

Immunodominant SARS Coronavirus Epitopes in Humans Elicited both Enhancing and Neutralizing Effects on Infection in Non-Human Primates

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Supplements

Table S1. Neutralization and enhancement of SARS-CoV infection of Vero E6 cells in the presence of human antisera of convalescent SARS patients*

#Enhancement of SARS-CoV infection						#Blockage of SARS-CoV infection	
SARS patient		SARS patient		SARS patient		SARS patient	
code	Neutralization	code	Neutralization	Code	Neutralization	code	Neutralization
W1	-117.00% ± 13.73	W49	-53.00% ± 11.14	S204	-58.00% ± 9.98	W33	29.00% ± 6.28
W9	-92.00% ± 10.15	W61	-49.00% ± 16.08	S101	-91.00% ± 8.50	W29	48.00% ± 6.00
W35	-100.00% ± 11.56	W53	-53.00% ± 8.91	S245	-46.00% ± 7.69	W73	50.00% ± 7.77
W38	-65.00% ± 8.41	S102	-43.00% ± 1.95	S98	-62.00% ± 12.58	S100	25.00% ± 11.92
W41	-48.00% ± 9.82	S196	-46.00% ± 10.29			3E7	36.00% ± 2.27
W45	-42.00% ± 10.92	S241	-50.00% ± 9.35			1G4	56.00% ± 0.77

*Total 116 antisera (1:500 dilution) were tested in the neutral red staining(NRS) assay.

#For neutralization, six antisera with > 20% inhibition were defined as significantly blocking SARS-CoV infection; for enhancement, 17 antisera with > 40% augmentation were defined as significantly enhancing SARS-CoV infection.

Table S2 Design and synthesized new peptides

Novel peptides from the Spike glycoprotein

S11–38: TSGSDLRCTTFDDVQAPNYTQHTSSMR

S119–132: NSTNVVIRACNFEL

S138–158: FAVSKPMGTQHTMIFDNAFN

S159–184: CTFEYISDAFSLDVSEKSGNFKHLRE

S162–180: EYISDAFSLDVSEKSGNFK

S188–211: KNKDGFLYVYKGYQPIDVVRDLPSG

S342–368: RKKISNCVADYSVLYNSTFFSTFKCYG (Disulfide)

S391–404: GDDVRQIAPGGTGV

S416–434: FMGCVLAWNTRNIDATSTG

S436–459: YNYKYRYLRHGKLRPFERDISNVP

S471–490: ALNCYWPLNDYGFYTTTGIG

S491–502: YQPYRVVVLSE

S522–535: NQCVNFNFNGLTGT

S597–625: LYQDVNCTDVSTAIHADQLTPAWRIYSTG

S679–695: MSLGADSSIAYSNNTIA

S686–713: SIAYSNNTIAIPTNFSISITTEVMPVSM

S803–828: LLFNKVTLADAGFMKQYGECLGDINA

S946–960: KQLSSNFGAISSVLN

S965–986: RLDKVEAEVQIDRLITGRLQSL

S993–1012: QLIRAAEIRASANLAATKM

S1084–1110: WFITQRNFFSPQIITTD

S1138–1154: FKNHTSPDVDLGDIG

S1155–1178: NASVVNIQKEIDRLNEVAKNLNE

Table S3. Generated anti-peptide mAbs * NF=not functional.

No.	Clone no.	Isotype	Function*	Binding affinity to the peptide	
1	S ₄₇₁₋₅₀₃	2F7	IgG1 κ	NF	1.10E-09
2		3E10	IgG1 κ	NF	2.95E-09
3		4E5	IgG1 κ	Neutralizing	4.67E-09
4		4A10	IgG1 κ	Neutralizing	2.96E-09
5		6G5	IgG2b κ	Neutralizing	5.82E-09
6		6A10	IgG1 κ	NF	7.57E-09
7		9A6	IgG1 κ	NF	4.94E-09
8	S ₆₀₄₋₆₂₅	2B4	IgG1 κ	NF	4.75E-06
9		3B10	IgG1 κ	NF	2.17E-07
10		4H11	IgG1 κ	NF	1.68E-08
11		9A6	IgG1 κ	NF	2.62E-09
12		3A2	IgG2a κ	NF	8.16E-09

