## RESEARCH ARTICLE

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# **Comparison of antibody persistency through one year between one-dose and two-dose regimens of Ad5-nCoV vaccine for COVID-19**

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#### **ABSTRACT**

This post-hoc analysis compared the receptor-binding domain (RBD)-specific and pseudovirus neutralizing antibodies against the wild-type SARS-CoV-2 strain elicited by one or two doses (56-d interval) of Ad5 nCoV vaccine regimen (NCT04341389 and NCT04566770). Both trials had low-dose and high-dose groups. Propensity score matching was used to adjust the baseline between one- and two-dose regimens. To predict the decrease in antibody titers 1 y after vaccination, half-lives of RBD-binding antibodies and pseudovirus neutralizing antibodies were computed. We obtained 34 and 29 pairs of participants in the low- and high-dose groups based on the propensity score matching. The two-dose regimen of Ad5 nCoV increased the peaking level of neutralizing antibodies compared to the one-dose regimen at day 28, but the responses of the neutralizing antibodies were not consistent with those of the RBD antibodies. Half-lives of the RBD-binding antibodies in the two-dose Ad5-nCoV regimen (202–209 days) were longer than those in the one-dose regimen (136–137 d); half-lives of the pseudovirus neutralizing antibody in the one-dose Ad5-nCoV regimen (177 d) were longer than those in the two-dose regimen (116–131 d). The predicted positive rates of RBD-binding antibodies in the one-dose regimen (34.1%–38.3%) would be lower than those in the two-dose Ad5-nCoV regimen (67.0%–84.0%), while the positive rates of pseudovirus neutralizing antibodies in the one-dose regimen (65.4%–66.7%) would be higher than those in the two-dose regimen (48.3%–58.0%). The two-dose Ad5-nCoV regimen with a 56-d interval had no effect on the persistence of neutralizing antibodies but slowed decay trend of RBD-binding antibodies.

# **Introduction**

<span id="page-0-6"></span><span id="page-0-5"></span><span id="page-0-4"></span>COVID-19 vaccines are the most effective weapon against the SARS-CoV-2 associated infections, severe diseases, and deaths.[1](#page-6-0)[–3](#page-6-1) Although many COVID-19 vaccines induce a quick protective response within a short time, antibody responses and vaccine effectiveness have been observed to decrease over time.<sup>4,5</sup> BNT162b2 and mRNA-1273 elicited excellent shortterm neutralizing antibody responses and protective effectiveness[.6](#page-6-4)[,7](#page-6-5) But these mRNA vaccines induced high serum neutralizing antibodies that waned by 3–6 months, with a half-life of 60 d.<sup>[8](#page-6-6)[,9](#page-6-7)</sup> Conversely, Ad26.COV2.S generated lower initial neutralizing antibody titers than BNT162b2 and mRNA-1273, $^{10}$ but these neutralizing antibody responses and clinical effective-ness were fairly durable for at least 8 months.<sup>[11](#page-6-9),12</sup> Different COVID-19 vaccines could have different profiles for antibody persistency, which needs to be explored.

