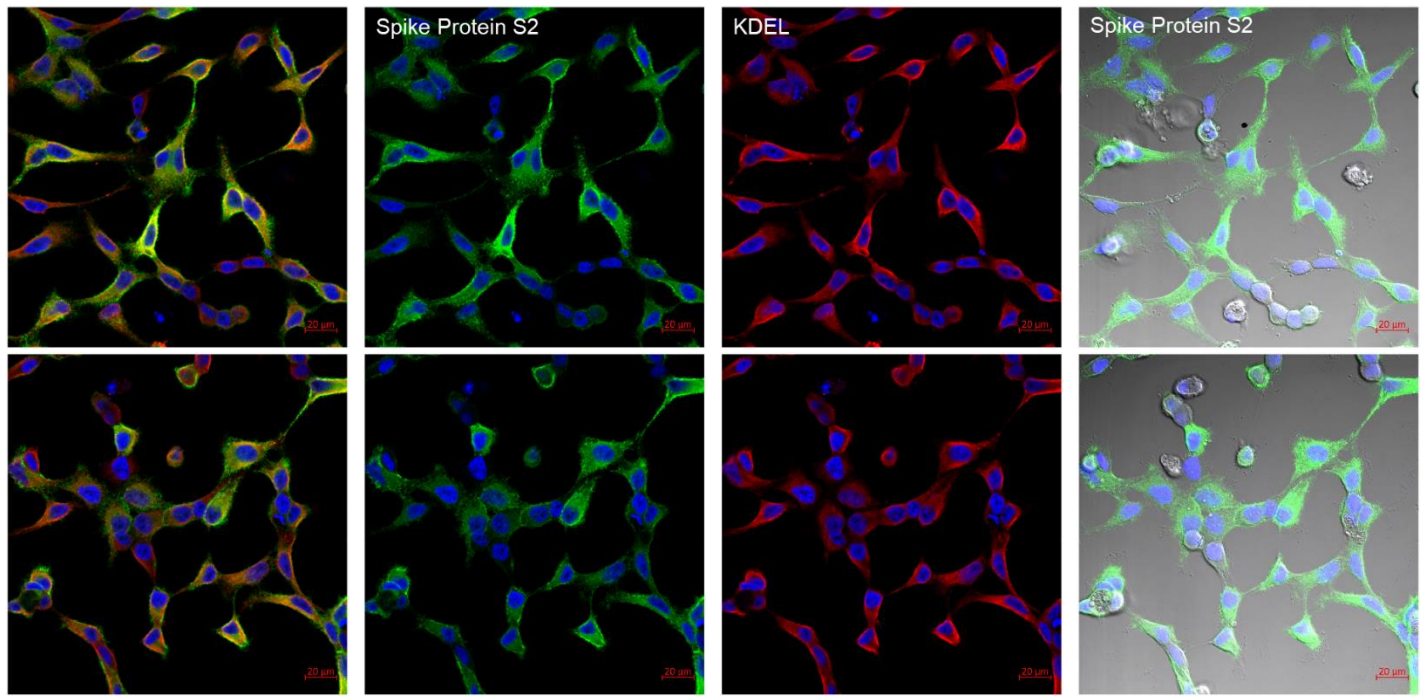


1 **Supplementary Fig.1: Intracellular localization S antigen expressed in HeLa cells**  
2 **transformed with WT mRNA.** Cells transformed with WT mRNA were fixed, permeabilized  
3 and stained with antibodies for Spike protein (green) and KDEL (red), an endoplasmic reticulum  
4 marker. Samples were then examined using confocal microscopy. The left column represents an  
5 overlay of Spike protein and KDEL, while the right column shows the spike protein signal  
6 superimposed on a transmitted light channel.

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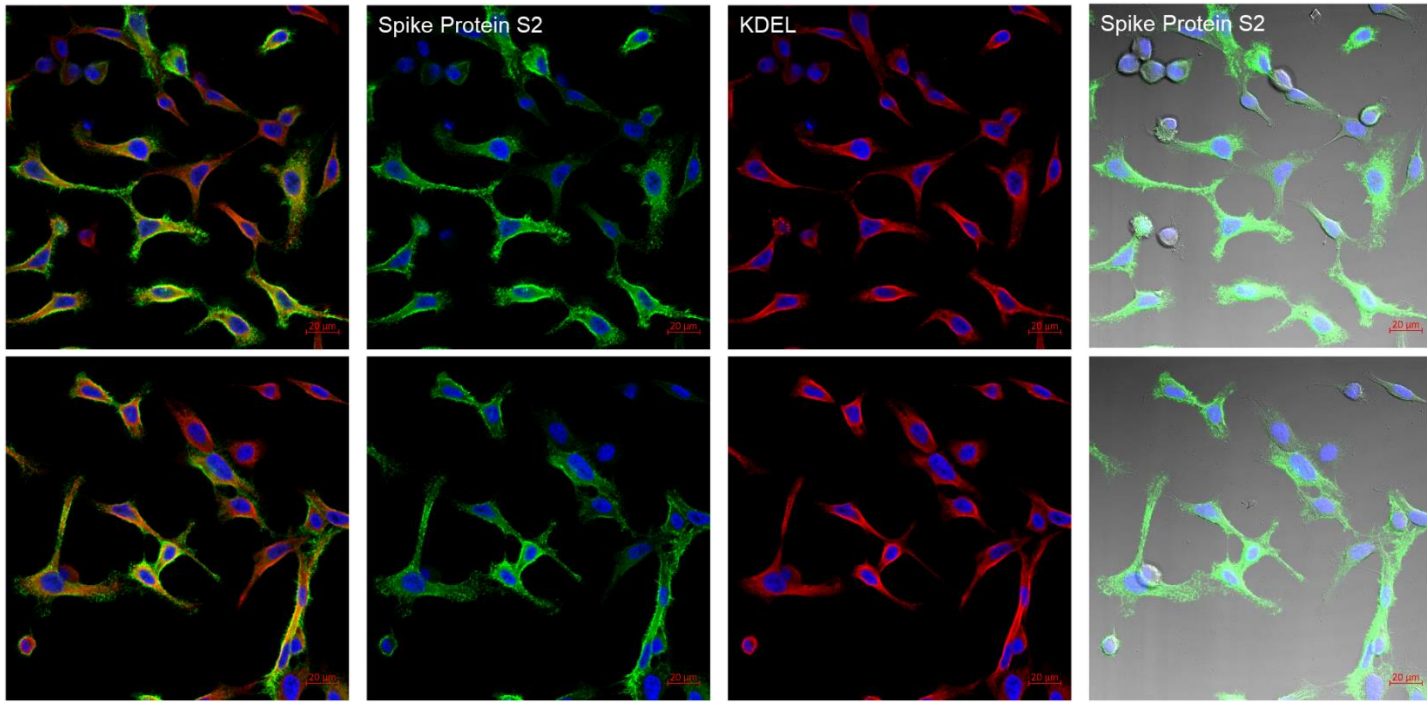


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10 **Supplementary Fig. 2: Intracellular localization S antigen expressed in HeLa cells**  
11 **transformed with 2P mRNA.** Cells transformed with 2P mRNA were fixed, permeabilized and  
12 stained with antibodies for Spike protein (green) and KDEL (red), an endoplasmic reticulum  
13 marker. Samples were then examined using confocal microscopy. The left column represents an  
14 overlay of Spike protein and KDEL signals, while the right column shows the spike protein  
15 signal superimposed on a transmitted light channel.

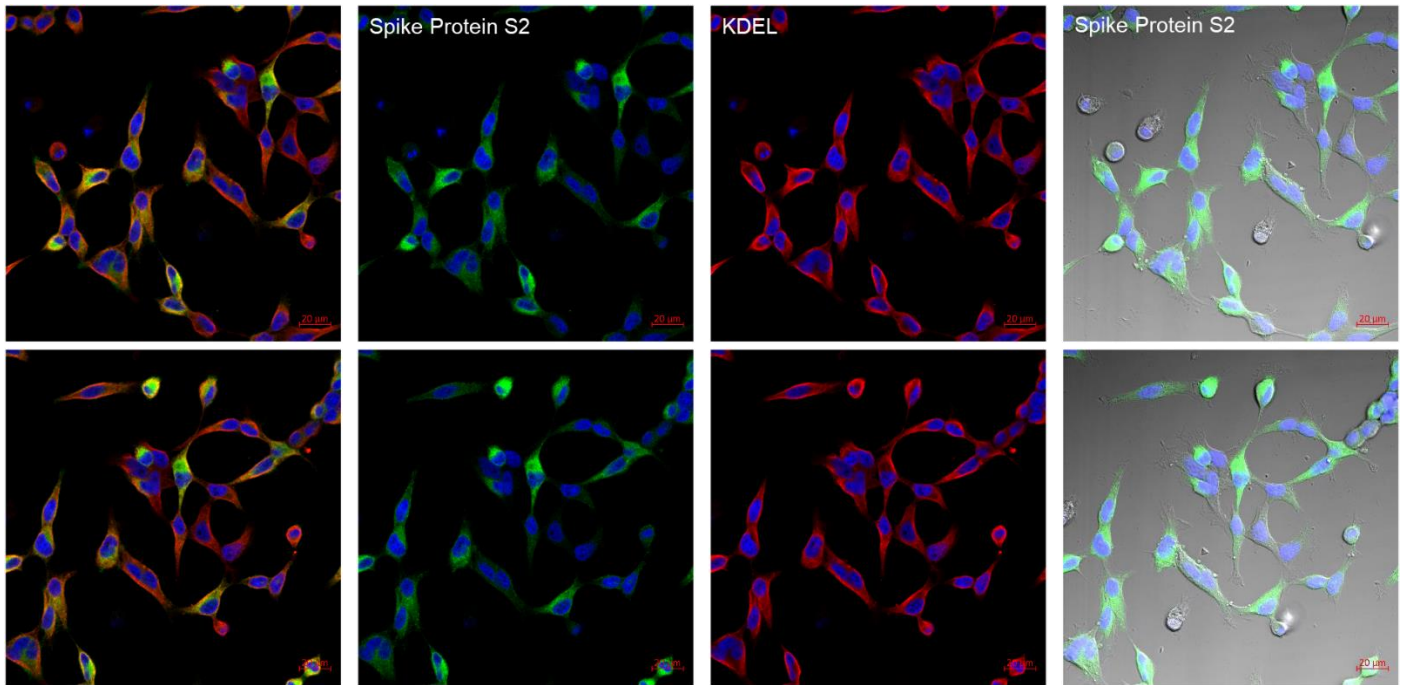
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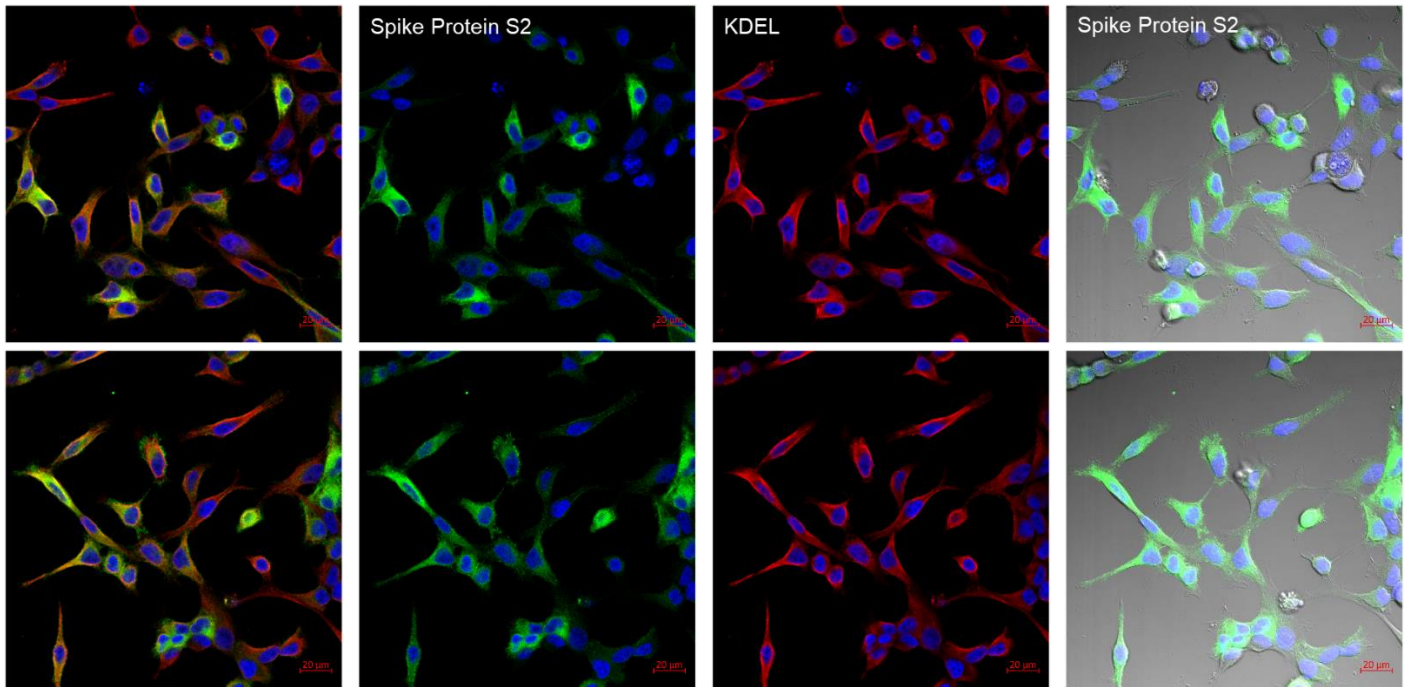
19 **Supplementary Figure 3: Intracellular localization S antigen expressed in HeLa cells**  
20 **transformed with GSAS mRNA.** Cells transformed with GSAS mRNA were fixed,  
21 permeabilized and stained with antibodies for Spike protein (green) and KDEL (red), an  
22 endoplasmic reticulum marker. Samples were then examined using confocal microscopy. The  
23 left column represents an overlay of Spike protein and KDEL signals, while the right column  
24 shows the spike protein signal superimposed on a transmitted light channel.