13		11B1	IgG2a	κ	Neutralizing	2.32E-09
14		11B3	IgG2a	κ	NF	6.94E-08
15		14F1	IgG2b	κ	NF	1.58E-06
16		6D5	IgG1	κ	NF	8.77E-09
17	S ₅₉₇₋₆₂₅	37-9-5	IgG1	κ	NF	3.94E-10
18		43-3-14	IgG1	κ	Enhancing	1.21E-11
19		65-5-10	IgG1	κ	NF	2.21E-10
20		219-10-28	IgG1	κ	Enhancing	1.20E-08
21	S ₁₁₆₄₋₁₁₉₁	126-10	IgG1	κ	Neutralizing	4.59E-9
22		581-39	IgG1	κ	NF	1.56E-9
23		608-25	IgG1	κ	NF	1.68E-9

Table S4 Alanine walking peptides

L597A: AYQDVNCTDVSTAIHADQLTPAWRIYSTG

Y598A: LAQDVNCTDVSTAIHADQLTPAWRIYSTG

Q599A: LYADVNCTDVSTAIHADQLTPAWRIYSTG

D600A: LYQAVNCTDVSTAIHADQLTPAWRIYSTG

V601A: LYQDANCTDVSTAIHADQLTPAWRIYSTG

N602A: LYQDVACTDVSTAIHADQLTPAWRIYSTG

C603A: LYQDVNATDVSTAIHADQLTPAWRIYSTG

T604A: LYQDVNCADVSTAIHADQLTPAWRIYSTG

D605A: LYQDVNCTAVSTAIHADQLTPAWRIYSTG

V606A: LYQDVNCTDASTAIHADQLTPAWRIYSTG

mAb43-3-14 specifically binding short peptides

Leu-6-Asn(S597–602): LYQDVN

Leu-7-Cys(S597–603): LYQDVNC

Leu-8-Thr(S597–604): LYQDVNCT

Table S5 Monkeys for peptide vaccine immunization against SARS-CoV

Vac1 (Control group)			Vac2 (MAP-S₅₉₇₋₆₂₅ formulated with FCA/IFA)			Vac3 (MAP-S₄₇₁₋₅₀₃, MAP-S₆₀₄₋₆₂₅, MAP-S₁₁₆₄₋₁₁₉₁ formulated with FCA/IFA)			Vac4 (MAP-S₄₇₁₋₅₀₃, MAP-S₅₉₇₋₆₂₅, MAP-S₁₁₆₄₋₁₁₉₁ formulated with FCA/IFA)		
Animal code	Gender	Body Weight (kg)	Animal code	Gender	Body weight (kg)	Animal Code	Gender	Body weight (kg)	Animal code	Gender	Body weight (kg)
041870 (A1)	F	5.2	040888 (B1)	F	5.6	041057 (C1)	M	6.1	041642 (D1)	F	4.6
031860 (A2)	F	4.8	040390 (B2)	F	5.2	030874 (C2)	F	5.5	030177 (D2)	M	5.4
040995 (A3)	M	6.0	040085 (B3)	M	6.5	040203 (C3)	M	5.1	041399 (D3)	M	4.8
041911 (A4)	M	4.3	031519 (B4)	M	5.0	031632 (C4)	F	6.2	041902 (D4)	F	6.1

	040669 M 5.5 (B5)	041811 M 5.1 (C5)	041659 M 4.5 (D5)
	040830 F 4.8 (B6)	041962 F 4.5 (C6)	041326 F 6.0 (D6)

Table S6 Vaccinated animals sacrificed at 2 DPI

Vac1		Vac2		Vac3		Vac4	
Animal code	Gender	Animal code	Gender	Animal code	Gender	Animal code	Gender
041870 (A1)	F	040888 (B1)	F	041057 (C1)	M	041642 (D1)	F
031860 (A2)	F	040390 (B2)	F	030874 (C2)	F	030177 (D2)	M
		040085 (B3)	M	040203 (C3)	M	041399 (D3)	M