<span id="page-0-10"></span><span id="page-0-9"></span><span id="page-0-8"></span><span id="page-0-7"></span>Ad5-nCoV, an adenoviral vector vaccine based on a replication-defective human adenovirus type 5 encoding the full spike of SARS-CoV-2, was developed by CanSino in China.<sup>13</sup> In the phase 2 trial of one-dose Ad5-nCoV, in the  $1 \times 10^{11}$  and  $5 \times 10^{10}$  viral particles dose groups, significant neutralizing antibody responses to pseudovirus SARS-CoV-2 were produced by both doses of vaccine, with GMTs of 61.4 (95% CI 53.0, 71.0) and 55.3 (95% CI 45.3, 67.5).<sup>14</sup> Besides, in the two-dose Ad5-nCoV regimen study, in participants aged 18 to 55, the second vaccination of Ad5-nCoV elicited higher pseudovirus neutralizing antibodies (76.8; 95% CI 52.4, 112.7) at day 28 than the first vaccination (45.1; 95% CI 24.8, 81.8).<sup>15</sup> In an efficacy trial of onedose Ad5-nCoV, protection against symptomatic, PCRconfirmed, COVID-19 infection at 28 d or more postvaccination was reported to be 57.5% (95% CI 39.7%, 70.0%).<sup>[16](#page-6-14)</sup> On May 19, 2022, the World Health Organization (WHO) issued an emergency use listing (EUL) for Ad5-nCoV by intramuscular injection, $17$  which was approved by more than 10 countries later.<sup>18</sup> However, previous studies about Ad5-nCoV presented above only reported antibody responses at day 28 after one-dose or two-dose vaccination; the antibody persistency induced by one or two doses of Ad5-nCoV is still unknown.<sup>14[,15](#page-6-13)</sup>

<span id="page-0-12"></span><span id="page-0-11"></span>In this study, the antibody persistency in one- and two-dose regimens of Ad5-nCoV were reported and whether two-dose Ad5-nCoV vaccines with the 56-d interval could yield benefits

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#### **ARTICLE HISTORY**

Received 31 January 2023 Revised 1 June 2023 Accepted 13 June 2023

#### **KEYWORDS**

Ad5-nCoV; persistence of immunity; RBD-binding antibody; pseudovirus neutralizing antibody; power-law model; exponential decay model

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<span id="page-0-1"></span>Supplemental data for this article can be accessed on the publisher's website at [https://doi.org/10.1080/21645515.2023.2230760.](https://doi.org/10.1080/21645515.2023.2230760)

of a longer duration of antibody responses than the one-dose regimen were explored.

## **Materials and methods**

#### *Study design, participants and procedures*

These two randomized, open-label, placebo-controlled trials evaluating the safety and immunogenicity of the one-dose and twodose regimens of Ad5-nCoV vaccine were conducted from April 2020 to December 2020 and from September 2020 to July 2021, respectively, in China.<sup>14,[15](#page-6-13)</sup> Two trial protocols and informed consents were approved by the institutional review board of the Jiangsu Provincial Center of Disease Control and Prevention. Written informed consent was obtained from all participants before screening. Detailed study designs have been reported previously. In brief, 253, 129 and 126 healthy adults aged 18 y and older were recruited to receive a single injection of the Ad5-nCoV vaccine of high-dose  $(1 \times 10^{11}$  viral particles per dose), low-dose  $(5 \times 10^{10}$  viral particles per dose), or placebo (only the vaccination excipients without viral particles) in the one-dose regimen trial (ClinicalTrials.gov number: NCT04341389, Supplementary Figure S1a). Blood samples were taken from participants at day 0 immediately before vaccination, at day 28, and month 6 post vaccination. While in the two-dose regimen trial (ClinicalTrials.gov number: NCT04566770, Supplementary Figure S1b), 100, 120, and 60 participants aged 18 y and older were randomly assigned to receive high-dose Ad5-nCoV ( $1 \times 10^{11}$ ) viral particles per dose), or low-dose Ad5-nCoV ( $5 \times 10^{10}$  viral particles per dose), or placebo, with a 56-d interval. For the second injection, the same dosage as for the first injection was given to each group. Blood samples were taken from participants at day 0 immediately before the first dose, at day 28 and month 6 after the second dose. We used participants in the vaccine groups in these two trials aged 18 y and above for the analysis (as described in [Figure 1](#page-1-0)).