25



28 **Supplementary Fig. 4: Intracellular localization S antigen expressed in HeLa cells**  
29 **transformed with 2P/GSAS mRNA.** Cells transformed with 2P/GSAS mRNA were fixed,  
30 permeabilized and stained with antibodies for Spike protein (green) and KDEL (red), an  
31 endoplasmic reticulum marker. Samples were then examined using confocal microscopy. The  
32 left column represents an overlay of Spike protein and KDEL signals, while the right column  
33 shows the spike protein signal superimposed on a transmitted light channel.

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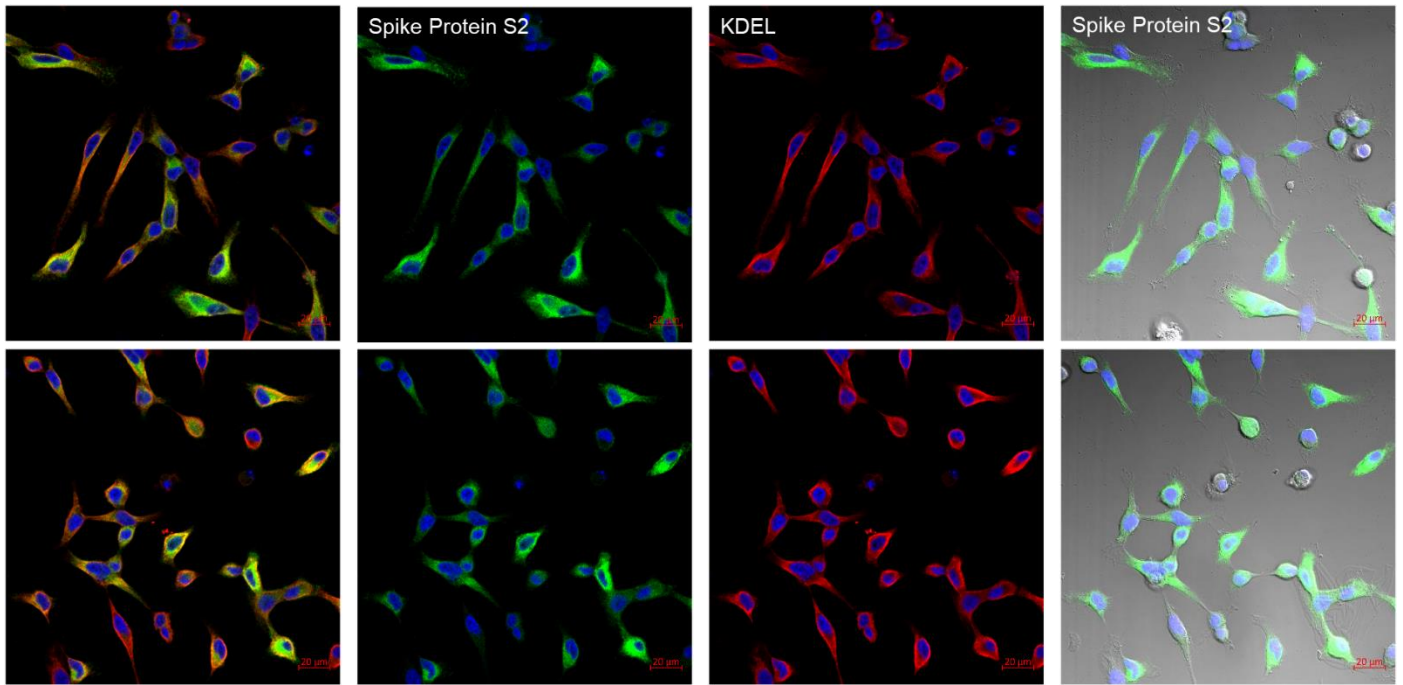
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37 **Supplementary Fig. 5: Intracellular localization S antigen expressed in HeLa cells**  
38 **transformed with 2P/GSAS/ALAYT mRNA.** Cells transformed with 2P/GSAS/ALAYT  
39 mRNA were fixed, permeabilized and stained with antibodies for Spike protein (green) and  
40 KDEL (red), an endoplasmic reticulum marker. Samples were then examined using confocal  
41 microscopy. The left column represents an overlay of Spike protein and KDEL signals, while the  
42 right column shows the spike protein signal along superimposed on a transmitted light channel.

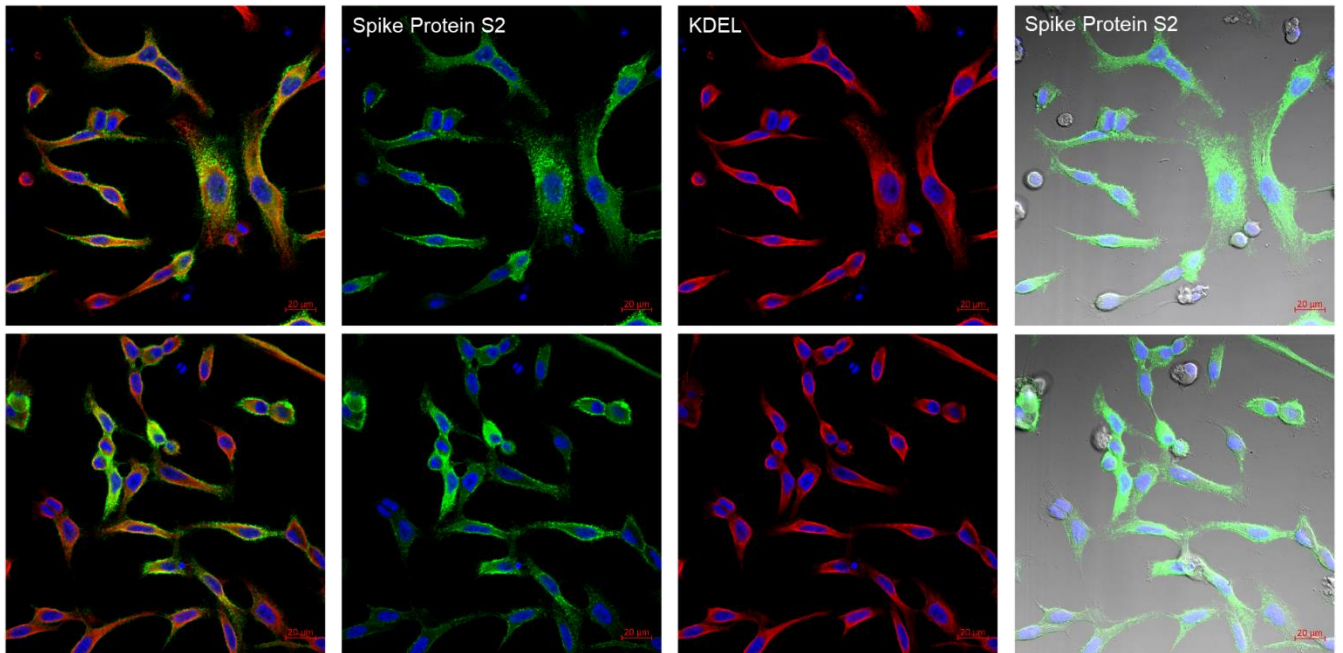
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46 **Supplementary Fig. 6: Intracellular localization S antigen expressed in HeLa cells**  
47 **transformed with 6P mRNA** Cells transformed with 6P mRNA were fixed, permeabilized and  
48 stained with antibodies for Spike protein (green) and KDEL (red), an endoplasmic reticulum  
49 marker. Samples were then examined using confocal microscopy. The left column represents an  
50 overlay of Spike protein and KDEL signals, while the right column shows the spike protein  
51 signal superimposed on a transmitted light channel.