Table S7 Vaccinated animals sacrificed at 6 DPI

Vac1		Vac2		Vac3		Vac4	
Animal code	Gender	Animal code	Gender	Animal code	Gender	Animal code	Gender
040995 (A3)	M	041519 (B4)	M	031632 (C4)	F	041902 (D4)	F
041911 (A4)	M	040669	M	041811	M	041659	M

		(B5)		(C5)		(D5)	
		040830 (B6)	F	041962 (C6)	F	041326 (D6)	F

Table S8 Pathologic classification of the severity of the lung damage in SARS-CoV-infected rhesus macaques

Grade	Pathologic classification of the severity of the lung damage in SARS-CoV-infected rhesus macaques
–	normal macaque lung section without SARS-CoV infection
±	minor inflammation, slight alveolar septal broadening and sparse monocyte infiltration
I	apparent inflammation, hemorrhage in septa, elastic fibers of alveolar wall distorted as shown by silver staining
II	early symptoms of acute diffuse alveolar damage (DAD), alveolar septal broadening with increasing infiltration of inflammatory cells
III	typical symptoms of acute DAD, extensive exudation and septal broadening, shrinkage of alveoli caused by pressure, restricted fusion of thick septa, ruptured elastic fibers of alveoli, variably filled with protein-rich edema fluid, fibrin, erythrocytes, cellular debris, and a moderate number of inflammatory cells in alveolar cavities
IV	severe acute DAD, massive cell infiltration and alveolar shrinking, sheets of septal fusion, necrotic lesions at the hemorrhagic septa and massive cell numbers in alveolar cavities

Table S9 Monkeys treated by mAb43-3-14 were sacrificed at 2 DPI or 6 DPI

	Control group(0.9% NaCl)			0.2mg/kg group			1.8mg/kg group		
	Group	Animal code	Body weight(kg)	Group	Animal code	Body weight(kg)	Group	Animal code	Body weight(kg)
2 DPI	A1	061920	3.3	B1	041986	4.2	D1	051848	3.85
	A2	050426	4.1	B2	040654	3.8	D2	051976	3.9
	A3	051926	3.6	B3	060512	3.15	D3	060578	3.3
6 DPI	A4	060090	3.5	B4	051192	4.1	D4	060060	3.7
	A5	051804	3.9	B5	051820	3.6	D5	040478	3.8
	A6	050104	4.0	B6	051282	3.8	D6	051144	4.1

Table S10 The conditions for preparation of multiple antigen peptides

MAP\Conditions	Lysine core/peptide amount	Reaction solvent	Reaction pH/Reaction time	Reaction monitoring gradient	Preparation condition	Molecular weight (Da, calculated/ observed)	Amount (Yield)	Purity
MAP-S ₆₀₄₋₆₂₅	0.4 mg (0.42 mmol)/5.1 mg (2.00 mmol)	CH ₃ CN/H ₂ O (25/75, v/v)	7.8 (saturated Na ₂ CO ₃ adjusting), 2 h	0–25% CH ₃ CN for 8 min, 25–45% CH ₃ CN for 50 min	Vydac (208TP510), 0–25% CH ₃ CN for 5 min, 25–40% CH ₃ CN for 50 min	10655.83/1 0656.1072	2.1 mg (40%)	98%
MAP-S ₅₉₇₋₆₂₅	0.35 mg (0.37 mmol)/5.0 mg (1.54	CH ₃ CN/H ₂ O (50/50, v/v)	8.5 (saturated Na ₂ CO ₃ adjusting), 5 h	0–25% CH ₃ CN for 8 min, 25–45% CH ₃ CN for	Vydac (214TP1022) , 0–25% CH ₃ CN for 5	13581.83/1 3587.2338	2.5 mg (50%)	95%