<span id="page-1-1"></span>The detailed methods of assays have been reported previously.<sup>19</sup> The receptor-binding domain (RBD)-specific antibodies against the wild-type SARS-CoV-2 strain were measured <span id="page-1-3"></span><span id="page-1-2"></span>by ELISA kits (Beijing Wantai BioPharm, Beijing, China) and pseudovirus neutralization antibodies against the wild-type SARS-CoV-2 strain were measured by neutralizing antibodies responding to pseudovirus (a vesicular stomatitis virus pseudovirus system expressing the spike glycoprotein).<sup>20</sup> Neutralizing antibodies against the vaccine vector Ad5 were measured by serum neutralizing assay.<sup>21</sup> The detection limits for the RBDspecific antibody test, neutralizing antibody test to pseudovirus, and neutralizing antibody test to vaccine vector Ad5 were 1:40, 1:10, and 1:12, respectively. Undetectable antibody titers were assigned values of one-half the limits of detection.

## *Outcomes*

We compared the vaccine elicited antibody titers (RBD-binding antibodies and pseudovirus neutralizing antibodies) in participants receiving one-dose or two-dose regimens of Ad5-nCoV at day 28 and month 6 after the vaccination. The half-life of antibodies were also assessed in one- and two-dose Ad5-nCoV regimens to predict positive rates (proportion of participants whose measured value was higher than the detection limits) and GMTs of antibodies 1 y later.

#### *Statistical analyses and models*

Multiple linear regression analysis was performed to evaluate the multivariate associations between independent variables in baseline (age, sex, BMI, and levels of preexisting adenovirus type-5 neutralizing antibody) and dependent variables (RBD-binding antibodies and pseudovirus neutralizing antibodies). An analysis of covariance was used to evaluate the consistency of the measurement of the RBD-binding or pseudovirus neutralizing antibody responses in two different laboratories (assays in the onedose regimen trial were conducted by National Institutes for Food and Drug Control and assays in the two-dose regimen trial were conducted by Vazyme Biotech Co., Ltd). Propensity score matching would be applied in these two Ad5-nCoV regimens in order to visualize the differences in

<span id="page-1-0"></span>

**Figure 1.** Trial profile. vp=viral particles. \*2 did not receive the second vaccination at day 56.

antibody titers when baseline characteristics varied between the one- and two-dose regimens and had an impact on antibody titers.

Levels of RBD-binding and pseudovirus neutralizing antibodies were presented as GMTs with 95% CIs, and fold decreases were calculated. Log-transformed antibody titers were tested for normality and homogeneity of variance using a Shapiro-Wilk and Levene's test, respectively. Comparisons were done between two groups by using Student's t-test for data with a normal distribution and homogeneous variance; otherwise, data were analyzed using the nonparametric Wilcoxon rank-sum test. Categorical data were analyzed using the Chi-square test or Fisher's exact test. Except for the part about decay models, subsequent analyses were based on the participants after propensity score matching. Statistical analysis other than model fitting was performed using SPSS (Version 26).

In order to estimate the antibody half-lives to predict positive rates and GMTs (positive participants) 1 y later, decay models were fitted based on data collected on day 28 and month 6 after the last Ad5-nCoV vaccination. The use of decay models was supported by prior results from the persistence of mRNA-1273 vaccination for the COVID-19 study, which demonstrated that the power-law model and exponential decay models, respectively, are the best matches for binding and neutralizing antibodies.<sup>[9](#page-6-7)</sup> Statistical analyses were done using the lme4 package in R 4.0.4 to compute models and 95% confidence intervals of decay rates.

#### *Exponential decay model*

The exponential decay model was assumed a steady decay rate over time. Exponential model was given by the following:<sup>[9](#page-6-7),[22](#page-6-20)</sup>

$$
Log_{10}(Titer) = \alpha + \beta \times (study\ day + c) \tag{1.1}
$$

<span id="page-2-0"></span>The half-life of exponential model was calculated by following  $formula: <sup>9,22</sup>$ 

$$
t_{\frac{1}{2}} = \frac{\log_{10}(0.5)}{\hat{\beta}} \tag{1.2}
$$

### *Power law model*

<span id="page-2-1"></span>The power law model was assumed decay rates decreased over time. Power law model in one-dose Ad5-nCoV trial was given by the following:<sup>[9](#page-6-7),23</sup>

$$
Log_{10}(Titer) = \alpha + \beta \times (log_{10}(study \, day) + c) \qquad (2.1.1)
$$