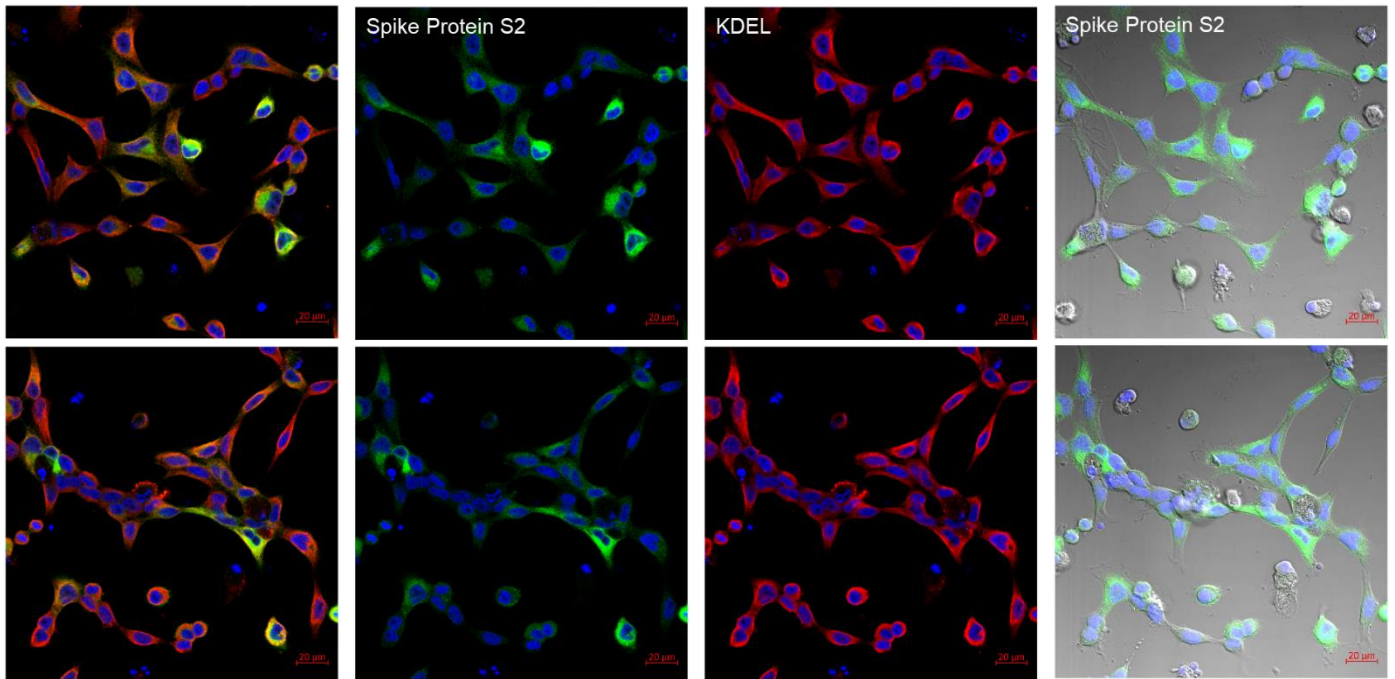
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55 **Supplementary Fig. 7: Intracellular localization S antigens expressed in HeLa cells**  
56 **transformed with 6P/GSAS mRNA.** Cells transformed with 6P/GSAS mRNA were fixed,  
57 permeabilized and stained with antibodies for Spike protein (green) and KDEL (red), an  
58 endoplasmic reticulum marker. Samples were then examined using confocal microscopy. The  
59 left column represents an overlay of Spike protein and KDEL signals, while the right column  
60 shows the spike protein signal superimposed on a transmitted light channel.

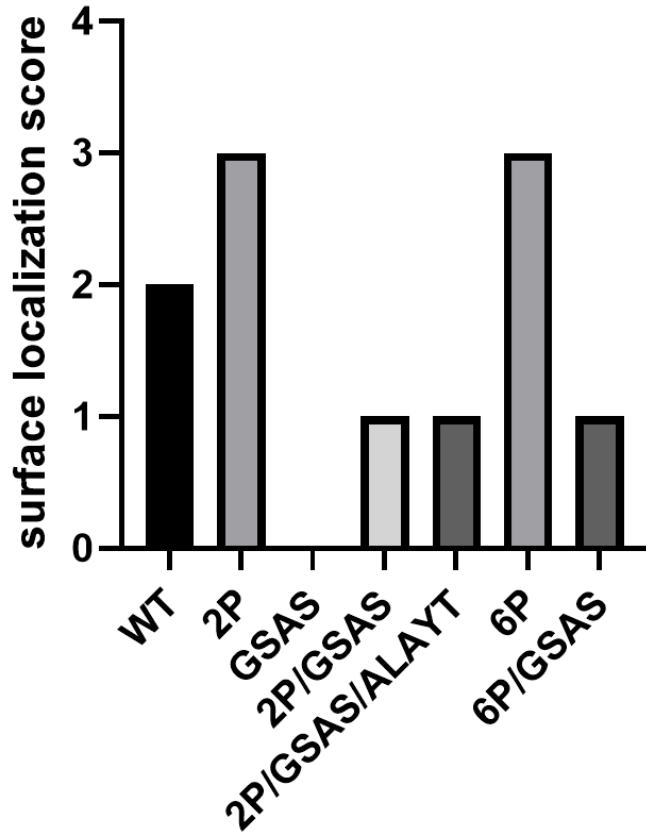
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64 **Supplementary Fig. 8: Relative degree (score) of surface localization of S-proteins based on**  
65 **visual inspection of HeLa cells transformed with mRNAs.**

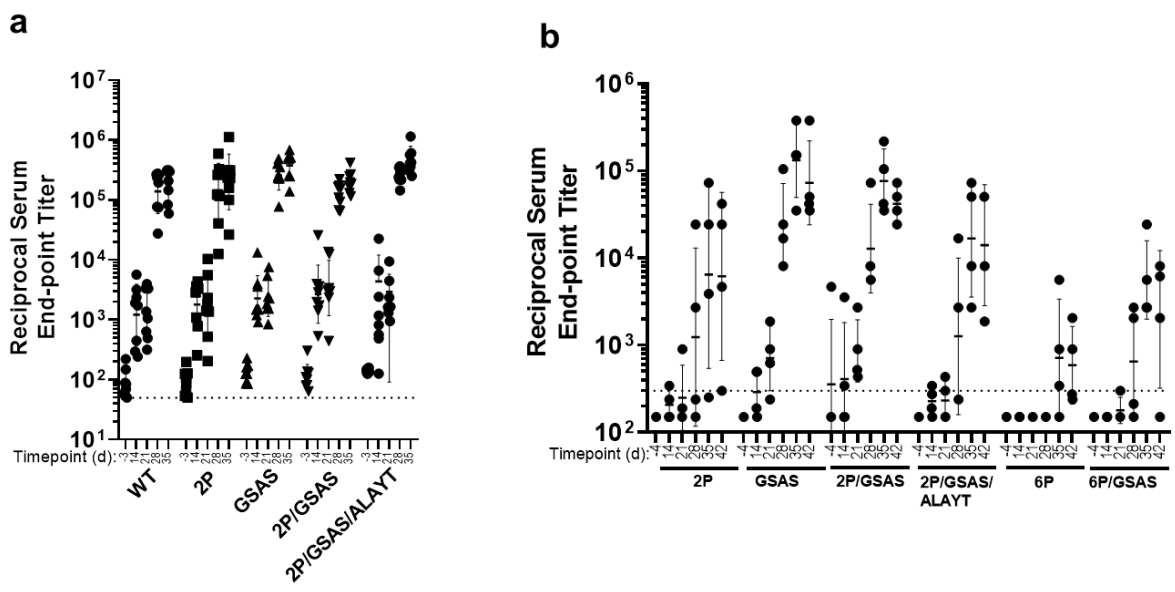


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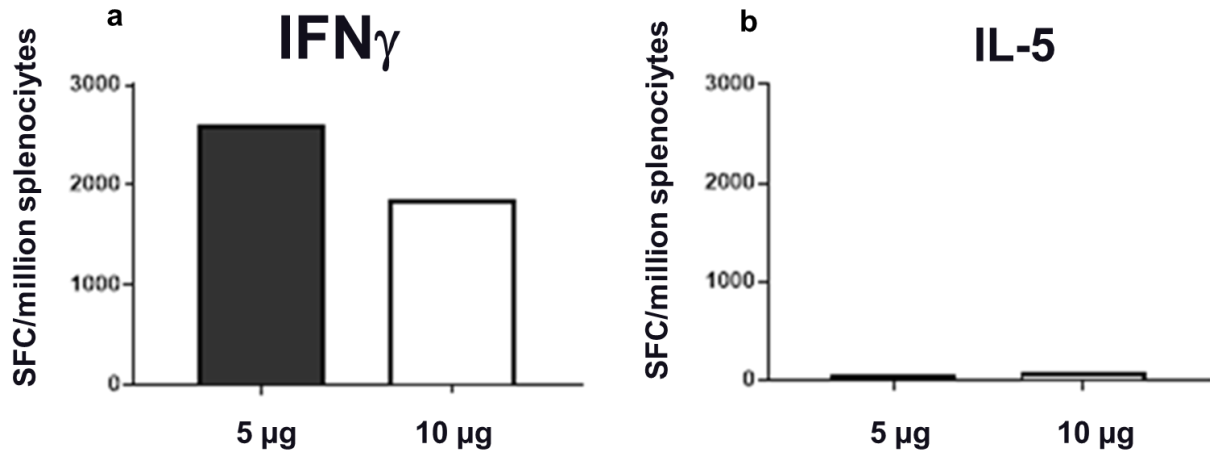
68 **Supplementary Fig.9 Comparison of antibody binding responses against S antigen**  
69 **formulations in mice and NHPs.** Serum binding antibody titers in mice (a) and NHPs (b)  
70 immunized with mRNA vaccines WT, 2P, GSAS, 2P/GSAS, 2P/GSAS/ALAYT, 6P, and  
71 6P/GSAS. Briefly, BALB/c female mice (n=8) or Cynomolgus monkeys (n=4) were immunized  
72 twice three weeks apart with 0.4 µg or 5 µg of mRNA vaccines formulations respectively. 2P,  
73 GSAS and GSAS/2P (MRT5500) were tested in both animal models, while WT was tested only  
74 in mice.  
75 Sera samples from pre-immunized animals (Pre-; D-4) as well as samples collected on D14, 21,  
76 28, 35, 42 (NHP only) were tested for reactivity to recombinant S protein in ELISA or tested in a  
77 pseudovirus neutralization assay (Fig. 2). Each dot represents an individual serum sample, and  
78 the line represents the geometric mean for the group. the dotted line below for each panel  
79 represents the lower limit of assay readout..



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81

82 **Supplementary Fig.10: MRT5500 induces TH1-biased T-cell responses in mice.** (a) IFN $\gamma$   
83 and (b) IL-5 ELISPOT D35 data for the 5 and 10  $\mu$ g doses. Following re-stimulation with S-  
84 protein peptides, pooled splenocytes MRT5500-immunized mouse groups secreted  
85 predominantly IFN $\gamma$  (TH1) whereas IL-5 (TH2) secretion was marginal.

