

THE LANCET

Supplementary appendix

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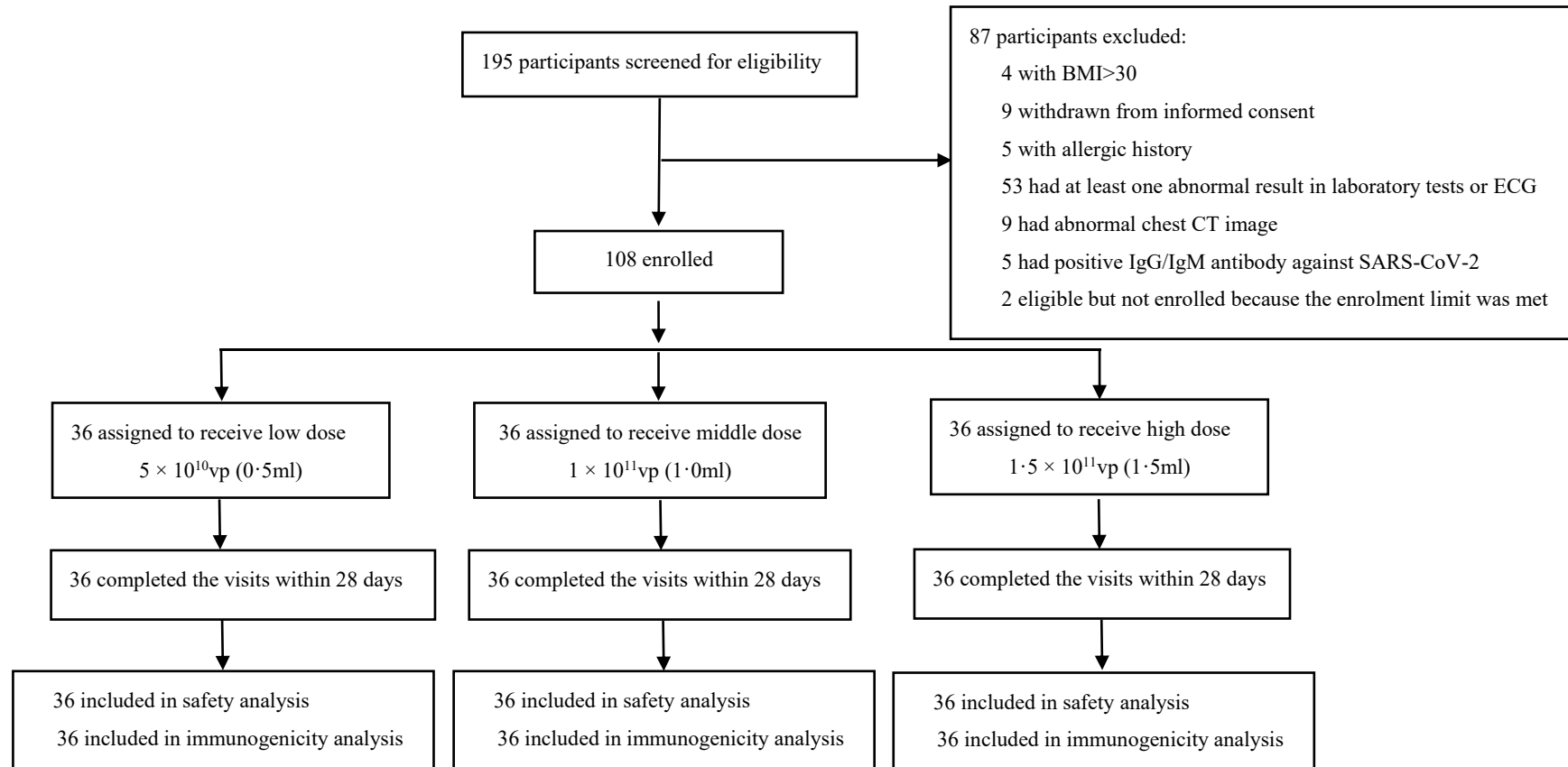
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Appendix 1. Trial profile.



BMI=body mass index. ECG=electrocardiograph. CT=computerized tomography. IgG=immunoglobulin G. IgM=immunoglobulin M. SARS-CoV-2= severe acute respiratory syndrome coronavirus 2.

Appendix 2. List of severe (grade 3) adverse reactions reported within 28 days of follow-up.

