

## Supplementary Information

### Passive transfer of Ad26.COV2.S-elicited IgG from humans attenuates SARS-CoV-2 disease in hamsters

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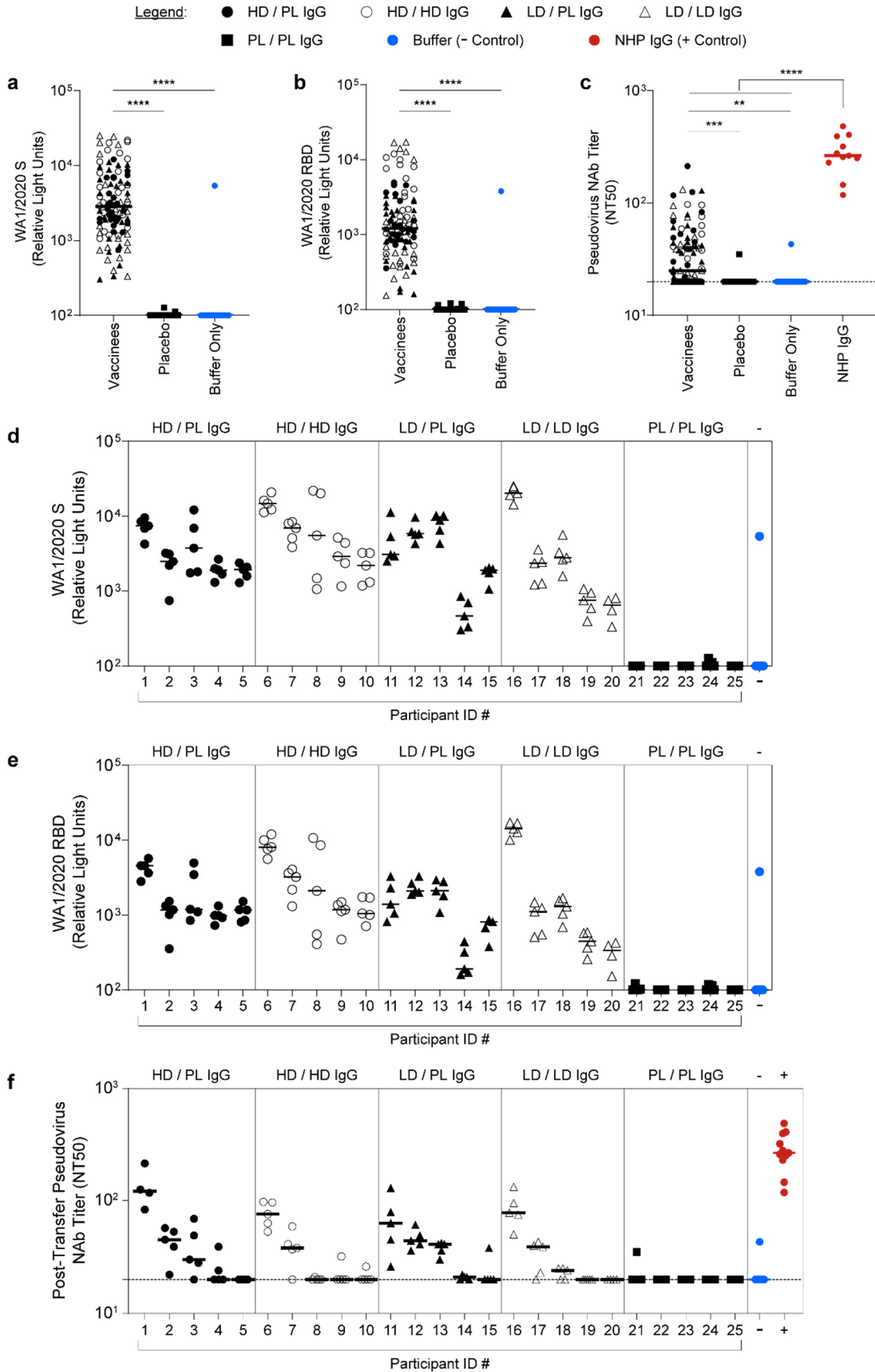
<sup>4</sup>Janssen Vaccines & Prevention, Leiden, The Netherlands.