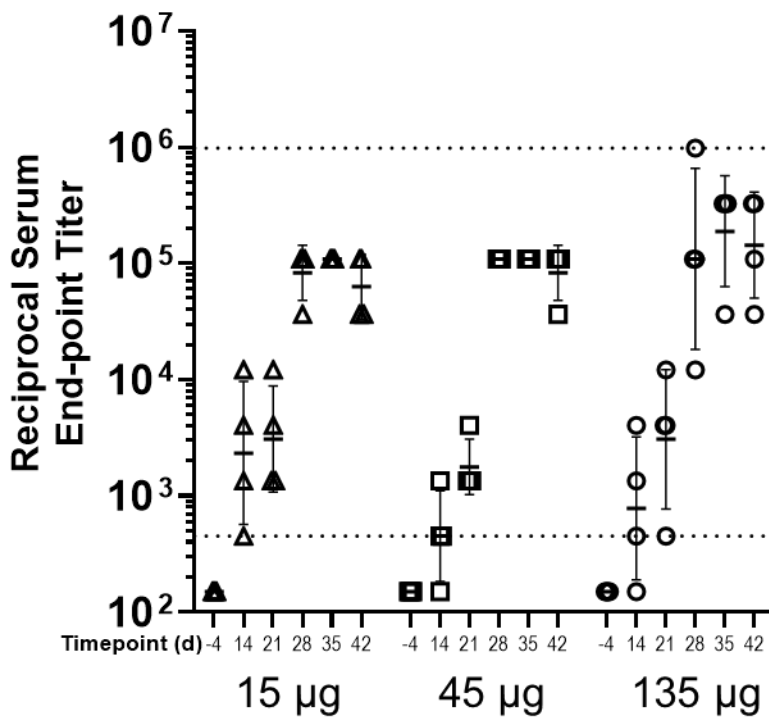


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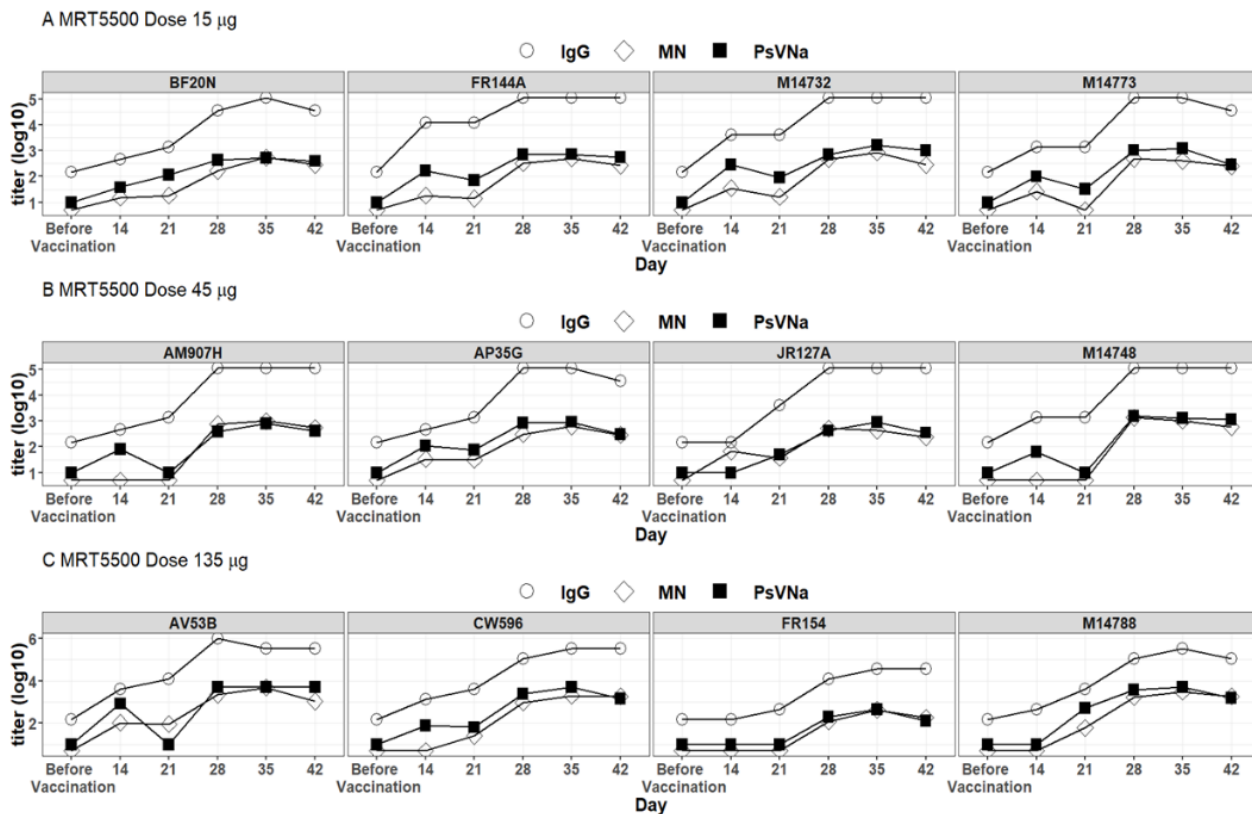
88 **Supplementary Fig.11: MRT5500 elicited strong anti-spike antibody binding response in**  
 89 **NHPs.** Briefly, Nunc MaxiSorb plates were coated with SARS-CoV S-GCN4 protein (custom  
 90 made at GeneArt) at 0.5  $\mu\text{g}/\text{mL}$  in PBS overnight at 4°C. Plates were washed 3 times with PBS-  
 91 Tween 0.1% before blocking with 1% BSA in PBS-Tween 0.1% for 1 h at ambient temperature.  
 92 Samples were plated with 1:450 initial dilution followed by 3-fold, 7-point serial dilution in  
 93 blocking buffer. Plates were washed 3 times after 1 h incubation at room temperature before  
 94 adding 50  $\mu\text{L}$  of 1:5000 Rabbit anti-human IgG (Jackson Immuno Research) to each well. Plates  
 95 were incubated at room temperature for 1hr and washed 3x. Plates were developed using Pierce  
 96 1-Step Ultra TMB-ELISA Substrate Solution for 0.1 h and stopped by TMB stop solution. Plates  
 97 were read at 450 nm in SpectraMax plate reader. Antibody titers were reported as the highest  
 98 dilution that is  $\geq 0.2$  Optical Density (OD) cutoff.

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100

101 **Supplementary Fig. 12: Strong correlations between individual NHP ELISA, PsV and MN**  
 102 **time-point titers (see also Supplementary Table 2).** Top panel A: 4 subjects in 15  $\mu$ g dose;  
 103 Middle panel B: 4 subjects in 45  $\mu$ g dose; Bottom panel C: 4 subjects in 135  $\mu$ g dose.

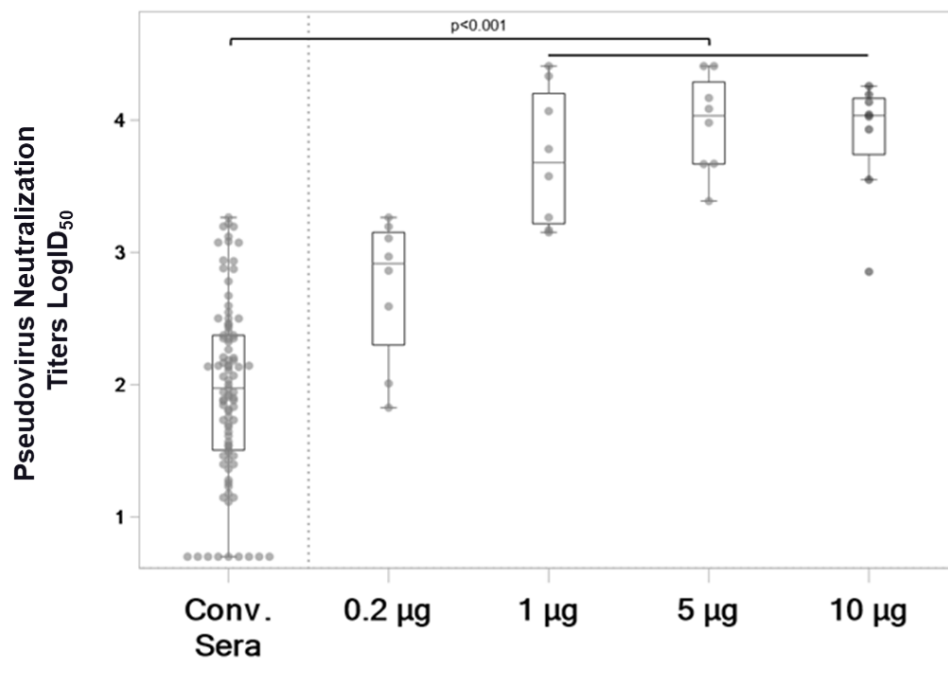


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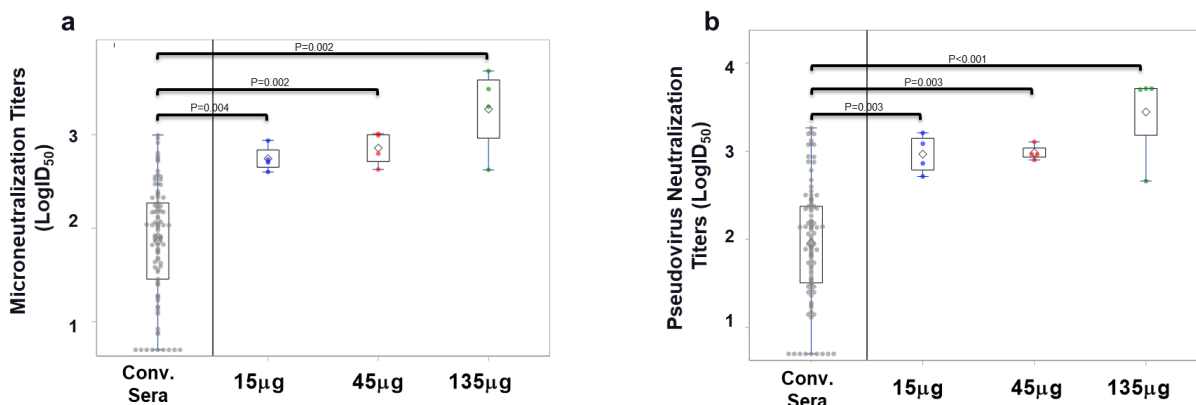
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107 **Supplementary Fig. 13: PsV titers in mice for the 1 µg, 5 µg and 10 µg dose levels of**  
 108 **MRT5500 were significantly different from the Human Convalescent sera PsV titers**



109  
 110 **Supplementary Fig. 14: Comparison of D35 MRT5500 titers to human convalescent sera**  
 111 **PsV titers in NHPs for the 15 µg, 45 µg and 135 µg dose levels of MRT5500 were**  
 112 **significantly different from the Human Convalescent sera PsV titers.**