The half-life of power law model in one-dose trial was calculated as:

$$
Log_{10}\left(t_{\frac{1}{2}}\right) = log_{10}(study\ day) + \frac{log_{10}(0.5)}{\hat{\beta}} \tag{2.1.2}
$$

Study day was offset by 56 d to account for the two-dose regimen and the power law model in the two-dose Ad5-nCoV trial was given by the following:<sup>9</sup>

$$
Log_{10}(Titer) = \alpha + \beta \times (log_{10}(study \, day - 56) + c) \quad (2.2.1)
$$

And the half-life of power law model in two-dose trial was calculated following the formula below:

$$
Log_{10}\left(t_{\frac{1}{2}}\right) = log_{10}(study\ day - 56) + \frac{log_{10}(0.5)}{\hat{\beta}} \tag{2.2.2}
$$

*α* and *β* were the intercept and the decay rate, representing fixed effect. *c* was an arbitrary small constant.

 $\hat{\beta}$  was the point estimate of decay rate.  $t_{1/2}$  was the day when titers decrease to half compared with the starting value. The 95% confidence intervals of half-life were computed by 1000 bootstrap simulations.

## **Results**

#### *Population characteristics*

This post-hoc analysis for the durability of the vaccine-elicited humoral immune response was limited to participants aged 18 and older receiving the one-dose or two-dose regimen of Ad5 nCoV. As described in the Supplementary Tables S1 and S2, multiple linear regression revealed that antibody titers were influenced by age and preexisting Ad5 neutralizing antibodies. And there were marked differences in age distribution between one-dose and two-dose regimens: in the low-dose groups, 116 (89.9%) participants aged 18–55 were in the one-dose regimen, while 20 (16.9%) were in the two-dose regimen; in the highdose groups, 223 (88.1%) participants aged 18–55 were in the one-dose regimen, while nobody aged 18–55 y was in the twodose regimen. Furthermore, regardless of the vaccine dosage, the antibody titers were comparable between laboratories from the two regimens when age and preexisting Ad5 neutralizing antibody titers were set as covariables using analysis of covariance ( $p = .303$  and .471 for the low-dose groups, and  $p = .103$ and .319 for the high-dose groups).

We obtained 34 and 29 pairs of participants in the low- and high-dose groups by propensity score matching in order to visualize the difference in antibody titers between participants in the one- and two-dose Ad5-nCoV regimens since age and preexisting adenovirus type-5 neutralizing antibody titers were different between these two regimens (Supplementary Table S2). The demographic characteristics of the matched participants were similar between the trials. The mean ages of the participants in the low-dose groups were 51.3 (SD 8.2) and 51.4 (SD 8.0) y in one- and two-dose regimens, respectively; in the high-dose groups, they were 60.9 and 60.5 y (As described in [Table 1](#page-3-0)). Both the RBD-binding and pseudovirus neutralizing antibody titers at day 28 following the first immunization were comparable between matching groups with the same vaccine dosages from the two trials (*p* = .464 and .320 for the low-dose groups, and  $p = .507$  and .652 for the high-dose groups).

# *Antibody responses through 6 months after the last vaccination*

At day 28 after the last vaccination, no difference in RBDbinding antibodies was found between one-dose and two-dose Ad5-nCoV regimens both for the low- or high-dose group. At month 6, the two-dose regimen of Ad5-nCoV (low-dose group: 285.3; 95% CI 179.1, 454.3; high-dose group: 178.0; 95% CI 126.7, 250.1) showed significantly higher RBDbinding antibodies in terms of GMTs than did the one-dose

<span id="page-3-0"></span>



<span id="page-3-2"></span>Data are mean (SD), number of participants (%) and p value. SD = standard deviation. BMI = body-mass index. N = the number of participants after propensity score matching.

