Q, Zhang D *et al.* Lymphopenia predicts disease severity of COVID-19: a descriptive and predictive study. *Signal transduction and targeted therapy* 2020; **5**:33.
- 5 Swanson PA, 2nd, Padilla M, Hoyland W *et al.* AZD1222/ChAdOx1 nCoV-19 vaccination induces a polyfunctional spike protein-specific TH1 response with a diverse TCR repertoire. *Science translational medicine* 2021; **13**:eabj7211.
- 6 Painter MM, Mathew D, Goel RR *et al.* Rapid induction of antigen-specific CD4(+) T cells is associated with coordinated humoral and cellular immunity to SARS-CoV-2 mRNA vaccination. *Immunity* 2021; **54**:2133-2142.e2133.
- 7 Madhi SA, Ihekweazu C, Rees H, Pollard AJ. Decoupling of omicron variant infections and severe COVID-19. *Lancet (London, England)* 2022; **399**:1047-1048.
- 8 Milne G, Hames T, Scotton C *et al.* Does infection with or vaccination against SARS-CoV-2 lead to lasting immunity? *The Lancet Respiratory medicine* 2021; **9**:1450-1466.
- 9 Servellita V, Syed AM, Morris MK *et al.* Neutralizing immunity in vaccine breakthrough infections from the SARS-CoV-2 Omicron and Delta variants. *Cell* 2022; **185**:1539-1548.e1535.
- 10 WHO. Living guidance for clinical management of COVID-19: living guidance. <https://apps.who.int/iris/handle/10665/349321> 2021.

Table S1. Initial symptoms and convalescent symptoms of Omicron infected patients *

Symptoms	Initial	Convalescent
Cough	159 (37.0)	36 (8.4)
Fever	130 (30.2)	13 (3.0)
Sore throat	83 (19.3)	24 (5.6)
Fatigue	55 (12.8)	21 (4.9)
Nasal congestion	48 (11.2)	20 (4.7)
Nasal discharge	38 (8.8)	19 (4.4)
Parageusia	6 (1.4)	22 (5.1)
Heterosmia	4 (0.9)	22 (5.1)
Diarrhea	4 (0.9)	21 (4.9)
Rash	1 (0.2)	24 (5.6)
Conjunctivitis	0 (0.0)	22 (5.1)
Mucosal inflammation	0 (0.0)	21 (4.9)
Hypotension	0 (0.0)	22 (5.1)

* Categorical variables were summarized as counts (percentages).

Table S2. Comorbidities of Omicron infected patients *

Disease		
Cardiovascular system	Hypertension	73 (17.0)
	Coronary artery disease	18 (4.2)
	Arrhythmia	9 (2.1)
	Other	4 (0.9)
Digestive system	Abnormal liver function	69 (16.0)
	Fatty liver	17 (4.0)
	Gastrointestinal diseases	12 (2.8)
	Gallstone	7 (1.6)
Endocrine system	Diabetes	31 (7.2)
	Electrolyte disorder	20 (4.7)
	Hyperlipidemia	19 (4.4)
	Thyroid dysfunction	12 (2.8)
Respiratory system	Other	11 (2.6)
	Lung disease	20 (4.7)
	Bronchiectasia	3 (0.7)
Nervous system	Cerebrovascular disease	9 (2.1)
	Viral encephalitis	1 (0.2)
	Malignant tumor of brain	1 (0.2)
Genitourinary system	Chronic nephritis	3 (0.7)
	Renal malignancy	1 (0.2)
Hematopathy	Anemia	11 (2.6)
	Leucopenia	4 (0.9)
	Thrombocytopenia	3 (0.7)
Dermatosis	Leukemia	1 (0.2)
		3 (0.7)
Other		14 (3.3)

* Categorical variables were summarized as counts (percentages).