ID of participants	Dose group	Symptom	Grade	Start time	Resolve time	Treatment
027	Low dose	Fever	3	Day 0 after vaccination	48h	No treatment
029	Low dose	Fever	3	Day 1 after vaccination	24h	No treatment
060	Middle dose	Fever	3	Day 1 after vaccination	48h	No treatment
067	Middle dose	Fever	3	Day 1 after vaccination	48h	No treatment
075	High dose	Fever	3	Day 1 after vaccination	48h	Self-purchased medicine
		Dyspnea	3	Day 1 after vaccination	24h	No treatment
		Fatigue	3	Day 1 after vaccination	48h	No treatment
		Muscle pain	3	Day 1 after vaccination	48h	No treatment
077	High dose	Fever	3	Day 1 after vaccination	24h	No treatment
078	High dose	Fatigue	3	Day 0 after vaccination	24h	No treatment
		Joint pain	3	Day 0 after vaccination	24h	No treatment
080	High dose	Fever	3	Day 1 after vaccination	48h	No treatment
082	High dose	Fever	3	Day 1 after vaccination	48h	No treatment
087	High dose	Fever	3	Day 1 after vaccination	48h	Self-purchased medicine

Appendix 3. Logistic regression analysis of the incidence of fever with the baseline characters, vaccine dose and pre-existing Ad5 antibodies.

Model parameters	Estimates	P value	OR	
			Point estimates	95%CI by Wald
Univariate analysis*				
Middle dose	0·00	>0·99	1·00	0·39, 2·55
High dose	0·56	0·24	1·75	0·69, 4·45
Age	0·939	0·049	2·56	1·01, 6·51
Sex	-0·52	0·18	0·60	0·28, 1·28
Pre-existing Ad5 antibodies (>200)	-1·31	0·0013	0·27	0·12, 0·60
Multivariate analysis†				
Intercept	0·09	0·85		
Middle dose	-0·04	0·95	0·97	0·35, 2·66
High dose	0·39	0·45	1·48	0·54, 4·07
Age	0·53	0·042	2·87	1·04, 7·94
Sex	-0·32	0·14	0·53	0·23, 1·23
Pre-existing Ad5 antibodies (>200)	-1·26	0·0028	0·28	0·12, 0·65

* The dose group (the low-dose group as the reference), age (stratified into 18-44, 45-60; using 45-60 years as reference), sex (female as reference), and the level of Ad5 antibodies pre-vaccination (≤ 200 as reference) are analysed in univariate logistic regression, respectively, before involved in a multivariable analysis. † The occurrence of adverse reactions (fever) is the dependent variable. The independent variables are the dose group (the low-dose group was the reference), age (stratified into 18-44, 45-60; using 45-60 years as reference), sex (female as reference), and the level of Ad5 antibodies pre-vaccination (≤ 200 as reference).

Appendix 4. Abnormal changes of laboratory tests on day 7 after vaccination.

Laboratory measures		Low dose (n=36)	Middle dose (n=36)	High dose (n=36)	Total
Platelet decrease	Any	1 (3)	1 (3)	2 (6)	4 (4)
	Grade 1	1 (3)	0 (0)	1 (3)	2 (2)
	Grade 2	0 (0)	1 (3)	1 (3)	2 (2)
ALT increase	Any	2 (6)	5 (14)	3 (8)	10 (9)
	Grade 1	2 (6)	5 (14)	3 (8)	10 (9)
AST increase	Any	0 (0)	0 (0)	1 (3)	1 (1)
	Grade 1	0 (0)	0 (0)	1 (3)	1 (1)
Hyperglycemia (fasting)	Any	1 (3)	1 (3)	2 (6)	4 (4)
	Grade 1	0 (0)	1 (3)	0 (0)	1 (1)
	Grade 2	1 (3)	0 (0)	2 (6)	3 (3)
Total bilirubin increase	Any	4 (11)	1 (3)	4 (11)	9 (8)
	Grade 1	4 (11)	0 (0)	3 (8)	7 (6)
	Grade 2	0 (0)	1 (3)	1 (3)	2 (2)