<sup>5</sup>Janssen Research & Development, Beerse, Belgium

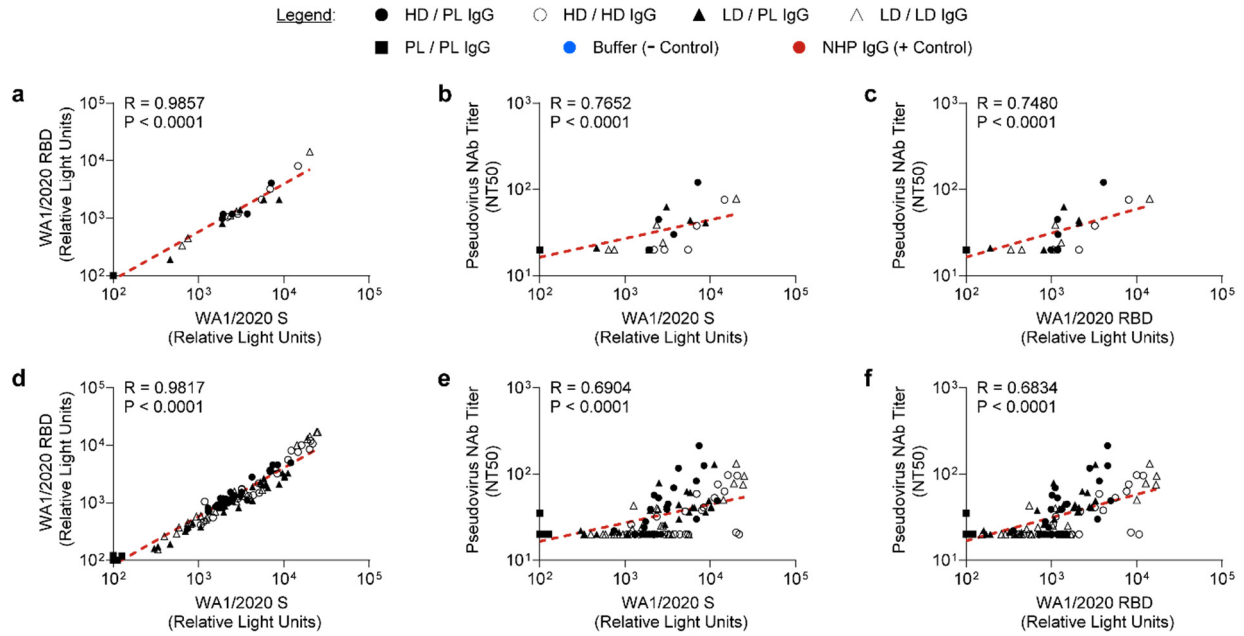
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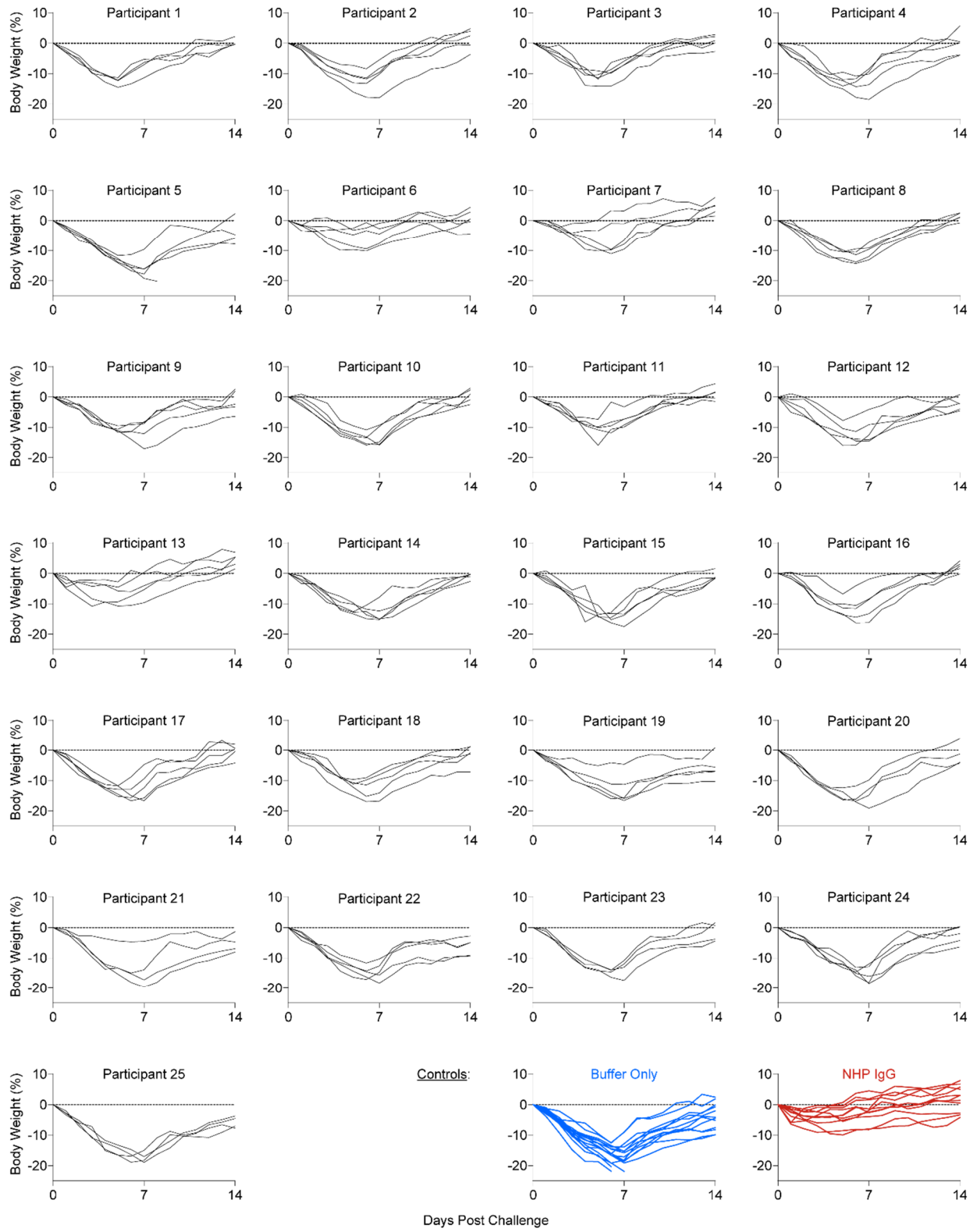
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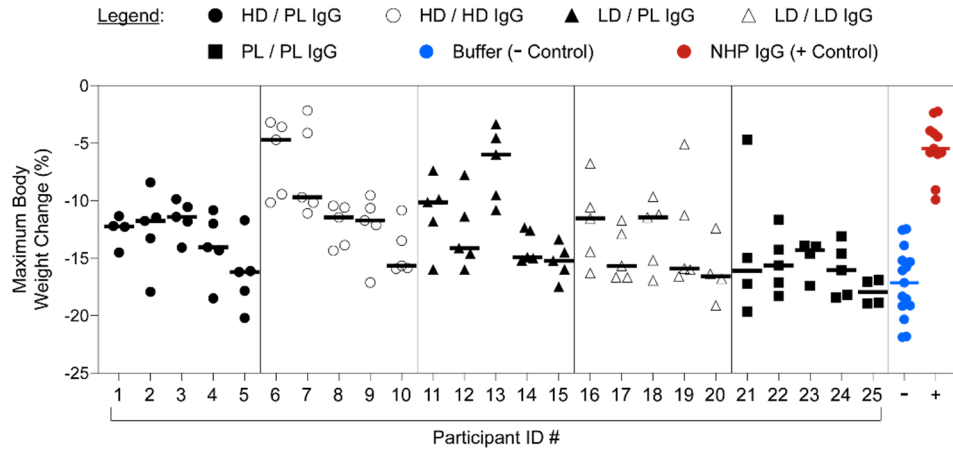
**Supplementary Figure 1.** Purified IgG from vaccine or placebo recipients was transferred to naïve hamsters via intraperitoneal injection and one-day post-transfer, serum was analyzed for **a)** WA1/2020 SARS-CoV-2 spike (S) binding, **b)** WA1/2020 SARS-CoV-2 receptor binding domain (RBD) binding, or **c)** WA1/2020 pseudovirus neutralization titers. Additional control groups included hamsters that received buffer alone and hamsters that received an equivalent dose (i.e., 25 mg) of IgG purified from convalescent non-human primates (NHPs). Data displayed in a-b were generated via electrochemiluminescence assays (ECLA) using an anti-human detection antibody, thus the NHP IgG recipient group was excluded from the analyses. In panels a-c, data points displayed correspond to the value of each individual recipient hamsters and horizontal lines indicate group medians. Statistics displayed are the results of Kruskal-Wallis tests with Dunn's multiple comparisons test. (\*\* =  $P \leq 0.01$ ; \*\*\* =  $P \leq 0.001$ ; \*\*\*\* =  $P \leq 0.0001$ ). Similarly, in panels **d-f)** the binding and neutralizing antibody titers of individual hamsters are displayed, separated into groups corresponding to the N=20 study participants who received the Ad26.COV2.S vaccine, the N=5 participants who received a placebo immunization, or the control hamsters that received either buffer alone or convalescent NHP IgG. Within each dosing regimen, participants were ordered and assigned a number 1-25 by magnitude of response for visualization purposes. As above, in panels d-e, the NHP IgG recipient group was excluded from ECLA analyses due to the human-specific detection antibody. Horizontal lines in panels d-f indicate the group medians.