<span id="page-3-1"></span>**Table 2.** RBD-binding and pseudovirus neutralizing antibody titers of participants after propensity score matching.

		Low-dose group				High-dose group			
Time point	n	One-dose trial	Two-dose trial	$p$ value	n	One-dose trial	Two-dose trial	p value	
RBD-binding antibody									
Day 0	34	21.5 (18.6, 24.9)	20.7 (19.3, 22.3)	.654	29	21.4 (18.7, 24.5)	20.0 (20.0, 20.0)	.326	
28 days after the last dose	34	456.9 (285.7, 730.7)	604.7 (411.0, 890.0)	.352	29	491.3 (320.2, 753.8)	422.3 (289.4, 616.1)	.589	
6 months after the last dose	34	59.5 (41.4, 85.5)	285.3 (179.1, 454.3)	$-.001$	29	57.7 (39.4, 84.6)	178.0 (126.7, 250.1)	$-.001$	
Pseudovirus neutralizing antibody									
Day 0	34	6.0 (4.9, 7.3)	5.6 (5.0, 6.3)	.538	29	5.5 (4.9, 6.2)	5.4 (4.9, 6.0)	.750	
28 days after the last dose	34	55.2 (34.8, 87.4)	96.4 (70.3, 132.0)	.026	29	46.3 (28.6, 75.1)	104.9 (77.0, 143.0)	.005	
6 months after the last dose	34	25.1 (17.6, 35.9)	32.2 (20.7, 50.1)	.376	29	25.7 (17.1, 38.7)	41.6 (29.6, 58.4)	.069	

Antibody titers were compared at timepoints of before and after the last vaccination. Data are GMT (95% CI) and p value. n = the number of participants after propensity score matching. Bold indicates *p* < 0.05.

regimen of Ad5-nCoV (low-dose group: 59.5; 95% CI 41.4, 85.5; high-dose group: 57.7; 95% CI 39.4, 84.6, [Table 2](#page-3-1) and Supplementary Figure S2).

However, pseudovirus neutralizing antibodies postvaccination showed different trends from RBD-binding antibodies. At day 28 after the last Ad5-nCoV shot, the GMTs of pseudovirus neutralizing antibodies in the two-dose regimen (low-dose group: 96.4; 95% CI 70.3, 132.0; high-dose group: 104.9; 95% CI 77.0, 143.0) were higher than those in the onedose regimen (low-dose group: 55.2; 95% CI 34.8, 87.4; highdose group: 46.3; 95% CI 28.6, 75.1). At month 6 after the last vaccination, pseudovirus neutralizing antibodies were not significantly different between the one-dose and two-dose regimens of Ad5-nCoV [\(Table 2](#page-3-1) and Supplementary Figure S2).

The reduction in the one-dose regimen of RBD-binding antibody titers for low-dose groups was 7.7 folds (95% CI 5.8, 10.3) between peak responses at day 28 and month 6, while in the two-dose regimen there was a gradual decline of a 2.1-fold (95% CI 1.5, 3.0) decrease. In the high-dose group, similar decreases in RBD antibodies were observed as in the low-dose group: in the one- and two-dose regimens, RBDbinding antibodies decreased by 8.5 (95% CI 6.0, 12.0) and 2.4 folds (95% CI 1.8, 3.1), respectively. Both the one- and twodose Ad5-nCoV regimens demonstrated a gradual decline in pseudovirus neutralizing antibodies from day 28 to month 6 following the last vaccination. These declines were 2.2 (95% CI 1.5, 3.2) and 3.0 (95% CI 2.2, 4.0) folds in the low-dose groups in one- and two-dose Ad5-nCoV regimens and 1.8 (95% CI

1.2, 2.8) and 2.5 (95% CI 1.9, 3.4) folds in the high-dose groups, respectively.