	mmol)			50 min	min, 25–40% CH ₃ CN for 50 min			
<i>MAP-S₄₇₁₋₅₀₃</i>	0.30 mg (0.31 mmol)/5.0 mg (1.29 mmol)	CH ₃ CN/H ₂ O/DMSO (40/50/10, v/v/v)	8.5 (saturated Na ₂ CO ₃ adjusting), overnight, or ultrasonicatio n	0–40% CH ₃ CN for 8 min, 40–55% CH ₃ CN for 50 min	Vydac (214TP1022), 0–25% CH ₃ CN for 5 min, 25–40% CH ₃ CN for 50 min	16084.23/1 6091.4292	1.5 mg (30%)	96%
<i>MAP-S₁₁₆₄₋₁₁₉₁</i>	0.35 mg (0.37 mmol)/5.0 mg (1.54 mmol)	CH ₃ CN/H ₂ O (25/75, v/v)	7.8 (saturated Na ₂ CO ₃ adjusting), 2 h	0–25% CH ₃ CN for 8 min, 25–45% CH ₃ CN for 50 min	Vydac (208TP510), 0–35% CH ₃ CN for 5 min, 35–45% CH ₃ CN for 50 min	14389.03/1 4392.1516	2.1 mg (42%)	98%

Chart S1. The protocol of immunization of rhesus monkeys

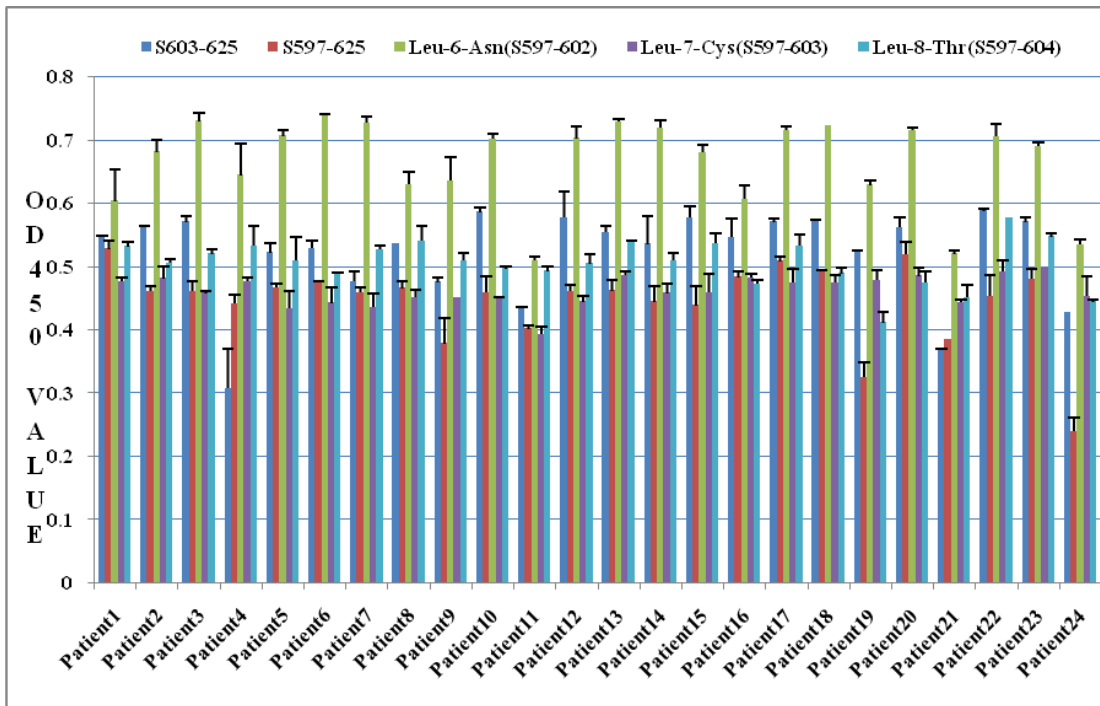
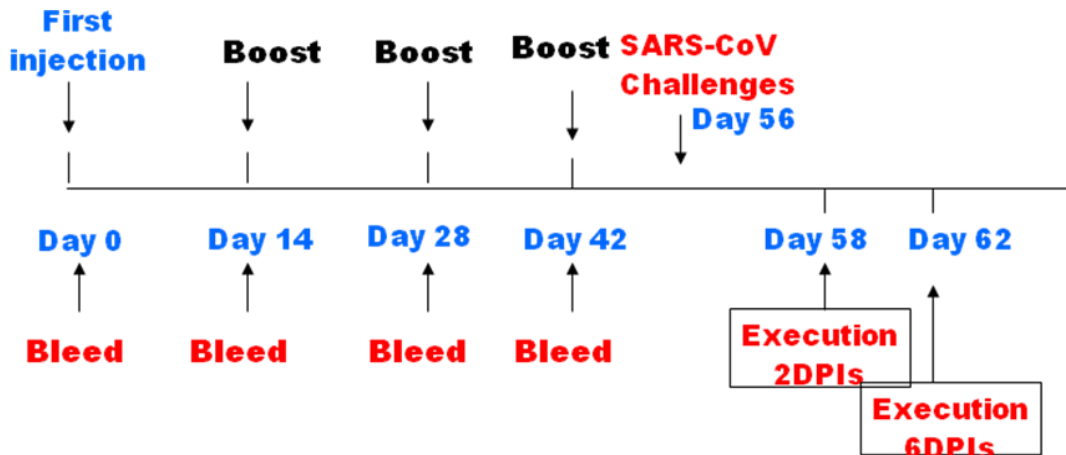