**Supplementary Figure 2. Post-transfer binding and neutralizing antibody activity correlation analyses.** Purified IgG from vaccine or placebo recipients was transferred to naïve hamsters via intraperitoneal injection and one-day post-transfer, serum was analyzed for **a)** WA1/2020 SARS-CoV-2 spike (S) binding, **b)** WA1/2020 SARS-CoV-2 receptor binding domain (RBD) binding, or **c)** WA1/2020 pseudovirus neutralization titers. Correlation analyses between these three assays were completed for the a-c) median values for each group of hamsters corresponding to one study participant and d-f) each individual hamster as a single data point. Statistics in all panels indicate the results of Spearman correlation analyses.



**Supplementary Figure 3.** One day post-IgG transfer, groups of hamsters were challenged with SARS-CoV-2 via the intranasal route. Post-challenge, hamsters were monitored for fourteen days for signs of clinical disease. Data shown represent the individual weight traces of each hamster in the groups corresponding to the study participants (N=25), as well as the negative control (buffer only) and positive control (NHP IgG) groups. Study participants received the following dosing regimens, as described in Table 1: i) Participants 1-5: HD/PL, Participants 6-10: HD/HD, Participants 11-15: LD/PL, Participants 16-20: LD/LD, and Participants 21-25: PL/PL. Incomplete body weight traces represent animals that exhibited greater than 20% body weight loss, a humane endpoint euthanasia criteria.



**Supplementary Figure 4.** Data displayed correspond to the median maximum body weight change, similar to the data displayed in Fig 2c-d. In these Supplementary, maximum body weight change is shown for each individual hamster in the groups corresponding to the study participants or the control groups, as indicated

**Supplementary Table 1.** Study overview of the Ad26.COV2.S dosing regimens tested in cohort

1b.

<b>Group</b>	<b>N</b>	<b>Day 1</b>	<b>Day 57</b>	<b>Code</b>
1	5	1 x 10 <sup>11</sup> VP Ad26.COV2.S	Placebo	HD / PL
2	5	1 x 10 <sup>11</sup> VP Ad26.COV2.S	1 x 10 <sup>11</sup> VP Ad26.COV2.S	HD / HD
3	5	5 x 10 <sup>10</sup> VP Ad26.COV2.S	Placebo	LD / PL
4	5	5 x 10 <sup>10</sup> VP Ad26.COV2.S	5 x 10 <sup>10</sup> VP Ad26.COV2.S	LD / LD
5	5	Placebo	Placebo	PL / PL