Table S3. Characteristics of Omicron infected patients *

Characteristic	All patients (n = 430)	Vaccination		P-value	Population		P-value
		Vaccinated (n = 392)	Unvaccinated (n = 38)		Adult (n = 316)	Child † (n = 114)	
Age — yr	36 (14-55)	36 (16-54.5)	30.5 (3-64)	0.616	47 (34-58)	10 (8-11)	< 0.001
Gender				0.325			0.162
Male	191 (44.4)	177 (45.2)	14 (36.8)		134 (42.4)	57 (50.0)	
Female	239 (55.6)	215 (54.8)	24 (63.2)		182 (57.6)	57 (50.0)	
COVID-19 disease severity ‡				0.188			< 0.001
Asymptomatic	7 (1.6)	6 (1.5)	1 (2.6)		1 (0.3)	6 (5.3)	
Mild	205 (47.7)	189 (48.2)	16 (42.1)		111 (35.1)	94 (82.5)	
Moderate	216 (50.2)	196 (50.0)	20 (52.6)		202 (63.9)	14 (12.3)	
Severe	2 (0.5)	1 (0.3)	1 (2.6)		2 (0.6)	0 (0.0)	
Critical	0 (0.0)	0 (0.0)	0 (0.0)		0 (0.0)	0 (0.0)	
Vaccination status				< 0.001			< 0.001
3 dose of IV	157 (36.5)	157 (40.1)	0 (0.0)		155 (49.1)	2 (1.8)	
2 dose of IV	178 (41.4)	178 (45.4)	0 (0.0)		84 (26.6)	94 (82.5)	
1 doses of IV	6 (1.4)	6 (1.5)	0 (0.0)		2 (0.6)	4 (3.5)	
2 dose of AVV	34 (7.9)	34 (8.7)	0 (0.0)		34 (10.8)	0 (0.0)	
1 dose of AVV	15 (3.5)	15 (3.8)	0 (0.0)		15 (4.7)	0 (0.0)	
3 doses of PSV	2 (0.5)	2 (0.5)	0 (0.0)		2 (0.6)	0 (0.0)	
Unvaccinated	38 (8.8)	(0.0)	38 (100.0)		24 (7.6)	14 (12.3)	
Admission of ICU				< 0.001			0.025
Yes	17 (4.0)	8 (2.0)	9 (23.7)		17 (5.4)	0 (0)	
No	413 (96.0)	384 (98.0)	29 (76.3)		299 (94.6)	114 (100.0)	
Re-positive during convalescence phase §				0.001			0.019
Yes	85 (19.8)	70 (17.9)	15 (39.5)		71 (22.5)	14 (12.3)	
No	345 (80.2)	322 (82.1)	23 (60.5)		245 (77.5)	100 (87.7)	
Comorbidities				0.001			< 0.001
None	258 (60.0)	239 (61.0)	19 (50.0)		148 (46.8)	110 (96.5)	
1	67 (15.6)	65 (16.6)	2 (5.3)		65 (20.6)	2 (1.7)	
2	40 (9.3)	37 (9.4)	3 (7.9)		39 (12.3)	1 (0.9)	
≥3	65 (15.1)	51 (13.0)	14 (36.8)		64 (20.3)	1 (0.9)	

* Continuous variables were shown in median (interquartile ranges), and the Mann-Whitney U test was used to analyze the differences between the two groups. Categorical variables were summarized as counts (percentages) and analyzed by Pearson's χ^2 test. Percentages may not total 100 because of rounding.

† Child was defined as younger than 18 years old.

‡ COVID-19 disease severity was defined according to WHO living guidance for clinical management of COVID-19.

§ PCR re-positive during convalescence phase was defined as PCR Ct value less than 40 after two independent PCR-negative results with an interval of more than 24 hours.

Table S4. Correlation between inactivated vaccine doses and COVID-19 disease severity and progression *

	All				Child †			
	3 Doses (n = 157)	2 Doses (n = 178)	1 Dose (n = 6)	Unvaccinated (n = 38)	3 Doses (n = 2)	2 Doses (n = 94)	1 Dose (n = 4)	Unvaccinated (n = 14)
COVID-19 disease severity ‡								
Asymptomatic	1 (0.6)	5 (2.8)	0 (0.0)	1 (2.6)	0 (0.0)	5 (5.4)	0 (0.0)	1 (7.1)
Mild	66 (42.0)	104 (58.4)	4 (66.7)	16 (42.1)	2 (100.0)	76 (80.9)	3 (75.0)	13 (92.9)
Moderate	90 (57.3)	68 (38.2)	2 (33.3)	20 (52.6)	0 (0.0)	13 (13.8)	1 (25.0)	0 (0.0)
Severe	0 (0.0)	1 (0.6)	0 (0.0)	1 (2.6)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)
Critical	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)
Admission of ICU								
Yes	1 (0.6)	4 (2.2)	1 (16.7)	9 (23.7)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)
No	156 (99.4)	174 (97.8)	5 (83.3)	29 (76.3)	2 (100.0)	94 (100.0)	4 (100.0)	14 (100.0)
PCR re-positive during convalescence phase §								
Yes	30 (19.1)	25 (14.0)	3 (50.0)	15 (39.5)	0 (0.0)	9 (9.6)	2 (50.0)	3 (21.4)
No	127 (80.9)	153 (86.0)	3 (50.0)	23 (60.5)	2 (100.0)	85 (90.4)	2 (50.0)	11 (78.6)
Days of hospital stay and rehabilitation								
	27 (25-30)	28 (26-30)	31 (30-31)	29 (26-33)	28 (26-30)	28 (25-30)	31 (31-31)	28 (26-29)

* Continuous variables were shown in median (interquartile ranges), and categorical variables were summarized as counts (percentages). Percentages may not total 100 because of rounding.