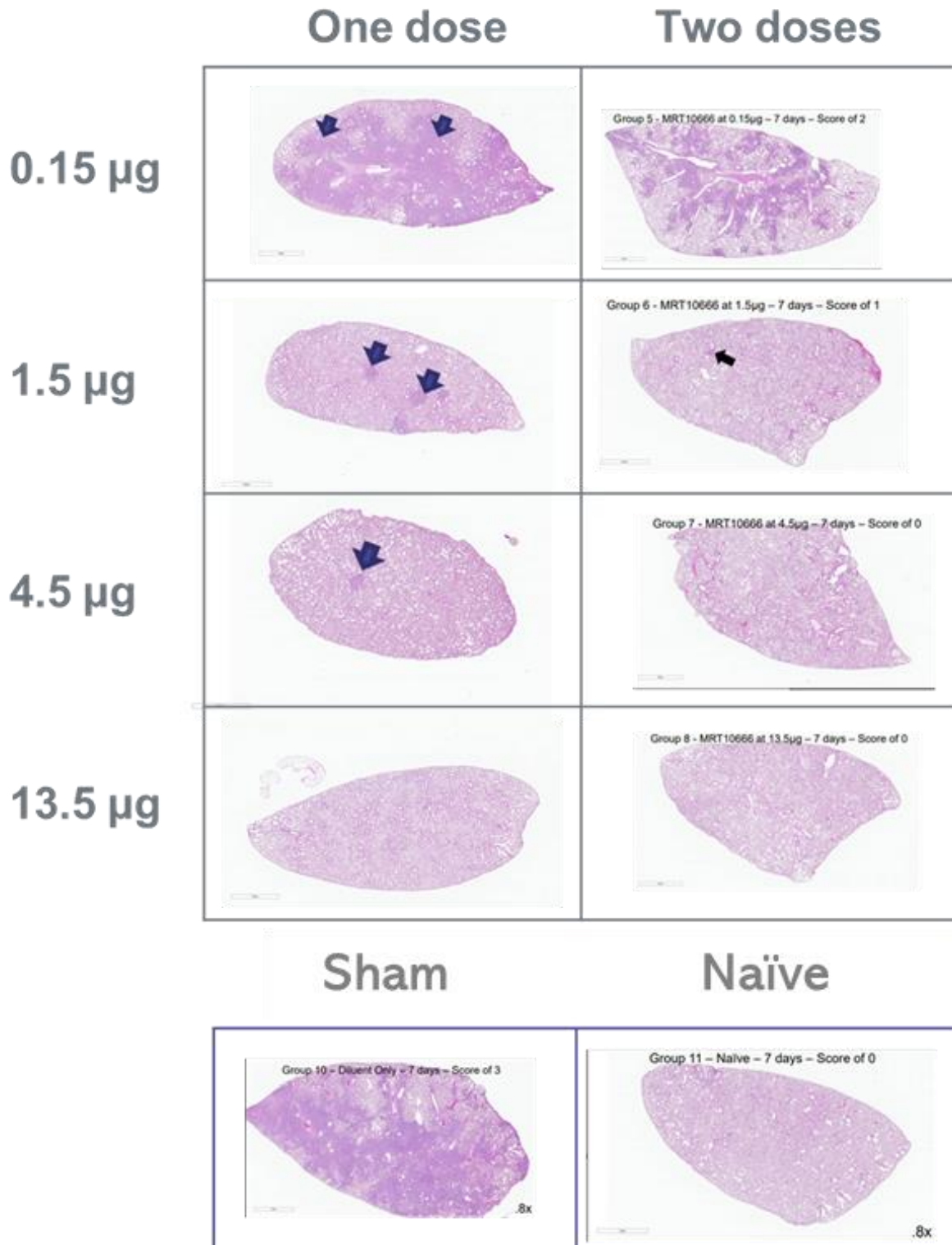


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115 **Supplementary Fig. 15: Protective efficacy of MRT5500 in hamster disease model.** Low  
116 magnification representative photomicrographs (H&E) of lungs from hamsters receiving one or  
117 two doses of MRT5500 at increasing doses. In Sham (placebo) animals, more than 50% of the  
118 lung parenchyma is disrupted by marked inflammatory cell infiltrate, type II pneumocyte  
119 hyperplasia, multifocal hemorrhage, syncytial cells and cellular debris in hamsters treated once  
120 with the lowest dose represented by multifocal dark purple regions (arrows). Note, substantially  
121 reduced affected areas in lungs of hamsters receiving two doses and increasing doses. More than  
122 50% of the lung parenchyma is disrupted in placebo treated mice as compared to naïve hamsters.  
123 Briefly, animals were immunized on D0, 21 with 0.15 µg, 1.5 µg, 4.5 µg, or 13.5 µg of  
124 MRT5500 and challenged on D42 with  $10^6$  pfu of SARS-CoV-2 intranasally (IN). Weight of

125 animals was monitored on daily basis.



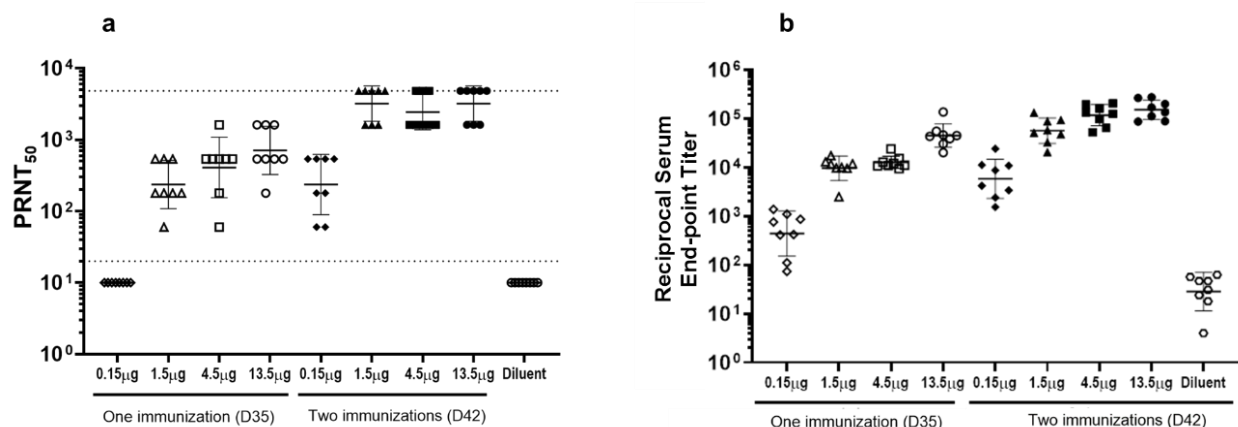
126

127 **Supplementary Fig. 15 (see legend page 14)**

128 **Supplementary Fig.16: Pre-challenge MRT5500 neutralizing and ELISA titers in hamsters.**

129 Plaque reduction neutralization (PRNT<sub>50</sub>; **a**) and ELISA (**b**) titers in sera of MRT5500  
130 vaccinated animals collected on D35 (one dose regimen) or D42 (two dose regimen). Briefly,  
131 hamsters were immunized either on D0 (one dose immunization) or D0, 21 (two doses  
132 immunizations) with 0.15  $\mu\text{g}$ , 1.5  $\mu\text{g}$ , 4.5  $\mu\text{g}$ , or 13.5  $\mu\text{g}$  of MRT5500 and challenged on D49  
133 with  $10^4$  pfu of SARS-CoV-2 intranasally (IN).

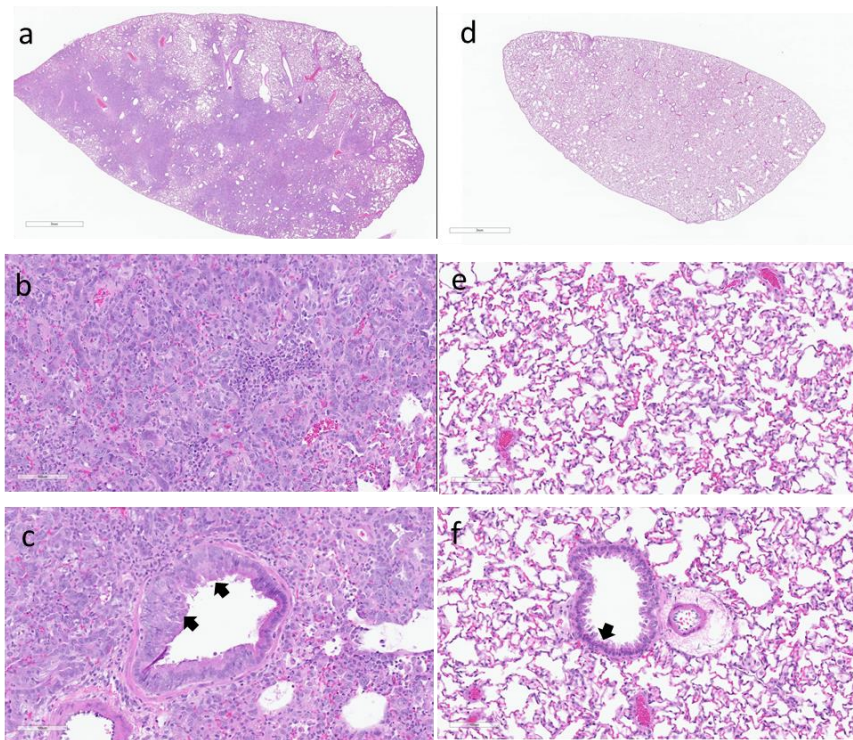
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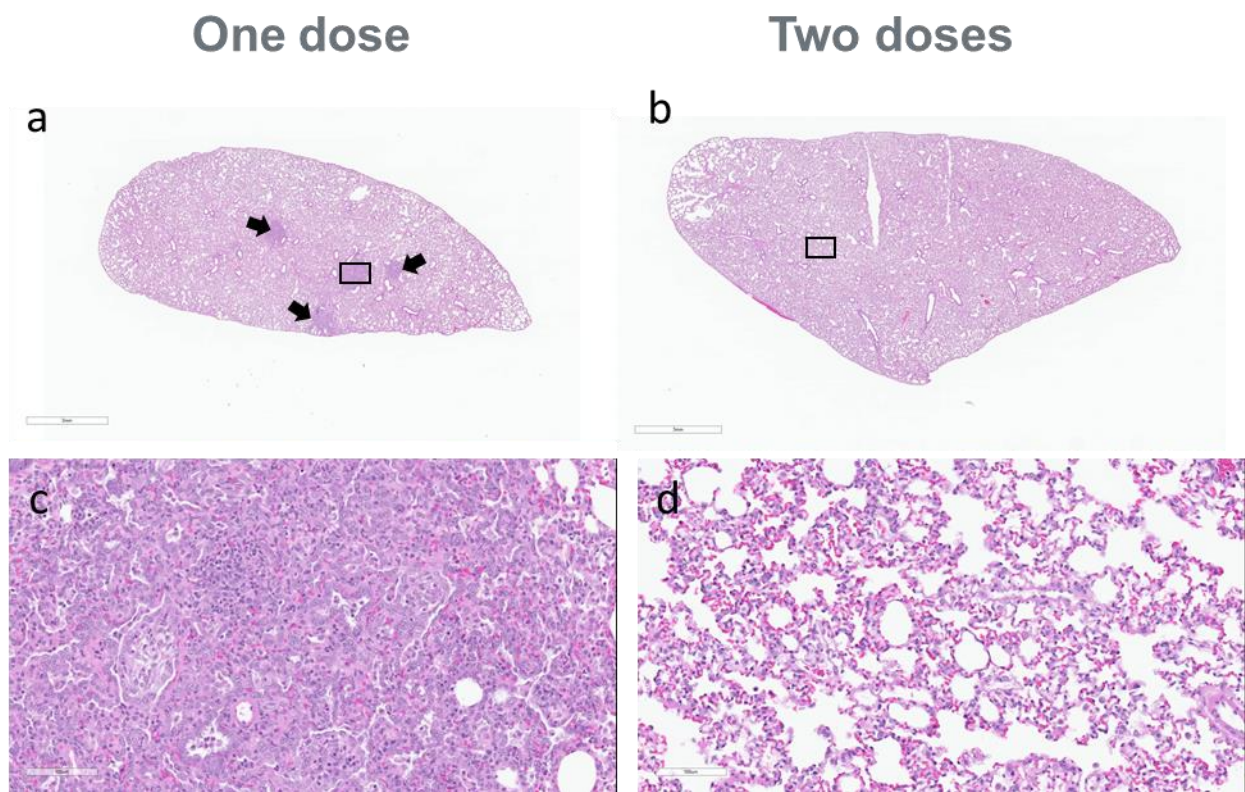
136

137 **Supplementary Fig. 17: Example of D7 post challenge histopathology findings in lungs of**  
138 **hamsters.** Representative photomicrographs (H&E) of lungs from a SARS-CoV-2 inoculated  
139 and placebo-treated hamster (a-c) and a naïve hamster (d-f). Normal lung parenchyma is  
140 disrupted by marked inflammatory cell infiltrate, type II pneumocyte hyperplasia, multifocal  
141 hemorrhage, syncytial cells, and cellular debris in infected hamster lung (b) as compared to  
142 normal parenchyma in naïve hamster lung (e). Bronchiolar epithelium (arrows) is markedly  
143 hyperplastic in infected lung sample (c) and histologically unremarkable in the naïve lung (f).