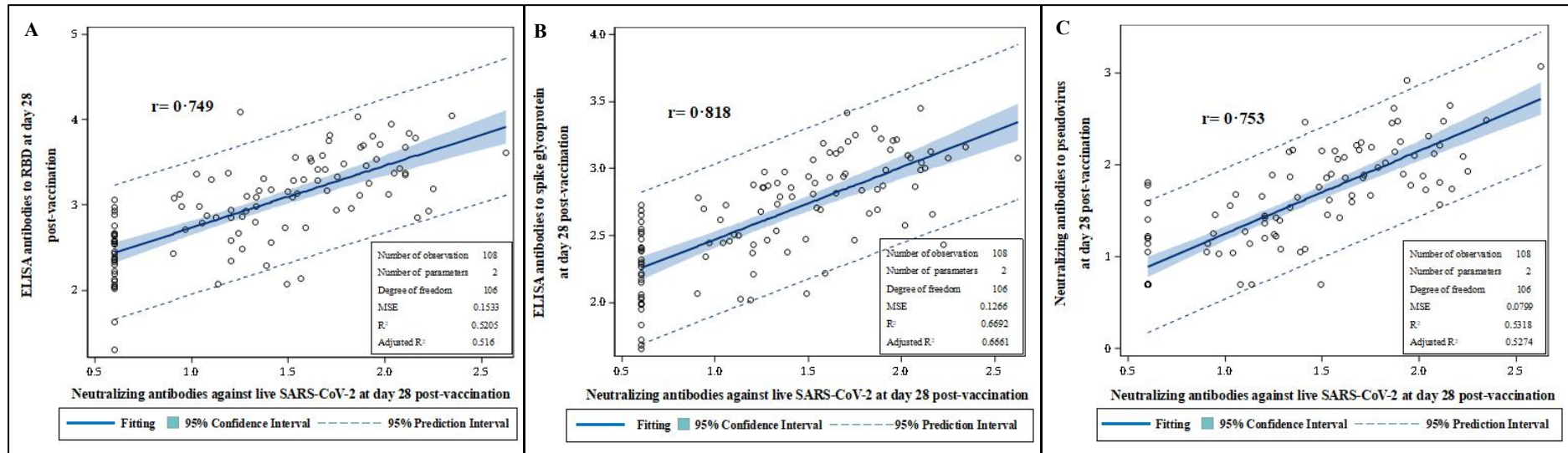
Data are n (%). ALT= alanine aminotransferase. AST= aspartate aminotransferase.

Appendix 5. Specific antibody responses to spike glycoprotein, neutralizing antibodies to pseudovirus at day 14 and 28 post-vaccination.

	Day 14				Day 28			
	Low dose (n=36)	Middle dose (n=36)	High dose (n=36)	P value	Low dose (n=36)	Middle dose (n=36)	High dose (n=36)	P value
ELISA antibodies to spike glycoprotein								
GMT	130.0 (83.0-203.7)	92.9 (64.9-132.9)	199.3 (141.7-280.2)	0.020*	385.1 (284.8-520.5)	345.6 (247.8-482.0)	596.4 (440.0-808.4)	0.034**
≥4-fold increase	15, 42%	22, 61%	25, 69%	0.050†	32, 89%	32, 89%	30, 83%	0.82
Neutralization antibodies to pseudovirus								
GMT	18.5 (11.1-31.1)	16.4 (10.8-24.9)	29.2 (18.9-45.3)	0.17	29.8 (18.4-48.1)	27.3 (16.9-44.3)	45.6 (28.4-73.0)	0.26
≥4-fold increase	16, 44%	17, 47%	21, 58%	0.46	21, 58%	19, 53%	25, 69%	0.34

Data are mean (95%CI) or n, %. n = number of participants. % = proportion of participants. ELISA=enzyme-linked immunosorbent assay. GMT= geometric mean antibody titer. The p values are the result of comparison between the three dose groups. The Student-Newman-Keuls test was used for multicomparison of GMTs. * The difference between high dose and middle dose was 149.2, with 95%CI 10.2~288.1. ** The difference between high dose and low dose was 280.2, with 95%CI 29.8~530.6; and the difference between high dose and low dose was 281.5, with 95%CI 31.1~531.9. † Only a margin of significance was found across the groups, and no significant difference was found after a Bonferroni correction for multicomparisons.

Appendix 6. The association between neutralizing antibodies to live SARS-CoV-2 and ELISA antibodies to receptor binding domain, spike glycoprotein, or pseudovirus-based neutralizing antibodies at day 28 post-vaccination.