### *Modeling the persistence of antibody responses*

In the low-dose groups, by using power-law models, the estimated half-life of RBD-binding antibodies for participants was 137 d (95% CI 129, 149) in one-dose Ad5-nCoV regimen and shorter than 202 d (95% CI 168, 250) in the two-dose Ad5 nCoV regimen. Likewise, the high-dose group showed similar decay trends as the low-dose group (one-dose regimen: 136; 95% CI 129, 145; two-dose regimen: 209; 95% CI 174, 256). While the estimated half-lives of pseudovirus neutralizing antibodies calculated by exponential decay models were 177 d (95% CI 143, 251) in the one-dose regimen, which was longer than 116 d (95% CI 104, 131) in the two-dose regimen (*p*  < .001); the high-dose group showed similar half-lives of pseudovirus neutralizing antibodies as the low-dose group (onedose regimen: 177 95% CI 143, 232; two-dose-regimen: 131; 95% CI 116, 151, as described in [Table 3\)](#page-4-0).

As shown in [Table 4](#page-4-1), 1 y following vaccination, the positive rates and GMTs for RBD-binding antibodies with low dosage in the one-dose and two-dose Ad5-nCoV regimens were estimated to be 34.1% (95% CI 32.5%, 39.7%) with a GMT of 98.9 (95% CI 77.6, 125.9), and 67.0% (95% CI 66.1%, 72.9%) with a GMT of 164.7 (95% CI 134.9, 204.2), respectively. In the high-dose groups, positive rates of RBDbinding antibodies would be 38.3% (95% CI 35.8%, 39.5%)

<span id="page-4-0"></span>**Table 3.** Estimates of antibody half-life.

Dose group	N	Regimen	Decay rate	Half-time				
Power-law model for RBD-binding antibody								
Low-dose group	126	one-dose	$-1.1594$	137				
			$(-1.2193, -1.0922)$	(129, 149)				
	118	two-dose	$-0.8782$	202				
			$(-0.9982, -0.7614)$	(168, 250)				
High-dose group	243	one-dose	$-1.1696$	136				
			$(-1.2181, -1.1167)$	(129, 145)				
	100	two-dose	$-0.8567$	209				
			$(-0.9766, -0.7479)$	(174, 256)				
Exponential decay model for pseudovirus neutralizing antibody								
Low-dose group	126	one-dose	$-0.0017$	177				
			$(-0.0021, -0.0012)$	(143, 251)				
	118	two-dose	$-0.0026$	116				
			$(-0.0029, -0.0023)$	(104, 131)				
High-dose group	243	one-dose	$-0.0017$	177				
			$(-0.0021, -0.0013)$	(143, 232)				
	100	two-dose	$-0.0023$	131				
			(-0.0026, -0.0020)	(116, 151)				

Half-life in power law models estimated at month 6 after the last vaccination in one- and two-dose Ad5-nCoV regimens. Decay rate in the table is the β in the power-law or exponential decay model. *N* = the number of participants aged 18 y above in one-dose and two-dose trials of Ad5-nCoV.

<span id="page-4-1"></span>



Positive rates in the power-law model were estimated based on titers at month 6 of the one- and two-dose regimen of Ad5-nCoV. *N* = the number of participants aged 18 y and older in the one-dose and two-dose Ad5-nCoV. Data are number of responders, positive rates (%) and GMT (95% CI) of positive participants. GMT = geometric mean titer. 95% CI = 95% confidence interval. Responders were participants whose measured value was higher than the detection limits.