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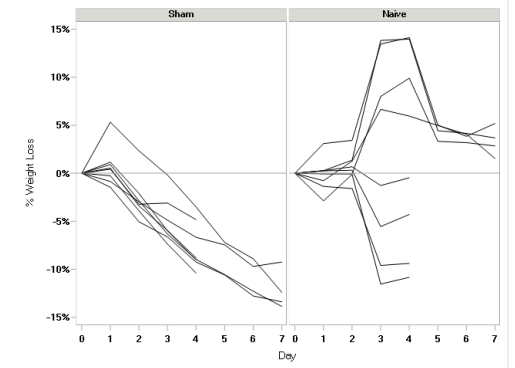
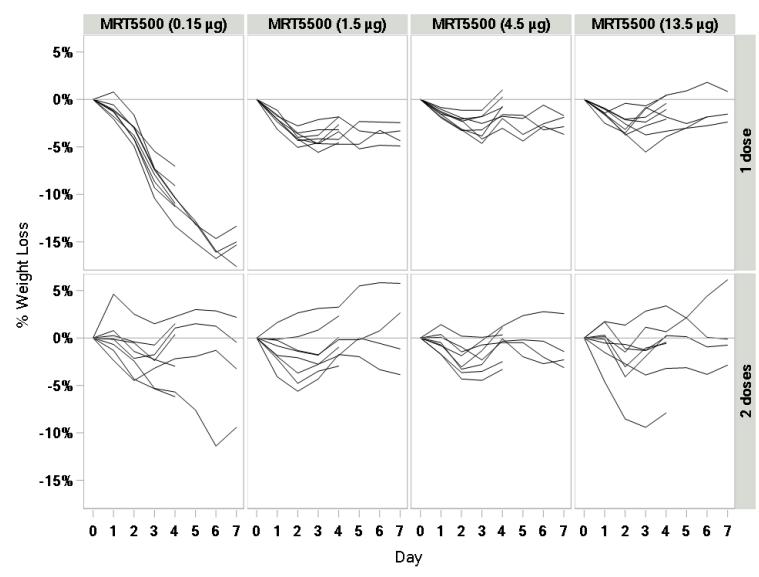
147 **Supplementary Fig.18: Example of D7 post challenge histopathology in lungs of hamsters**  
148 **receiving one dose or two doses at 1.5 µg MRT5500.** Representative lung photomicrographs  
149 (H&E). (a) Multiple foci of inflammatory cell infiltrate, type II pneumocyte hyperplasia,  
150 syncytial cells and cellular debris (arrows) are present in a hamster treated with a single 1.5 µg  
151 dose. (c) Higher magnification of boxed area in image a. Lung sections from a hamster treated  
152 with two 1.5 µg doses (b and d) appear histologically unremarkable.



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156 **Supplementary Fig.19: Hamster challenge study. Individual weight loss by groups**



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160 **Supplementary Table 1: Pairwise dose comparison in PsV neutralization titers on D35 in**  
 161 **mice.** There were no statistically significant differences in PsV titers among the 1 µg, 5 µg and  
 162 10 µg dose levels, while at the lowest dose level (0.2 µg) PsV titers were significantly different  
 163 from those obtained with the higher dose levels.

Pairwise dose comparisons PsVNa titers on D35 in mice	Fold Difference (95% CI)	p-value
0.2 µg vs. 1 µg	/9.8 (4.4, 21.7)*	p<0.0001
0.2 µg vs. 5 µg	/17.6 (7.9, 38.9)	p<0.0001
0.2 µg vs. 10 µg	/14.0 (6.3, 31.0)	p<0.0001
1 µg vs. 5 µg	/1.8 (0.8, 4.0)	p=0.1490
1 µg vs. 10 µg	/1.4 (0.6, 3.2)	p=0.3761
5 µg vs. 10 µg	x1.3 (0.6, 2.8)**	p=0.5734

Comparison: Group1 versus Group2

\* /X.X: Group1 is X.X-fold lower than group2

\*\* xX.X: Group1 is X.X-fold higher than group2

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165

166 **Supplementary Table 2: Spearman Correlation Coefficients (SCC) between ELISA (IgG),**  
 167 **Pseudoviral (PsV) and Microneutralization (MN) titers.** SCC were conducted per individual  
 168 animals (Suppl. Fig.4) and Means (95% CI) were calculated per dose (N=4) or all NHPs (N=12)

		Spearman Correlation Coefficient*		
Dose	N	ELISA and MN	ELISA and PsV	MN and PsV
15 µg	4	0.94 (0.90, 0.99)	0.95 (0.90, 1.00)	0.95 (0.90, 1.00)
45 µg	4	0.88 (0.82, 0.94)	0.90 (0.82, 0.97)	0.92 (0.81, 1.00)
135 µg	4	0.93 (0.83, 1.00)	0.88 (0.77, 0.98)	0.91 (0.84, 0.99)
Total	12	0.92 (0.89, 0.95)	0.91 (0.87, 0.94)	0.93 (0.90, 0.96)

\* Mean (95% CI)

169

170 **Supplementary Table 3: Hamster challenge data (1 dose regimen). Descriptive weight data**  
 171 **analysis during 7 DPI**

		1 Group			
		MRT5500 (0.15 µg)	MRT5500 (1.5 µg)	MRT5500 (4.5 µg)	MRT5500 (13.5 µg)
Weight D0 (g)	N	8	8	8	8
	NMiss	0	0	0	0
	Mean (SD)	142.8 (±18.6)	150.3 (±15.3)	149.7 (±10.5)	148.9 (±10.7)
	Min;Max	115.3;175.0	132.6;177.7	134.5;165.9	129.6;160.5
	Median [Q1;Q3]	141.9 [131.2;153.0]	146.6 [138.3;161.2]	150.7 [140.9;157.1]	151.3 [141.9;157.2]
Weight D1 (g)	N	8	8	8	8
	NMiss	0	0	0	0
	Mean (SD)	141.2 (±18.1)	147.2 (±14.7)	147.5 (±10.0)	146.7 (±10.5)
	Min;Max	116.2;172.9	129.8;173.5	133.1;163.3	127.6;157.9
	Median [Q1;Q3]	140.2 [129.0;151.2]	143.1 [136.3;157.8]	148.2 [139.4;154.4]	148.4 [140.2;155.5]
Weight D2 (g)	N	8	8	8	8
	NMiss	0	0	0	0
	Mean (SD)	137.9 (±18.2)	144.3 (±14.3)	146.1 (±9.5)	144.9 (±10.4)
	Min;Max	113.4;169.9	127.8;170.3	131.9;160.7	125.5;155.4
	Median [Q1;Q3]	137.1 [125.2;147.6]	140.3 [133.6;154.3]	146.5 [138.5;152.9]	148.4 [137.8;153.0]
Weight D3 (g)	N	8	8	8	8
	NMiss	0	0	0	0
	Mean (SD)	131.5 (±18.3)	144.1 (±13.6)	145.3 (±9.6)	145.3 (±9.3)
	Min;Max	106.9;162.2	126.4;167.8	131.1;159.5	128.5;155.0
	Median [Q1;Q3]	131.0 [118.3;142.3]	140.9 [134.2;154.1]	147.0 [137.1;152.0]	147.9 [138.6;152.9]
Weight D4 (g)	N	8	8	8	8
	NMiss	0	0	0	0
	Mean (SD)	128.1 (±18.5)	145.2 (±13.4)	148.1 (±10.7)	146.6 (±10.3)
	Min;Max	103.4;159.1	128.1;169.6	132.1;164.7	127.2;157.1
	Median [Q1;Q3]	126.8 [115.1;139.2]	142.3 [135.7;154.0]	149.7 [139.4;154.8]	149.0 [140.1;155.3]
Weight D5 (g)	N	4	4	4	4
	NMiss	4	4	4	4
	Mean (SD)	113.8 (±12.5)	142.1 (±15.6)	142.0 (±10.3)	147.4 (±14.3)
	Min;Max	100.1;127.5	125.7;155.5	131.8;153.8	126.3;157.0
	Median [Q1;Q3]	113.9 [103.5;124.2]	143.6 [128.7;155.5]	141.2 [133.5;150.5]	153.2 [138.5;156.4]
Weight D6 (g)	N	4	4	4	4
	NMiss	4	4	4	4
	Mean (SD)	110.7 (±11.1)	142.7 (±16.2)	142.9 (±9.7)	148.6 (±14.6)
	Min;Max	98.4;123.0	126.2;157.8	133.7;155.6	127.2;158.4
	Median [Q1;Q3]	110.7 [101.6;119.8]	143.4 [128.8;156.6]	141.2 [135.6;150.3]	154.3 [139.1;158.0]
Weight D7 (g)	N	4	4	4	3
	NMiss	4	4	4	5
	Mean (SD)	111.3 (±10.6)	142.3 (±15.6)	142.6 (±10.9)	145.4 (±15.6)
	Min;Max	99.9;124.6	126.1;156.0	132.2;156.7	127.6;156.9
	Median [Q1;Q3]	110.4 [103.3;119.4]	143.5 [128.9;155.7]	140.8 [134.2;151.1]	151.6 [127.6;156.9]

173 **Supplementary Table 4. Hamster challenge data (2 dose regimen). Descriptive weight data**  
 174 **analysis during 7 DPI**