RBD= receptor binding domain. Panel A shows the association between binding antibodies to RBD and neutralizing antibodies to live virus at day 28 post-vaccination. Panel B shows the association between binding antibodies to spike glycoprotein and neutralizing antibodies to live SARS-CoV-2 at day 28 post-vaccination. Panel C shows the association between neutralizing antibodies to pseudovirus and live SARS-CoV-2 at day 28 post-vaccination.

Appendix 7. Specific antibody responses to receptor binding domain, spike glycoprotein, neutralizing antibodies to live SARS-CoV-2 or to pseudovirus at day 14 and 28 post-vaccination according to pre-existing Ad5.

	Day 14				Day 28			
	Low dose	Middle dose	High dose	P value	Low dose	Middle dose	High dose	P value
ELISA antibodies to RBD								
Pre-existing Ad5 \leq 200								
n	16	17	20		16	17	20	
GMT	225.4 (92.9-546.8)	224.5 (139.1-362.4)	270.9 (142.5-514.9)	0.89	1013.3 (468.2-2192.7)	1444.7 (882.8-2364.3)	2275.9 (1231.3-4206.6)	0.16
\geq 4-fold increase	12, 75%	14, 82%	16, 80%	0.92	15, 94%	17, 100%	20, 100%	0.30
Pre-existing Ad5 $>$ 200								
n	20	19	16		20	19	16	
GMT	32.2 (21.0-49.4)	40.7 (21.1-78.3)	54.3 (30.9-95.4)	0.39	413.5 (273.4-625.2)	478.2 (259.5-881.0)	820.0 (473.4-1420.5)	0.15
\geq 4-fold increase	4, 20%	4, 21%	6, 38%	0.50	20, 100%	17, 89%	16, 100%	0.20
ELISA antibodies to spike glycoprotein								
Pre-existing Ad5 \leq 200								
n	16	17	20		16	17	20	
GMT	326.5 (191.5-556.8)	164.7 (109.6-247.5)	306.2 (196.4-477.4)	0.064	619.1 (392.6-976.2)	499.8 (324.0-771.0)	818.3 (529.5-1264.4)	0.24
\geq 4-fold increase	11, 69%	14, 82%	15, 75%	0.61	13, 81%	16, 94%	17, 85%	0.60
Pre-existing Ad5 $>$ 200								
n	20	19	16		20	19	16	
GMT	62.3	55.7	116.5	0.073	263.3	248.4	401.6	0.19

	Day 14				Day 28			
	Low dose	Middle dose	High dose	P value	Low dose	Middle dose	High dose	P value
	(37.6-103.2)	(34.1-90.9)	(75.2-180.4)		(186.5-371.9)	(153.3-402.7)	(276.5-583.2)	
≥4-fold increase	4, 20%	8, 42%	10, 63%	0.034*	19, 95%	16, 84%	13, 81%	0.36
Neutralizing antibodies to live SARS-CoV-2								
Pre-existing Ad5≤200								
n	16	17	20		16	17	20	
GMT	16.1	17.3	25.3	0.40	30.6	27.8	50.2	0.26
	(9.0-28.8)	(9.5-31.7)	(15.8-40.6)		(16.9-55.6)	(14.6-53.0)	(30.0-84.0)	
≥4-fold increase	9, 56%	10, 59%	13, 65%	0.86	13, 81%	11, 65%	17, 85%	0.35
Pre-existing Ad5>200								
n	20	19	16		20	19	16	
GMT	4.8	5.7	5.3	0.72	8.0	10.0	20.8	0.031†
	(3.8-5.9)	(3.9-8.5)	(3.5-8.2)		(5.2-12.2)	(5.6-17.7)	(11.2-38.7)	
≥4-fold increase	1, 5%	1, 5%	2, 13%	0.67	5, 25%	7, 37%	10, 63%	0.070
Neutralization antibodies to pseudovirus								
Pre-existing Ad5≤200								
n	16	17	20		16	17	20	
GMT	46.1	34.1	58.9	0.43	61.2	45.5	63.8	0.71
	(19.1-111.5)	(19.5-59.6)	(36.0-96.6)		(31.4-119.3)	(23.4-88.7)	(34.4-118.5)	
≥4-fold increase	11, 69%	12, 71%	18, 90%	0.21	13, 81%	12, 71%	16, 80%	0.77
Pre-existing Ad5>200								
n	20	19	16		20	19	16	
GMT	8.9	8.5	12.2	0.51	16.8	17.3	29.9	0.39
	(5.8-13.9)	(5.4-13.5)	(7.1-20.9)		(9.1-30.7)	(8.8-34.2)	(14.1-63.3)	
≥4-fold increase	5, 25%	5, 26%	3, 19%	0.85	8, 40%	7, 37%	9, 56%	0.47