and 84.0% (95% CI 82.0, 85.0) in the one-dose and two-dose Ad5-nCoV regimens. It suggested that positive rate and GMT for RBD-binding antibodies in the two-dose Ad5 nCoV regimen were higher than those in the one-dose regimen (low-dose group: 67.0% vs. 34.1%, *p* < .001; high-dose group: 84.0% vs. 38.3%,  $p < .001$ ). The positive rates and GMTs for pseudovirus neutralizing antibodies with low dosage in one- and two-dose regimens were 66.7% (95% CI 58.7%, 72.2%) with a GMT of 27.7 (95% CI 23.4, 33.1), and 48.3% (95% CI 44.9%, 52.5%) with a GMT of 24.2 (95% CI 20.4, 28.8), respectively. In the high-dose groups, the positive rate in the one-dose regimen would be 65.4% (95% CI 57.6%, 71.6%), with a GMT of 31.8 (95% CI 28.2, 36.3); in the twodose regimen, the positive rate would be 58.0% (95% CI 54.0%, 69.0%), with a GMT of 26.4 (95% CI 21.9, 32.4). And findings showed that the positive rate for pseudovirus neutralizing antibodies with a low dosage in the two-dose Ad5-nCoV regimen was lower than it was in the one-dose

regimen (low-dose group: 48.31% vs. 66.67%, *p* = .004; highdose group: 58.0% vs. 65.4%, *p* = .194).

## **Discussion**

In this study, we discovered that the two-dose Ad5-nCoV regimen produced higher peaked pseudovirus neutralizing antibodies than the one-dose regimen, but it had no effect on the peaking levels of RBD-binding antibody titers. Compared with one-dose Ad5-nCoV, although the decay trend of pseudoviruses was not slowed by the two-dose regimen, it did slow the decline of RBD-binding antibodies. We speculate that the phenomenon that there is no difference in pseudovirusneutralizing antibodies between the two-dose immunization group and the one-dose immunization group at 6 months after immunization may be due to the small sample size obtained after propensity score matching, resulting in "false negative" results.

The half-lives of RBD antibodies (approximately 7 months in two-dose Ad5-nCoV regimen and 5 months in one-dose regimen) and pseudovirus-neutralizing antibodies of Ad5 nCoV (approximately 4 months and 6 months in two-dose and one-dose Ad5-nCoV regimen) were longer than those of two-dose mRNA-1273, whose estimated half-lives for binding antibodies and pseudovirus neutralizing antibodies were 109 d (3.9 months) and 69 d (2.5 months), respectively. These findings were also supported by a comparison of the durability of the Ad26.COV2.S and mRNA COVID-19 vaccines, where Ad26.COV2.S exhibited a greater durability.<sup>[4](#page-6-2)</sup> It suggested that antibodies elicited by adenovirus-vectored COVID-19 vaccines may be more durable than mRNA COVID-19 vaccines. However, in our study, antibody titers 12 months after vaccination exhibited a greater reduction relative to peak antibody titers, and neutralizing antibody positivity was reduced by half by 1 y after vaccination. Among these positive participants, the antibody titers of some participants were only higher than the detection limit, and such critical antibody titers may not be enough to prevent breakthrough infection with COVID-19. Therefore, from the perspective of antibody titers, homologous immunization by two doses of Ad5-nCoV vaccine with a 56-d interval has limited benefit compared with one dose of Ad5-nCoV vaccine. In addition, the level of antibodies produced by high doses of this vaccine is similar to that of low doses, which means that if you consider increasing the dose of this vaccine to induce a stronger level of antibody immunity, it will also have little effect, and low doses of vaccine are sufficient to induce immune levels.