Fig. S1 The serologic reactivity of peptides with the human IgG of convalescent SARS patients.

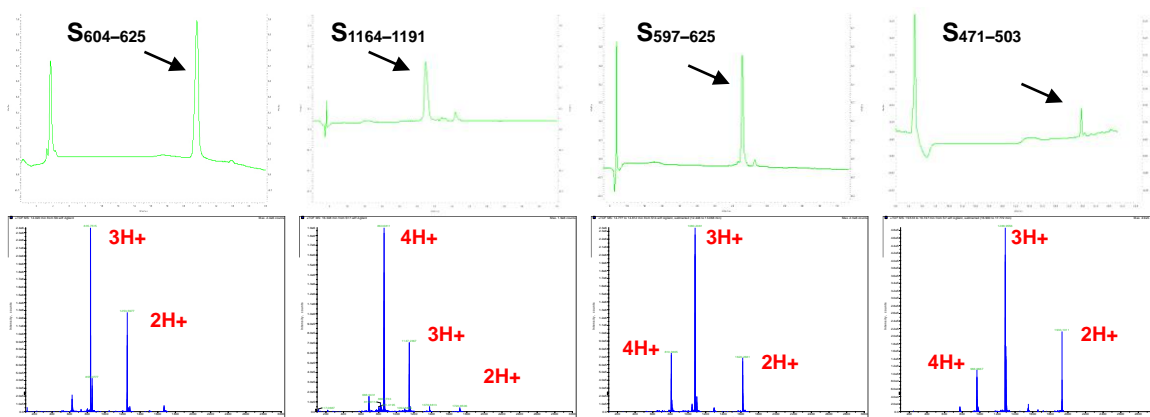


Fig. S2 Characterization of S₆₀₄₋₆₂₅, S₁₁₆₄₋₁₁₉₁, S₅₉₇₋₆₂₅ and S₄₇₁₋₅₀₃. RP-HPLC profiles and MS data.

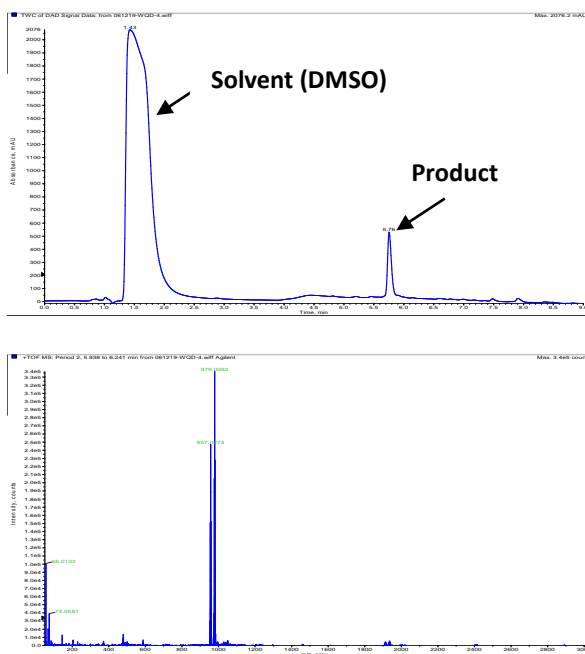


Fig. S3 Characterization of lysine core by RP-HPLC analysis and mass spectrum.