Data are mean (95%CI) or n, %. n = number of participants. % = proportion of participants. ELISA= enzyme-linked immunosorbent assay. RBD= receptor binding domain. GMT= geometric mean antibody titer. The p values are the result of comparison between the three dose groups. * The difference between high dose and low dose was 0.43%, with 95%CI 0.13%~0.72%. † The difference between high dose and low dose was 23.4, with 95%CI 4.3~42.5.

Appendix 8. Logistic regression analysis of seroconversion rate of neutralizing antibodies to live SARS-CoV-2 at day 28 post-vaccination with the baseline characters, vaccine dose and pre-existing Ad5 antibodies.

Model parameters	Estimates	P value	OR	
			Point estimates	95%CI by Wald
Univariate analysis*				
Middle dose	0.00	>0.99	1.00	0.40, 2.52
High dose	1.10	0.031	3.00	1.11, 8.14
Age	1.17	0.011	3.22	1.30, 7.96
Sex	-0.47	0.23	0.62	0.29, 1.35
Pre-existing Ad5 antibodies (>200)	-1.63	0.0001	0.20	0.08, 0.45
Multivariable analysis†				
Intercept	0.65	0.17		
Middle dose	-0.06	0.91	0.94	0.33, 2.67
High dose	1.02	0.074	2.76	0.91, 8.44
Age	0.68	0.011	3.88	1.37, 11.01
Sex	-0.32	0.17	0.53	0.21, 1.31
Pre-existing Ad5 antibodies (>200)	-1.67	0.0003	0.19	0.08, 0.47

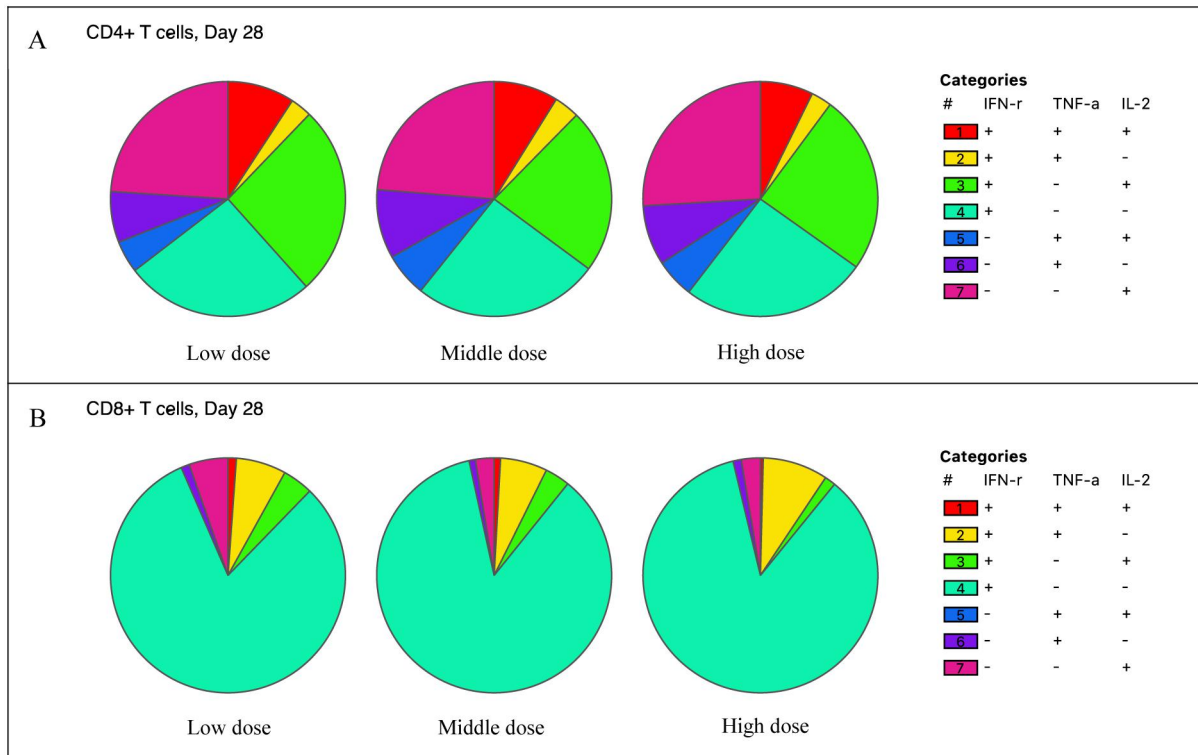
*The dose group (the low-dose group as the reference), age (stratified into 18-44, 45-60; using 45-60 years as reference), sex (female as reference), and the level of Ad5 antibodies pre-vaccination (≤ 200 as reference) are analysed in univariate logistic regression, respectively, before involved in a multivariable analysis. † The independent variables are the dose group (the low-dose group as the reference), age (stratified into 18-44, 45-60; using 45-60 years as reference), sex (female as reference), and the level of Ad5 antibodies pre-vaccination (≤ 200 as reference) included in a multivariable analysis.