<span id="page-5-0"></span>The vaccine-elicited immune responses against SARS-CoV-2 may also be compromised by preexisting neutralizing antibodies against adenovirus type-5 vaccine vectors. In the study of twodose regimen of Ad5-nCoV, the first dose increased anti-Ad5 neutralizing antibodies by 2.4–7.5 times before the second vaccination.[15](#page-6-13) The high levels of anti-vectored antibodies induced by the first vaccination had an impact on the twodose regimen of adenovirus-vectored COVID-19 vaccines. However, ChAdOx1 nCoV-19 demonstrated that a primeboost interval of over 12 weeks provided higher protective efficacy than one of about 6 weeks. $24$  This confirmed that a longer interval between homologous vaccine doses can improve the negative effect of the response to the vectors. In order to minimize the negative effect of preexisting anti-Ad5 antibodies, a wider prime-boost interval for homologous or heterologous prime-boost regimens needs to be considered. The immune persistence and booster effects of Ad5-nCoV were studied (NCT04568811), and it was discovered that when boosted at 6 months, neutralizing antibody titers increased more than 7 times compared to the peak level following the first dose.<sup>25</sup> However, according to our findings, the peak neutralizing antibody titers only increased by 1.7–2.3 times after the booster immunizations with an interval of 56 d, which was much lower than the 7-fold peak neutralizing antibody titers increased by the booster immunizations with an interval of 6 months. Therefore, boosting homologous immunization with Ad5 nCoV every 6 months is recommended.

Our study had some limitations. Firstly, we just recorded antibody titers within 6 months after the last immunization, decay models need to be validated with real data of longer time <span id="page-5-1"></span>points to verify their accuracy. Secondly, we did not measure neutralizing antibodies to live virus because neutralization assays against live virus can only be done in specialist laboratories under category 3 biological safety conditions. Pseudovirus neutralizing antibodies, however, typically correlate with live virus neutralizing antibodies, so they can be used in place of those in the absence of the P3 laboratory.<sup>26–29</sup> Thirdly, because immunological surrogate endpoints of COVID-19 vaccines have not been established, the predicted antibody titers at future time points may not be directly related to the vaccine efficacy. Fourth, the Omicron strain, which is currently the most prevalent epidemic in the world, has a number of spike mutations, is highly transmissible, and is partially immune to partial neutralizing antibodies. Some investigations indicated that SARS-CoV-2 variants of concerns (VOCs), particularly Omicron, decreased the antibody responses generated by COVID-19 vaccinations.<sup>30-32</sup> However, responses to these VOCs were not assessed in our study.

## <span id="page-5-2"></span>**Conclusion**

The results of this study have extended our knowledge of the antibody persistency of the Ad5-vectored COVID-19 vaccines (Ad5-nCoV) in one-dose and two-dose regimens. Given the titers and model predictions, the two-dose Ad5-nCoV regimen did not bring a significant benefit over than the one-dose regimen; other strategies of boosting, such as extending the interval time between two Ad5-nCoV injections require further study.

#### **Acknowledgments**

We thank the China National Institutes for Food and Drug Control and the Vazyme Biotech Co., Ltd. in China, for their independent serologic assays.

#### **Author contributions**

Jia-Lu Feng and Wen-Juan Wang drafted the manuscript. Jing-Xin Li and Feng-Cai Zhu contributed to critical review and revising of the report. Jia-Lu Feng, Hui Zheng and Lai-Run Jin were responsible for statistical analysis. Wen-Juan Wang, Peng-Fei Jin, Xin Xia, Xiao-Yin Zhang and Zhuo-Pei Li participated in the site work, including the follow-up, and data collection. All authors contributed to drafting/critically revising the manuscript for intellectual content, provided final approval of the version to be published.

#### **Disclosure statement**

No potential conflict of interest was reported by the author(s).

## **Funding**

This work was supported by the National Science Foundation for Excellent Young Scholars of China (grant number 82222062 to Jing-xin Li); the General Program of the National Natural Science Foundation of China (grant number 82173584 to Jing-xin Li); Jiangsu Provincial Science Foundation for Distinguished Young Scholars (grant number BK20220064 to Jing-xin Li) and the Key Program of Jiangsu Provincial Science and Technology Plan (grant number BE2021738 to Feng-cai Zhu).

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#### **Data availability statement**

Data reported in this study are not publicly shared. The data that support the findings of this study are available from the authors under reasonable request and with permission of the Jiangsu Provincial Center for Disease Control and Prevention. Individuals wishing to access data should submit requests for access to the corresponding author (jszfc@vip.sina.com).

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