† Child was defined as younger than 18 years old.

‡ COVID-19 disease severity was defined according to WHO living guidance for clinical management of COVID-19.

§ PCR re-positive during convalescence phase was defined as PCR Ct value less than 40 after two independent PCR-negative results with an interval of more than 24 hours.

Table S5. Laboratory Findings of Omicron infected patients*

	3 Doses (n = 157)	2 Doses (n = 178)	1 Dose (n = 6)	Unvaccinated (n = 38)	P-value †	
White-cell count — per mm³	6320 (5490-7230)	6310 (5300-7510)	7995 (7720-10680)	7420 (5130-8680)	0.016	3 V.S. 1: 0.004 2 V.S. 1: 0.009
Lymphocyte count — per mm³	2080 (1650-2430)	2350 (1910-2900)	2930 (2640-3340)	2600 (1530-4090)	0.001	3 V.S.2: < 0.001 3 V.S.1: 0.031
Neutrophil count — per mm³	3590 (2900-4230)	3260 (2440-4050)	4390 (3070-5720)	3590 (2510-4690)	0.045	3 V.S.2: 0.011
Monocyte count — per mm³	410 (350-500)	410 (340-520)	530 (390-590)	490 (420-580)	0.021	3 V.S. Un: 0.002 2 V.S. Un: 0.013
Platelet count — per mm³	262,000 (227,000-307,000)	288,000 (238,000-326,000)	316,000 (306,000-329,000)	290,000 (237,000-386,000)	0.008	3 V.S.2: 0.010 3 V.S.1: 0.011
SII	372 (239-642)	492 (314-578)	390 (248-550)	468 (330-625)	0.032	3 V.S.2: 0.003
NLR	1.72 (1.41-2.11)	1.40 (1.00-1.96)	1.40 (1.03-1.72)	1.52 (0.93-2.36)	<0.001	3 V.S.2: < 0.001
PLR	131 (104-162)	114 (96-141)	102 (97-118)	107 (84-159)	0.031	3 V.S.2: 0.009
MLR	0.20 (0.17-0.24)	0.17 (0.14-0.22)	0.17 (0.12-0.23)	0.21 (0.14-0.30)	0.003	3 V.S.2: < 0.001
C-reactive protein ≥10 mg/liter					<0.001	3 V.S. Un: < 0.001 2 V.S. Un: 0.007
Yes	2 (1.3)	6 (3.4)	1 (16.7)	6 (16.2)		
No	155 (98.7)	171 (96.6)	5 (83.3)	31 (83.8)		
Interleukin 6 ≥1.50 pg/milliliter					0.309	
Yes	45 (28.7)	54 (30.5)	1 (16.7)	16 (43.2)		
No	112 (71.3)	123 (69.5)	5 (69.5)	21 (56.8)		
CD4⁺ T cell count — per microliter	0.71(0.44-1.08)	0.75(0.47-1.15)	0.78(0.61-1.01)	0.65(0.39-1.16)	0.841	
CD8⁺ T cell count — per microliter	0.77(0.48-1.30)	0.80(0.51-1.45)	0.48(0.34-3.32)	0.97(0.48-2.30)	0.508	
CD4⁺/CD8⁺	1.68(1.27-2.14)	1.47(1.17-1.78)	1.67(1.47-1.99)	1.56(1.35-2.29)	0.034	3 V.S.2: 0.034
Treg cell count — per microliter	3.41(2.65-4.32)	3.54(2.67-4.60)	4.10(2.79-6.98)	2.69(1.93-3.36)	0.116	
Activated Treg cell count — per microliter	1.22(0.76-1.67)	0.96(0.59-1.35)	1.82(0.75-2.71)	0.74(0.39-1.03)	0.003	3 V.S. 2: 0.044 3 V.S. Un: 0.035
Th1 cell count — per microliter	25.59(21.32-31.84)	22.17(17.96-27.86)	23.72(16.80-30.07)	28.23(21.05-33.22)	0.001	3 V.S. 2: 0.001
Th2 cell count — per microliter	49.23(39.66-56.55)	60.58(49.10-67.72)	50.01(20.44-68.58)	45.81(38.87-60.84)	<0.001	3 V.S. 2: < 0.001
Th1/Th2	0.55(0.38-0.79)	0.37(0.28-0.55)	0.59(0.24-1.60)	0.62(0.36-0.76)	<0.001	3 V.S. 2: < 0.001