		2 Group			
		MRT5500 (0.15 µg)	MRT5500 (1.5 µg)	MRT5500 (4.5 µg)	MRT5500 (13.5 µg)
Weight D0 (g)	N	8	8	8	8
	NMiss	0	0	0	0
	Mean (SD)	146.2 (±17.0)	152.9 (±13.0)	149.0 (±14.3)	158.6 (±9.3)
	Min;Max	120.7;169.3	131.8;173.3	121.5;163.7	143.4;170.9
	Median [Q1;Q3]	144.2 [134.9;160.8]	152.1 [145.3;161.8]	151.4 [141.6;160.2]	158.8 [153.0;165.4]
Weight D1 (g)	N	8	8	8	8
	NMiss	0	0	0	0
	Mean (SD)	146.3 (±16.1)	151.1 (±13.5)	148.3 (±14.7)	158.0 (±10.8)
	Min;Max	121.0;165.9	126.4;170.1	120.9;163.9	141.2;171.4
	Median [Q1;Q3]	147.7 [134.4;159.7]	152.8 [143.4;160.0]	150.1 [139.8;160.9]	160.1 [148.9;166.9]
Weight D2 (g)	N	8	8	8	8
	NMiss	0	0	0	0
	Mean (SD)	143.6 (±14.9)	149.9 (±13.9)	145.6 (±14.8)	154.5 (±10.3)
	Min;Max	120.2;161.7	124.4;169.7	117.5;160.9	139.5;165.7
	Median [Q1;Q3]	144.6 [133.0;155.9]	153.8 [141.1;157.7]	147.9 [137.3;158.1]	154.7 [145.7;165.2]
Weight D3 (g)	N	8	8	8	8
	NMiss	0	0	0	0
	Mean (SD)	142.6 (±15.9)	150.4 (±13.3)	146.2 (±15.2)	155.7 (±12.5)
	Min;Max	119.8;163.9	126.1;168.5	118.1;162.1	137.8;169.9
	Median [Q1;Q3]	143.7 [129.8;154.9]	154.2 [141.8;158.6]	147.3 [138.0;159.4]	156.2 [144.6;168.2]
Weight D4 (g)	N	8	8	8	8
	NMiss	0	0	0	0
	Mean (SD)	143.9 (±15.8)	152.5 (±13.1)	148.2 (±14.8)	157.1 (±11.9)
	Min;Max	122.5;165.6	129.5;170.1	121.4;165.5	138.8;169.9
	Median [Q1;Q3]	144.6 [130.6;156.4]	155.5 [143.5;161.2]	149.2 [140.2;160.1]	158.7 [146.8;168.7]
Weight D5 (g)	N	4	4	4	4
	NMiss	4	4	4	4
	Mean (SD)	148.8 (±19.0)	154.2 (±18.4)	144.0 (±18.1)	158.9 (±14.3)
	Min;Max	121.7;166.0	129.2;173.3	119.1;162.5	138.9;171.6
	Median [Q1;Q3]	153.7 [136.7;160.9]	157.2 [142.6;165.9]	147.1 [132.8;155.2]	162.5 [148.9;168.9]
Weight D6 (g)	N	4	4	4	4
	NMiss	4	4	4	4
	Mean (SD)	147.7 (±21.7)	154.0 (±19.0)	143.2 (±17.7)	158.3 (±15.6)
	Min;Max	116.7;167.1	127.4;172.3	118.2;160.0	137.9;175.4
	Median [Q1;Q3]	153.4 [134.1;161.2]	158.2 [141.9;166.1]	147.3 [132.2;154.2]	160.0 [147.5;169.1]
Weight D7 (g)	N	4	4	4	4
	NMiss	4	4	4	4
	Mean (SD)	146.6 (±19.1)	154.3 (±19.4)	142.4 (±16.8)	159.4 (±16.1)
	Min;Max	119.3;163.8	126.7;171.3	118.7;158.2	139.3;178.3
	Median [Q1;Q3]	151.6 [134.9;158.3]	159.6 [141.5;167.1]	146.4 [131.7;153.2]	160.0 [148.4;170.4]

176 **Supplementary Table 5. Hamster challenge data (Control groups). Descriptive weight data**  
 177 **analysis during 7 DPI observation**

	Sham	Naive
Weight D0 (g)		
N	8	8
NMiss	0	0
Mean (SD)	149.2 ( $\pm$ 14.3)	146.6 ( $\pm$ 14.8)
Min;Max	129.3;167.6	119.6;170.7
Median [Q1;Q3]	152.7 [135.5;160.1]	147.1 [140.6;153.4]
Weight D1 (g)		
N	8	8
NMiss	0	0
Mean (SD)	150.2 ( $\pm$ 14.2)	146.2 ( $\pm$ 13.0)
Min;Max	130.0;168.3	123.3;165.8
Median [Q1;Q3]	152.8 [136.9;162.0]	147.2 [139.1;153.8]
Weight D2 (g)		
N	8	8
NMiss	0	0
Mean (SD)	145.2 ( $\pm$ 13.9)	147.4 ( $\pm$ 14.0)
Min;Max	124.9;162.2	123.7;170.5
Median [Q1;Q3]	148.4 [132.2;156.8]	148.0 [140.4;154.2]
Weight D3 (g)		
N	8	8
NMiss	0	0
Mean (SD)	141.7 ( $\pm$ 14.9)	148.3 ( $\pm$ 13.0)
Min;Max	121.6;162.4	125.2;167.0
Median [Q1;Q3]	144.2 [127.2;153.5]	150.4 [141.1;155.7]
Weight D4 (g)		
N	8	8
NMiss	0	0
Mean (SD)	137.8 ( $\pm$ 14.9)	149.2 ( $\pm$ 13.3)
Min;Max	117.9;159.5	125.5;167.2
Median [Q1;Q3]	140.5 [123.3;148.8]	151.5 [142.1;157.1]
Weight D5 (g)		
N	4	4
NMiss	4	4
Mean (SD)	137.1 ( $\pm$ 12.6)	145.2 ( $\pm$ 13.2)
Min;Max	118.6;146.9	125.6;153.2
Median [Q1;Q3]	141.5 [129.4;144.9]	151.1 [137.7;152.8]
Weight D6 (g)		
N	4	4
NMiss	4	4
Mean (SD)	134.1 ( $\pm$ 12.2)	144.4 ( $\pm$ 13.6)
Min;Max	116.3;143.3	124.2;152.8
Median [Q1;Q3]	138.5 [126.5;141.8]	150.3 [136.4;152.5]
Weight D7 (g)		
N	4	4
NMiss	4	4
Mean (SD)	132.2 ( $\pm$ 12.4)	143.6 ( $\pm$ 12.3)
Min;Max	114.2;142.3	125.8;152.1
Median [Q1;Q3]	136.1 [124.5;139.9]	148.3 [135.4;151.9]

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180 **Supplementary Table 6. Hamster challenge data (1 dose regimen). % weight loss during 7**  
 181 **DPI observation**

		1 Group			
		MRT5500 (0.15 µg)	MRT5500 (1.5 µg)	MRT5500 (4.5 µg)	MRT5500 (13.5 µg)
% Weight Loss (D1)					
N	8	8	8	8	8
NMiss	0	0	0	0	0
Mean (SD)	-1.1 (±0.9)	-2.0 (±0.6)	-1.4 (±0.4)	-1.4 (±0.5)	
Min;Max	-2.1;0.8	-3.1;-1.1	-1.9;-0.9	-2.5;-0.9	
Median [Q1;Q3]	-1.2 [-1.6;-0.8]	-2.1 [-2.2;-1.7]	-1.5 [-1.7;-1.2]	-1.4 [-1.6;-1.0]	
% Weight Loss (D2)					
N	8	8	8	8	8
NMiss	0	0	0	0	0
Mean (SD)	-3.4 (±1.0)	-4.0 (±0.7)	-2.4 (±0.7)	-2.7 (±1.1)	
Min;Max	-4.9;-1.6	-5.1;-2.8	-3.3;-1.1	-3.7;-0.4	
Median [Q1;Q3]	-3.4 [-4.2;-2.9]	-4.1 [-4.3;-3.6]	-2.2 [-3.2;-2.0]	-2.9 [-3.6;-2.1]	
% Weight Loss (D3)					
N	8	8	8	8	8
NMiss	0	0	0	0	0
Mean (SD)	-8.0 (±1.5)	-4.1 (±1.1)	-2.9 (±1.3)	-2.4 (±1.7)	
Min;Max	-10.4;-5.5	-5.6;-2.1	-4.6;-1.1	-5.5;-0.7	
Median [Q1;Q3]	-7.7 [-9.0;-7.3]	-4.4 [-4.7;-3.5]	-2.9 [-4.0;-1.8]	-2.1 [-3.3;-0.9]	
% Weight Loss (D4)					
N	8	8	8	8	8
NMiss	0	0	0	0	0
Mean (SD)	-10.4 (±1.8)	-3.3 (±1.1)	-1.1 (±1.3)	-1.5 (±1.6)	
Min;Max	-13.3;-7.0	-4.7;-1.8	-3.0;1.0	-3.9;0.4	
Median [Q1;Q3]	-10.7 [-11.2;-9.7]	-3.3 [-4.4;-2.3]	-1.2 [-1.9;-0.2]	-1.5 [-2.7;-0.0]	
% Weight Loss (D5)					
N	4	4	4	4	4
NMiss	4	4	4	4	4
Mean (SD)	-13.5 (±1.0)	-3.9 (±1.3)	-2.9 (±1.3)	-1.9 (±1.9)	
Min;Max	-15.1;-12.8	-5.2;-2.3	-4.4;-1.7	-3.0;0.9	
Median [Q1;Q3]	-13.1 [-14.1;-12.9]	-4.0 [-5.0;-2.8]	-2.9 [-4.0;-1.8]	-2.8 [-3.0;-0.8]	
% Weight Loss (D6)					
N	4	4	4	4	4
NMiss	4	4	4	4	4
Mean (SD)	-15.9 (±0.9)	-3.5 (±1.0)	-2.3 (±1.2)	-1.2 (±2.0)	
Min;Max	-16.8;-14.7	-4.8;-2.4	-3.2;-0.6	-2.8;1.8	
Median [Q1;Q3]	-16.0 [-16.4;-15.3]	-3.4 [-4.2;-2.8]	-2.7 [-3.0;-1.6]	-1.8 [-2.3;-0.0]	
% Weight Loss (D7)					
N	4	4	4	3	3
NMiss	4	4	4	5	5
Mean (SD)	-15.3 (±1.7)	-3.8 (±1.1)	-2.5 (±0.9)	-1.0 (±1.7)	
Min;Max	-17.6;-13.4	-4.9;-2.4	-3.7;-1.7	-2.4;0.8	
Median [Q1;Q3]	-15.2 [-16.5;-14.2]	-3.8 [-4.6;-2.9]	-2.4 [-3.3;-1.8]	-1.5 [-2.4;0.8]	

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185 **Supplementary Table 7. Hamster challenge data (2 dose regimen). % weight loss during 7**

186 **DPI observation**

		2 Group			
		MRT5500 (0.15 µg)	MRT5500 (1.5 µg)	MRT5500 (4.5 µg)	MRT5500 (13.5 µg)
<b>% Weight Loss (D1)</b>					
N		8	8	8	8
NMiss		0	0	0	0
Mean (SD)		0.1 (±2.1)	-1.2 (±1.7)	-0.5 (±1.1)	-0.4 (±2.0)
Min;Max		-2.4;4.6	-4.1;1.6	-1.7;1.4	-4.6;1.7
Median [Q1;Q3]		-0.1 [-1.0;0.5]	-1.3 [-2.1;-0.2]	-0.6 [-1.3;0.2]	-0.0 [-1.0;1.0]
<b>% Weight Loss (D2)</b>					
N		8	8	8	8
NMiss		0	0	0	0
Mean (SD)		-1.7 (±2.3)	-2.0 (±2.7)	-2.3 (±1.5)	-2.5 (±2.9)
Min;Max		-4.5;2.5	-5.6;2.6	-4.3;0.2	-8.5;1.4
Median [Q1;Q3]		-1.8 [-3.4;-0.5]	-1.7 [-4.2;-0.6]	-2.5 [-3.5;-1.2]	-2.1 [-3.6;-0.9]
<b>% Weight Loss (D3)</b>					
N		8	8	8	8
NMiss		0	0	0	0
Mean (SD)		-2.4 (±2.3)	-1.6 (±2.5)	-1.9 (±1.6)	-1.9 (±3.7)
Min;Max		-5.4;1.5	-4.3;3.1	-4.5;0.1	-9.4;2.8
Median [Q1;Q3]		-2.3 [-4.3;-1.3]	-2.3 [-3.1;-0.5]	-1.8 [-3.2;-0.5]	-1.3 [-2.9;0.0]
<b>% Weight Loss (D4)</b>					
N		8	8	8	8
NMiss		0	0	0	0
Mean (SD)		-1.5 (±3.3)	-0.3 (±2.1)	-0.5 (±1.6)	-1.0 (±3.3)
Min;Max		-6.2;2.2	-3.0;3.2	-3.3;1.2	-7.9;3.4
Median [Q1;Q3]		-0.9 [-4.3;1.3]	-0.6 [-1.8;1.2]	-0.2 [-1.5;0.7]	-0.2 [-1.9;0.5]
<b>% Weight Loss (D5)</b>					
N		4	4	4	4
NMiss		4	4	4	4
Mean (SD)		-1.3 (±4.7)	0.8 (±3.2)	-0.1 (±1.8)	0.3 (±2.5)
Min;Max		-7.6;3.0	-2.0;5.5	-2.0;2.4	-3.1;2.1
Median [Q1;Q3]		-0.2 [-4.8;2.2]	-0.1 [-1.1;2.7]	-0.3 [-1.2;1.1]	1.1 [-1.5;2.1]
<b>% Weight Loss (D6)</b>					
N		4	4	4	4
NMiss		4	4	4	4
Mean (SD)		-2.1 (±6.4)	0.7 (±3.8)	-0.6 (±2.4)	-0.1 (±3.4)
Min;Max		-11.4;2.9	-3.3;5.8	-2.7;2.8	-3.8;4.4
Median [Q1;Q3]		-0.0 [-6.3;2.0]	0.1 [-2.0;3.3]	-1.2 [-2.4;1.2]	-0.4 [-2.4;2.2]
<b>% Weight Loss (D7)</b>					
N		4	4	4	4
NMiss		4	4	4	4
Mean (SD)		-2.7 (±5.0)	0.8 (±4.2)	-1.1 (±2.5)	0.6 (±3.9)
Min;Max		-9.4;2.2	-3.9;5.8	-3.1;2.6	-2.9;6.1
Median [Q1;Q3]		-1.9 [-6.3;0.9]	0.7 [-2.5;4.2]	-1.9 [-2.7;0.6]	-0.4 [-1.8;3.0]