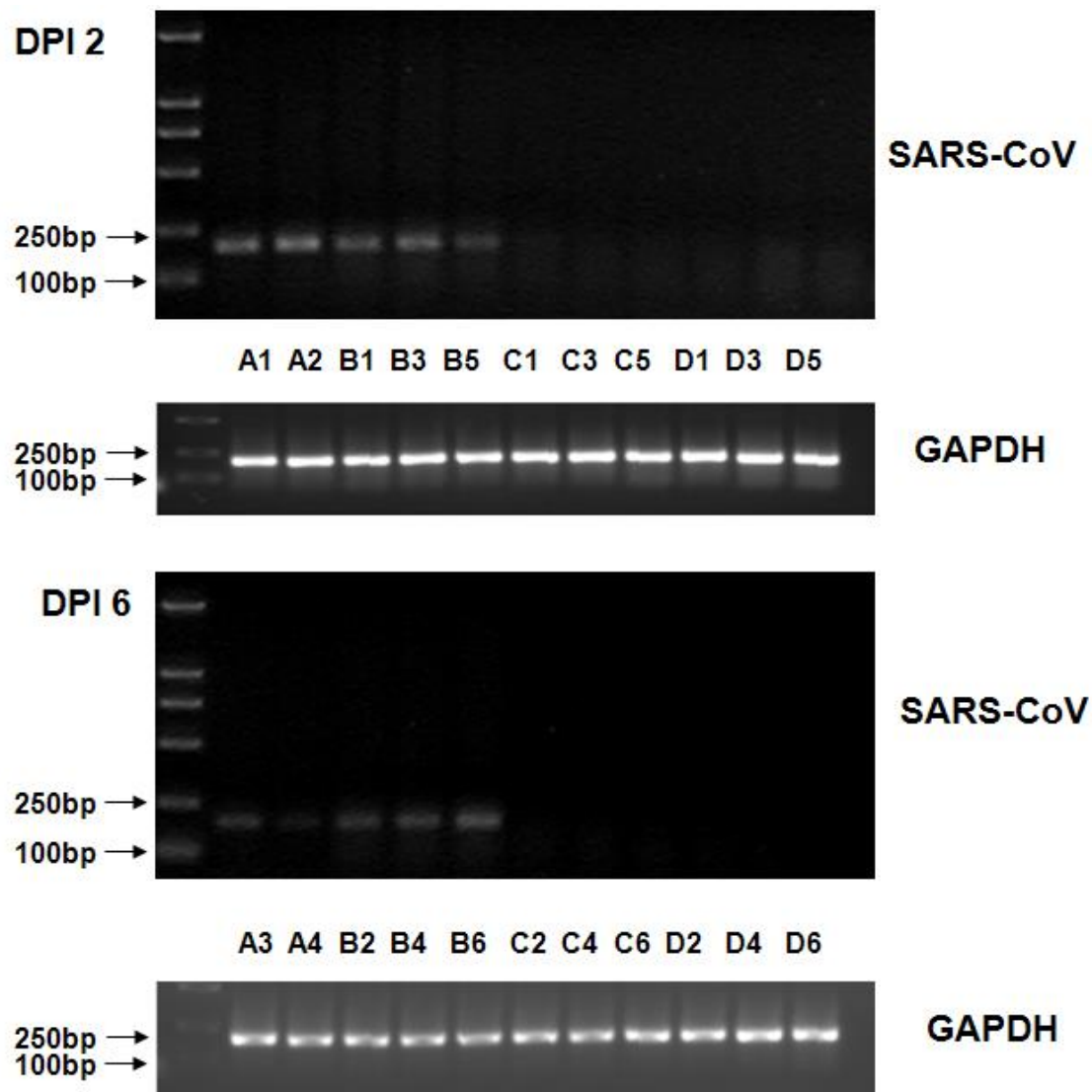


Fig. S4 Lung tissue viral burden analyzed by RT-PCR (A1-A4: Vac1 group, B1-B6: Vac 2 group, C1-C6: Vac 3 group, D1-D6: Vac 4 group)