* Data regarding white-cell, lymphocyte, neutrophil, monocyte, platelet count, SII, NLR, PLR, MLR, and the measurement of C-reactive protein and Interleukin 6 were missing for 2 patients (0.5%). Data were missing for the proportion of T cell subsets in 106 patients (28.0%). SII was defined as follows: SII = P x N/L, where P, N and L are the preoperative peripheral blood platelet, neutrophil and lymphocyte counts per liter, respectively. NLR was defined as the absolute neutrophil count divided by the absolute lymphocyte count. PLR was defined as the platelet count divided by the number of lymphocytes. MLR was defined as the ratio of absolute monocyte count to absolute

lymphocyte count. Continuous variables were shown in median (interquartile ranges). Statistical significance was determined by Kruskal-Wallis H test between groups and Mann-Whitney U test between the two groups. Categorical variables were summarized as counts (percentages) and analyzed by Pearson's χ^2 test. Percentages may not total 100 because of rounding.

† Only statistically significant results are shown here.

Table S6. Correlation between hepatic and renal function and vaccination*

	3 Doses (n = 157)	2 Doses (n = 178)	1 Dose (n = 6)	Unvaccinated (n = 38)	P-value
Glutamic pyruvic transaminase — IU/liter	32.29 (20.32-67.33)	18.91 (12.53-40.79)	15.92 (12.06-28.17)	16.94 (13.85-29.29)	< 0.001
Glutamic oxalacetic transaminase — IU/liter	29.00 (23.14-37.88)	26.24 (22.96-34.05)	28.76 (27.16-31.73)	31.72 (25.79-39.11)	0.072
Total bilirubin — µmol/liter	12.33 (9.78-13.81)	11.68 (9.61-14.00)	-	10.62 (8.47-13.61)	0.322
Gamma-glutamyl transpeptidase — IU/liter	25.16 (17.33-54.26)	25.80 (15.14-51.63)	-	18.01 (14.47-38.44)	0.499
Alkaline phosphatase — IU/liter	68.12 (56.43-81.84)	74.89 (61.82-94.30)	-	79.67 (54.00-124.76)	0.133
Albumin — g/liter	42.73 (40.91-45.46)	41.43 (38.62-44.72)	-	36.12 (31.97-43.31)	0.008
Creatinine — µmol/liter	61.33 (50.86-71.81)	46.66 (38.31-58.80)	38.11 (37.61-59.12)	43.52 (32.30-58.82)	< 0.001
Urea nitrogen — mmol/liter	3.97 (3.35-4.69)	3.94 (3.10-4.68)	5.19 (4.06-5.65)	4.53 (3.48-5.87)	0.011
UREA/CREA †	0.07 (0.06-0.08)	0.08 (0.06-0.10)	0.12 (0.10-0.13)	0.11 (0.07-0.14)	< 0.001

* Data were missing for the measurement of glutamic pyruvic transaminase, glutamic oxalacetic transaminase, creatinine, urea nitrogen and UREA/CREA in 2 patients (0.5%), for total bilirubin, glutamyl transpeptidase, alkaline phosphatase and albumin in 322 patients (74.9%). Continuous variables were shown in median (interquartile ranges), and the Kruskal-Wallis H test was used to analyze the differences between groups.

† UREA/CREA was defined as the ratio of plasma urea nitrogen to plasma creatinine.

Table S7. Multivariate analysis of risk factors for COVID-19 disease severity*

	All		Child	
	OR (95% CI)	P-value	OR (95% CI)	P-value
Age	1.063 (1.046-1.080)	< 0.001	0.908 (0.755-1.093)	0.310
Gender				
Female	0.897 (0.566-1.422)	0.643	0.491 (0.175-1.374)	0.175
Male	Reference		Reference	
Vaccination status				
3 doses of IV	0.540 (0.220-1.326)	0.179	4.039 (0.052-314.191)	0.530
2 doses of IV	0.668 (0.278-1.600)	0.365	5.942 (0.799-44.124)	0.082
1 dose of IV	0.566 (0.061-5.217)	0.615	13.531 (0.829-220.964)	0.067
Unvaccinated	Reference		Reference	
Comorbidities				
With	1.014 (0.576-1.784)	0.962	3.216 (0.319-33.348)	0.319
Without	Reference		Reference	

* Statistical significance was determined by ordered multi-class logistic regression.