188 **Supplementary Table 8. Hamster challenge data (Control groups). % weight loss during 7**

189 **DPI observation**

	Sham	Naive
% Weight Loss (D1)		
N	8	8
NMiss	0	0
Mean (SD)	0.7 ( $\pm 2.1$ )	-0.1 ( $\pm 1.7$ )
Min;Max	-1.5;5.3	-2.9;3.1
Median [Q1;Q3]	0.5 [-0.6;1.0]	0.1 [-1.1;0.3]
% Weight Loss (D2)		
N	8	8
NMiss	0	0
Mean (SD)	-2.6 ( $\pm 2.2$ )	0.7 ( $\pm 1.5$ )
Min;Max	-5.1;2.3	-1.6;3.4
Median [Q1;Q3]	-3.1 [-3.7;-2.5]	0.5 [-0.1;1.3]
% Weight Loss (D3)		
N	8	8
NMiss	0	0
Mean (SD)	-5.1 ( $\pm 2.4$ )	1.7 ( $\pm 10.1$ )
Min;Max	-7.4;-0.2	-11.5;13.8
Median [Q1;Q3]	-6.0 [-6.5;-4.0]	2.7 [-7.6;10.7]
% Weight Loss (D4)		
N	8	8
NMiss	0	0
Mean (SD)	-7.7 ( $\pm 2.4$ )	2.4 ( $\pm 10.1$ )
Min;Max	-10.4;-3.5	-10.8;14.1
Median [Q1;Q3]	-8.9 [-9.1;-5.8]	2.7 [-6.8;11.9]
% Weight Loss (D5)		
N	4	4
NMiss	4	4
Mean (SD)	-9.0 ( $\pm 1.9$ )	4.4 ( $\pm 0.8$ )
Min;Max	-10.6;-7.2	3.3;5.0
Median [Q1;Q3]	-9.0 [-10.6;-7.3]	4.7 [3.9;5.0]
% Weight Loss (D6)		
N	4	4
NMiss	4	4
Mean (SD)	-10.9 ( $\pm 1.9$ )	3.8 ( $\pm 0.4$ )
Min;Max	-12.8;-8.9	3.2;4.2
Median [Q1;Q3]	-11.0 [-12.5;-9.3]	4.0 [3.5;4.1]
% Weight Loss (D7)		
N	4	4
NMiss	4	4
Mean (SD)	-12.2 ( $\pm 2.1$ )	3.3 ( $\pm 1.5$ )
Min;Max	-13.9;-9.2	1.5;5.2
Median [Q1;Q3]	-12.9 [-13.6;-10.8]	3.3 [2.2;4.4]

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193 **Supplementary Table 9. Pairwise comparisons of PsVNa titers on D35 in NHPs (5 µg dose,**  
 194 **see also Fig.2)**

Group pairs	p-value (Wilcoxon Exact Rank Test)
<b>2P/GSAS vs 2P</b>	<b>0.3143**</b>
<b>2P/GSAS vs GSAS</b>	<b>0.4857</b>
<b>2P/GSAS vs 2P/GSAS/ALAYT</b>	<b>0.4857</b>
<b>2P/GSAS vs 6P</b>	<b>0.0286</b>
<b>2P/GSAS vs 6P/GSAS</b>	<b>0.1143</b>
<b>2P/GSAS vs Convalescent Sera</b>	<b>0.0105*</b>

\* For comparison versus Convalescent Sera, Wilcoxon Rank Test was used (not the exact test)

\*\* Two animals out of four in 2P group demonstrated PsVNa titers below lower limit of detection (see Fig.2)

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198 **Supplementary Table 10. sgmRNA copies in lungs and nares of MRT5500 vaccinated**  
 199 **hamsters as compared to Sham (diluent) group (4 and 7 DPI)**

**4 DPI**

Dose number	1 dose				2 doses			
	0.15 µg	1.5 µg	4.5 µg	13.5 µg	0.15 µg	1.5 µg	4.5 µg	13.5 µg
sgRNA copies in <b>Lungs</b>	> Sham p=0.0286	> Sham p=0.4857	= Sham p=1	< Sham p=0.0286	< Sham p=0.0571	< Sham p=0.0286	< Sham p=0.0286	< Sham p=0.0286
sgRNA copies in <b>Nares</b>	> Sham p=0.0571	< Sham p=0.8857	> Sham - p=0.1143	= Sham p=1	< Sham p=0.8857	< Sham p=0.4	< Sham p=0.4857	< Sham p=0.5429

**7 DPI**

Dose number	1 dose				2 doses			
	0.15 µg	1.5 µg	4.5 µg	13.5 µg	0.15 µg	1.5 µg	4.5 µg	13.5 µg
sgRNA copies in <b>Lungs</b>	> Sham p=0.1429	< Sham p=0.1429	< Sham - p=0.1429	< Sham p=0.1429	< Sham p=0.2571	< Sham p=0.1429	< Sham p=0.1429	< Sham p=0.1429
sgRNA copies in <b>Nares</b>	< Sham p=0.8857	< Sham - p=0.1143	< Sham - p=0.0571	< Sham p=0.0286	< Sham p=0.0286	< Sham p=0.3143	< Sham p=0.6	< Sham p=0.0286

- Tables represent the results of Wilcoxon Exact Test comparisons against Sham regarding sgRNA copies in Lung and Nares.
- “=” (equal), <’ (less then) or >’ (more then) signs represent the directions of the difference as compared to Sham group
- DPI – days post infection (challenge) with SARS-CoV-2

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203 **Supplementary Table 11: List of acronyms**

204	mRNA	messenger RNA
205	LNP	Lipid Nanoparticles
206	MRT5500	mRNA/LNP vaccine formulation
207	COVID-19	Coronavirus disease 2019
208	2019-nCoV	The 2019-novel coronavirus
209	SARS	Severe Acute Respiratory Syndrome
210	MERS-CoV	Coronavirus causing Middle East Respiratory Syndrome
211	HCoV-HKU1	Human coronavirus HKU1
212	SARS-CoV-2	Severe Acute Respiratory Syndrome Coronavirus 2, virus causing
213		COVID-19
214	S-protein or S	Structural Spike glycoprotein of SARS-CoV-2
215	S-GCN4 protein	Recombinant S-protein containing a fusion of S-ectodomain with artificial
216		trimerization domain GCN4 (custom made in GeneArt)
217	E-, M-, N-proteins	Proteins E, M, N of SARS-CoV-2
218	S1, S2	S1 and S2 subunits of S-protein
219	RBD	Receptor Binding Domain of S-protein
220	ACE2	Angiotensin-Converting Enzyme 2, cellular receptor of SARS CoV-1, 2
221	WT S-protein	Wild Type full length S-protein of SARS-CoV-2
222	2P S-protein	Pre-fusion stabilized mutant of full length S-protein of SARS-CoV-2
223	GSAS S-protein	Furin cleavage-site mutant of full length S-protein of SARS-CoV-2
224	2P/GSAS	S-protein double mutant containing 2P and GSAS
225	nAb	Neutralizing Antibodies
226	mAb	Monoclonal Antibodies
227	HEK293	Immortalized Human Embryonic Kidney 293 cells
228	HEK293T	The cell line, derived from the HEK293 cell line, that expresses
229		a mutant version of the SV40 large T antigen
230	HeLa	Immortalized Human cervical cell line
231	kDa	kilo-Daltons, unit of protein mass

232	ER	Endoplasmic Reticulum
233	KDEL	Molecular marker of ER
234	ERGIC	Endoplasmic Reticulum–Golgi Intermediate Compartment
235	KLHYT	Intracellular ER retention signal of S—proteins of SARS-CoV-1, 2
236	BALB/c	Albino Laboratory-Bred/c strain of mice
237	ELISA	Enzyme-Linked Immunosorbent Assay
238	MN	Microneutralization
239	IgG	Immunoglobulin G
240	IgA	Immunoglobulin A
241	GMT	Geometric Mean Titers
242	GFP	Green Fluorescence Protein
243	RVP	GFP Reporter pseudoViral Particles
244	PsVNa	Pseudoviral neutralization assay
245	ID <sub>50</sub>	Serum dilution providing 50% inhibition of RVP or WT SARS-CoV-2
246		entry
247	NHP	Non-Human Primate
248	PsV	Pseudovirus, pseudoviral
249	VAERD	Vaccine Associated Enhanced Respiratory Disease
250	RSV	Respiratory Syncytial Virus
251	RSV F protein	Major structural protein F of RSV
252	T <sub>H</sub> 1, T <sub>H</sub> 2	T-helper cells type 1 and 2
253	IFN- $\gamma$	Interferon gamma
254	IL-13	Interleukin 13
255	IL-5	Interleukin 5
256	rhIL-2	recombinant human Interleukin 2 (IL-2)
257	PBMC	Peripheral Blood Mononuclear Cell
258	ELISPOT	Enzyme-Linked Immune Absorbent Spot assay
259	RPMI 1640	The growth medium used in cell culture

260	Cap 1	Specially altered nucleotide on the 5' end of some primary transcripts such
261		as precursor messenger RNA
262	Poly(A) tail	A stretch of multiple adenosine monophosphates at 3' end of some
263		primary transcripts such as precursor messenger RNA
264	PBS	Phosphate Buffer
265	Triton X-100	Nonionic surfactant used for permeabilization of HeLa cells
266	IACUC	Institutional Animal Care and Use Committee
267	D	Day
268	IM	Intramuscular
269	TCID <sub>50</sub>	50% Tissue Culture Infectious Dose
270	MN ID <sub>50</sub> Titer	OD of 50% neutralization point - intercept)/slope
271	CovA	Concanavalin A
272	OD	Optical Density
273	CTL	Contact Laboratory Services
274	SFC	Spot Forming Cells
275	h	h
276	µg	Microgram
277	ng	Nanogram
278	µl	Microliter
279	mL	Milliliter