Appendix 9. Specific neutralizing antibodies to Ad5 vector at day 14 and 28 post-vaccination according to pre-existing Ad5.

	Day 14				Day 28			
	Low dose (n=36)	Middle dose (n=36)	High dose (n=36)	P value	Low dose (n=36)	Middle dose (n=36)	High dose (n=36)	P value
Total								
GMT	840·6 (332·7-2123·5)	1429·6 (642·5-3181·0)	1208·9 (595·5-2454·0)	0·64	672·0 (254·9-1771·2)	1041·2 (467·6-2318·1)	898·7 (412·6-1957·2)	0·76
Pre-existing Ad5≤200								
GMT<	71·9 (20·7-249·3)	340·6 (82·7-1403·6)	323·7 (130·5-802·6)	0·10	49·3 (14·1-172·4)	238·4 (60·5-939·5)	202·0 (76·4-534·2)	0·12
Pre-existing Ad5>200								
GMT	6008·9 (4537·2-7957·9)	5159·1 (3745·1-7106·9)	6275·9 (4833·1-8149·5)	0·59	5434·1 (4029·6-7328·0)	3893·5 (2625·4-5774·1)	5808·1 (4600·0-7333·6)	0·16

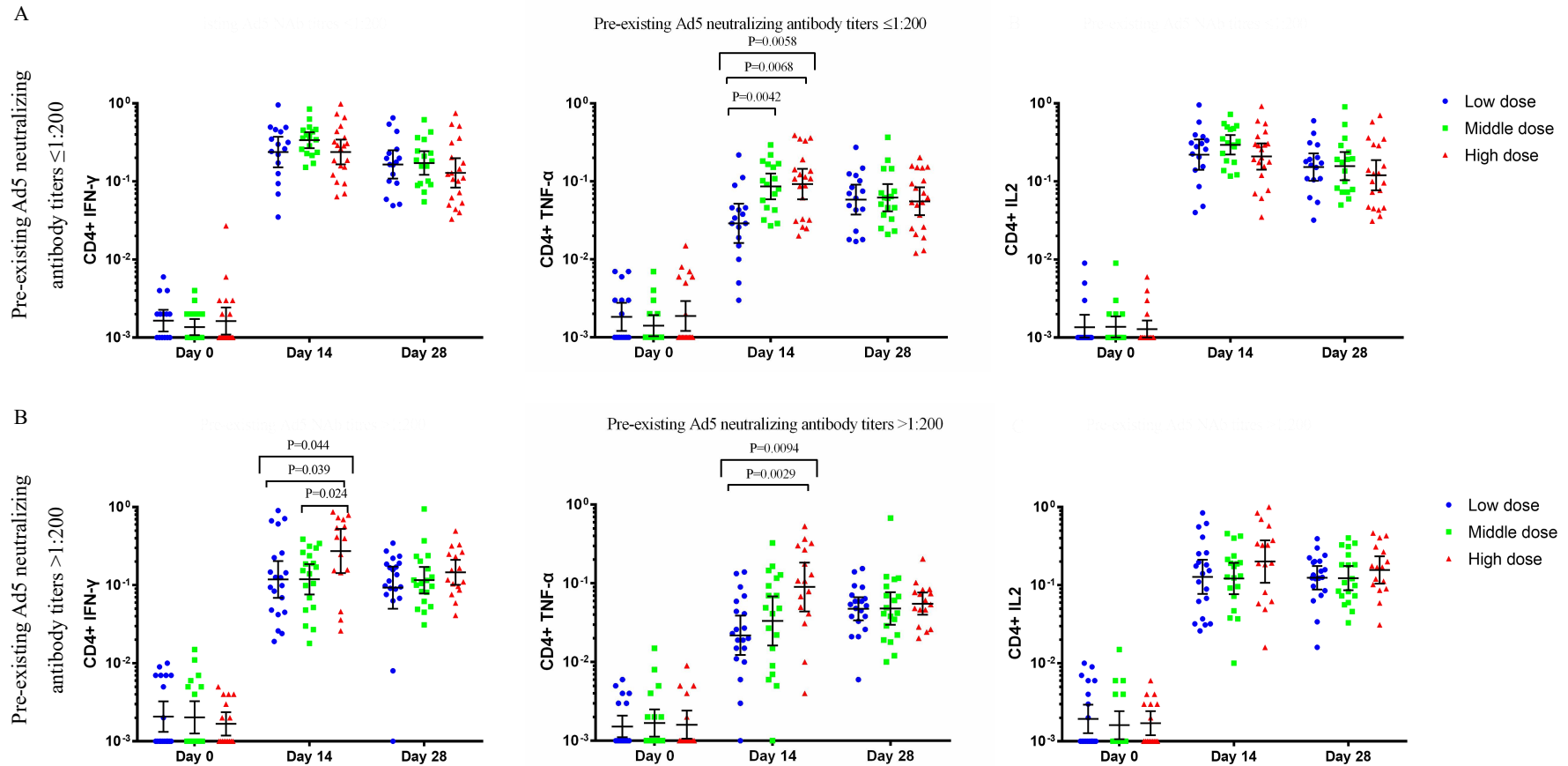
Data are mean (95%CI) or n, %. n = number of participants. % = proportion of participants. Ad5= adenovirus type 5. GMT= geometric mean antibody titer. The p values are the result of comparison between the three dose groups.