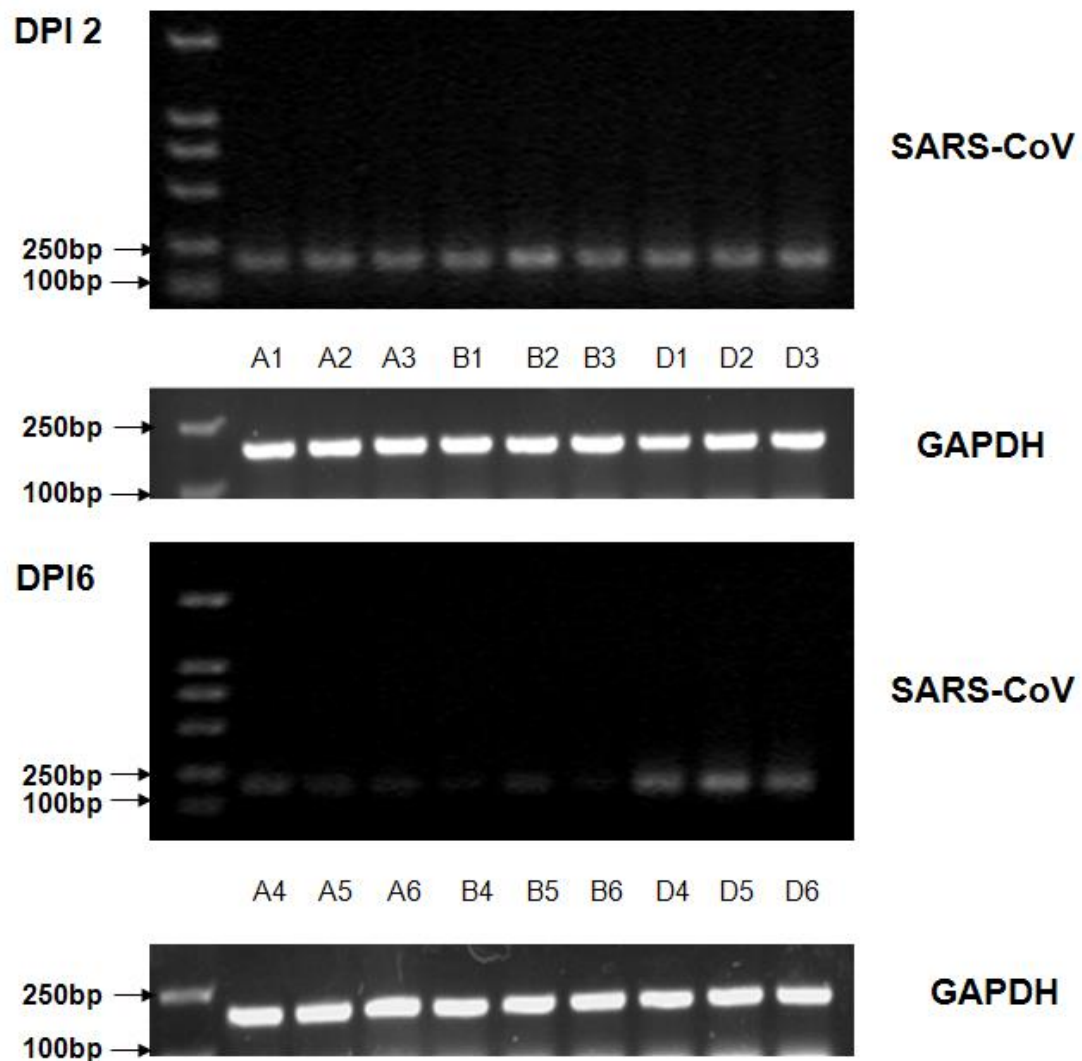


Fig. S5 Lung tissue viral burden analyzed by RT-PCR (A1-A6: 0.9% NaCl group, B1-B6: 0.2mg/kg group, C1-C6: 1.8mg/kg group)