Table S8. Multivariate analysis of risk factors for ICU admission*

	All		Adult	
	OR (95% CI)	P-value	OR (95% CI)	P-value
Age	1.061 (1.012-1.112)	0.014	1.059 (1.009-1.112)	0.021
Gender				
Female	-		0.963 (0.223-4.155)	0.960
Male	Reference		Reference	
Vaccination status				
3 doses of IV	0.023 (0.002-0.224)	0.001	0.023 (0.002-0.222)	0.001
2 doses of IV	0.167 (0.038-0.739)	0.018	0.165 (0.037-0.732)	0.018
1 dose of IV	0.996 (0.045-21.970)	0.998	0.962 (0.038-24.604)	0.981
Unvaccinated	Reference		Reference	
Comorbidities				
With	30101052.05 (0.000-)	0.994	46178823.18 (0.000-)	0.996
Without	Reference		Reference	

* Statistical significance was determined by ordered multi-class logistic regression.

Table S9. Multivariate analysis of risk factors for PCR re-positive during convalescence phase*

	All		Adult	
	OR (95% CI)	P-value	OR (95% CI)	P-value
Age	1.019 (1.007-1.032)	0.002	1.023 (1.002-1.046)	0.035
Gender				
Female	-		1.457 (0.777-2.731)	0.241
Male	Reference		Reference	
Vaccination status				
3 doses of IV	0.317 (0.144-0.700)	0.004	0.301 (0.117-0.771)	0.012
2 doses of IV	0.297 (0.132-0.665)	0.003	0.279 (0.102-0.761)	0.013
1 dose of IV	1.985 (0.327-12.056)	0.456	1.201 (0.063-23.051)	0.903
Unvaccinated	Reference		Reference	
Comorbidities				
With	-		0.656 (0.329-1.308)	0.231
Without	Reference		Reference	

* Statistical significance was determined by ordered multi-class logistic regression.

Table S10. Multivariate analysis of risk factors for hospital stay and rehabilitation duration*

	All		Adult	
	OR (95% CI)	P-value	OR (95% CI)	P-value
Age	-		1.002 (0.985-1.020)	0.818
Gender				
Female	-		1.070 (0.642-1.780)	0.796
Male	Reference		Reference	
Vaccination status				
3 doses of IV	0.461 (0.225-0.946)	0.035	0.242 (0.092-0.635)	0.004
2 doses of IV	0.676 (0.334-1.367)	0.276	0.405 (0.149-1.097)	0.075
1 dose of IV	4.048 (0.431-38.029)	0.221	0.426 (0.023-8.032)	0.569
Unvaccinated	Reference		Reference	
Comorbidities				
With	-		0.992 (0.570-1.726)	0.992
Without	Reference		Reference	

* Statistical significance was determined by ordered multi-class logistic regression.

Table S11. COVID-19 vaccination status of plasma samples prior to infection *

	3 doses of IV (<i>n</i> = 80)	2 doses of IV (<i>n</i> = 39)	2 doses of AVV (<i>n</i> = 16)
CoronaVac	42 (52.5)	23 (59.0)	-
BBIBP-CorV	38 (47.5)	16 (41.0)	-
Ad5-nCoV	-	-	16 (100.0)

* Categorical variables were summarized as counts (percentages). IV, inactivated vaccine; AVV, adenovirus-vectored vaccine.

Table S12. Correlation between COVID-19 disease severity and vaccination-infection interval *

	All kinds of vaccine			Inactivated vaccine		
	All (n = 392)	Adult (n = 292)	Child (n = 100)	All (n = 341)	Adult (n = 241)	Child (n = 100)
Severity						
Asymptomatic	44.5 (43.3-46.5)	38	45 (44-47)	44.5 (43.3-46.5)	38	45 (44-47)
Mild	55 (45-136)	75.5 (43.5-182.5)	49 (45-64)	54 (45-136)	75 (45-182.5)	49 (45-64)
Moderate	78.5 (44-194.5)	81 (44.5-197)	47 (44-52)	75 (43-191)	80 (42-195)	47 (44-52)
Severe	172	172	-	172	172	-
Critical	-	-	-	-	-	-
P-value	0.020	0.504	0.186	0.053	0.597	0.186

* COVID-19 disease severity was defined according to WHO living guidance for clinical management of COVID-19. Continuous variables were shown in median (interquartile ranges). Statistical significance was determined by Kruskal-Wallis H test between groups.