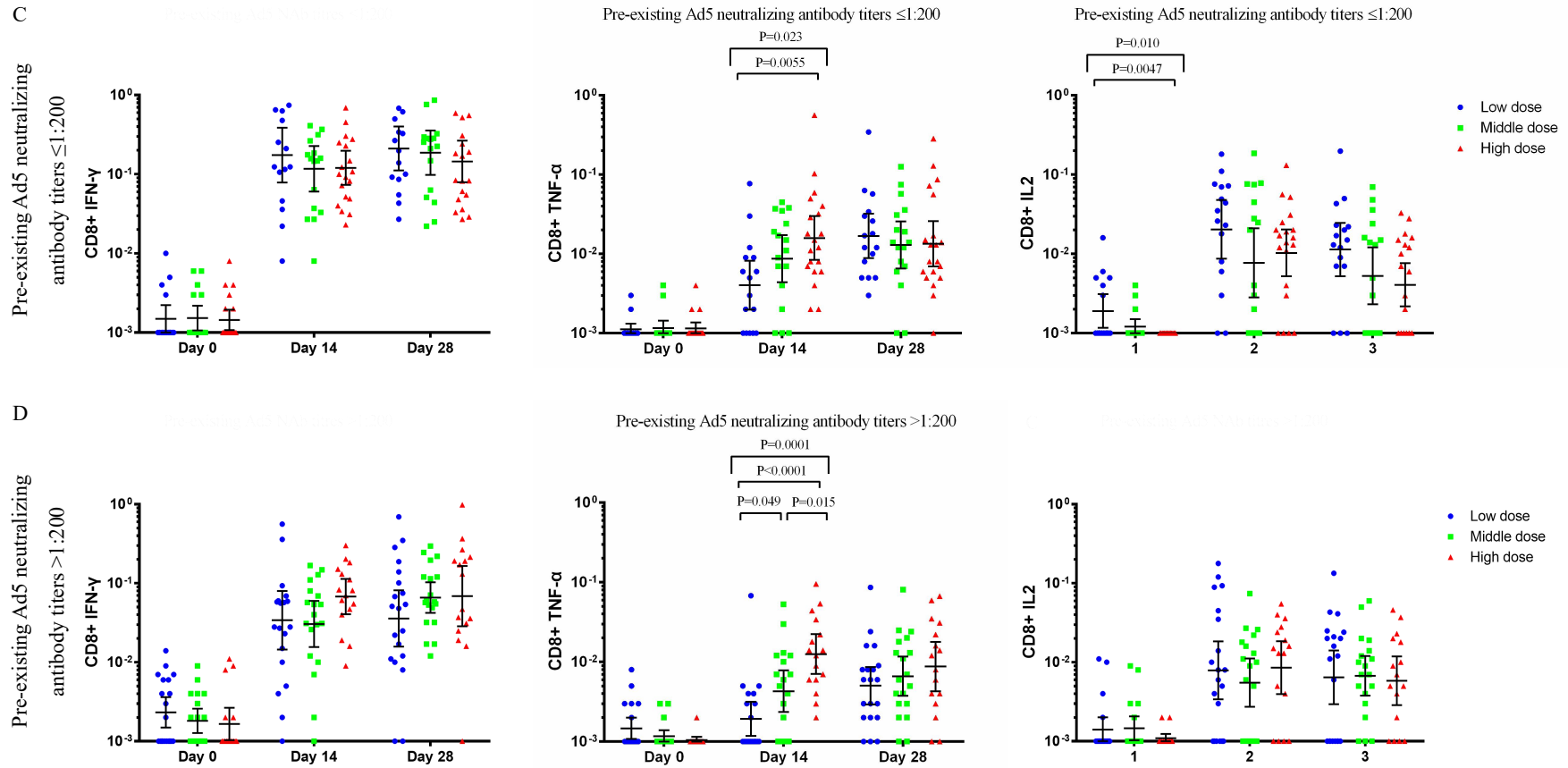
Appendix 10. T cells producing combination of INF- γ , TNF- α , and IL-2 at day 28 post-vaccination measured by flow cytometry.



Panel A shows the proportions of CD4+ T cells producing any combination of INF- γ , TNF- α , and IL-2 at day 28 post-vaccination. Panel B shows the proportions of CD8+ T cells producing any combination of INF- γ , TNF- α , and IL-2 at day 28 post-vaccination.

Appendix 11. Stratified analyses of the T cell responses based on the pre-existing Ad5 neutralizing antibody titers measured by flow cytometry.





Panel A shows the specific T cell response with secretion of IFN- γ , tumor necrosis factor α (TNF- α), and interleukin-2 from CD4+ T cells at day 0, 14, and 28 in participants with low pre-existing Ad5 neutralizing antibody titers ($\leq 1:200$) in three dose groups. Panel B shows the specific T cell response with secretion of IFN- γ , TNF- α , and interleukin-2 from CD4+ T cells at day 0, 14, and 28 in participants with high pre-existing Ad5 neutralizing antibody titers ($> 1:200$) in three dose groups. Panel C shows the specific T cell response with secretion of IFN- γ , TNF- α , and interleukin-2 from CD8+ T cells at day 0, 14, and 28 in participants with low pre-existing Ad5 neutralizing

antibody titers ($\leq 1:200$) in three dose groups. Panel D shows the specific T cell response with secretion of IFN- γ , TNF- α , and interleukin-2 from CD8⁺ T cells at day 0, 14, and 28 in participants with high pre-existing Ad5 neutralizing antibody titers ($>1:200$) in three dose groups.

Appendix 12. Proportion of participants with either seroconversion of neutralizing antibodies to live SARS-CoV-2 or a positive T cell response measured by INF- γ ELISpot at day 14 and 28 post-vaccination.

		Low dose (n=36)	Middle dose (n=36)	High dose (n=36)	Total (n=108)
Day 14 post-vaccination					
Seroconversion	n (%)	30 (83)	35 (97)	35 (97)	100 (93)
Seroconversion rate CI	95%CI	68.1, 92.1	85.8, 99.5	85.8, 99.5	86.1, 96.2
Day 28 post-vaccination					
Seroconversion	n (%)	28 (78)	33 (92)	36 (100)	97 (90)
Seroconversion rate CI	95%CI	61.9, 88.3	78.2, 97.1	90.4, 100.0	82.7, 94.2

n (%), n = number of participants. % = proportion of participants. A positive antibody response (seroconversion) was defined as at least a 4-fold increase in post-vaccination titer from baseline. The ELISpot T cell responses were considered positive if at least 2-fold increase in the numbers of IFN- γ -secreting T cells was found post-vaccination compared to the baseline.