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1 **Summary**

2 **Efficacy of a Broadly Neutralizing SARS-CoV-2 Ferritin Nanoparticle Vaccine in Nonhuman Primates**

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31 The emergence of novel severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) variants
32 stresses the continued need for next-generation vaccines that confer broad protection against
33 coronavirus disease 2019 (COVID-19). We developed and evaluated an adjuvanted SARS-CoV-2
34 Spike Ferritin Nanoparticle (SpFN) vaccine in nonhuman primates (NHPs). High-dose (50 μ g)
35 SpFN vaccine, given twice within a 28 day interval, induced a Th1-biased CD4 T cell helper
36 response and a peak neutralizing antibody geometric mean titer of 52,773 against wild-type virus,
37 with activity against SARS-CoV-1 and minimal decrement against variants of concern.
38 Vaccinated animals mounted an anamnestic response upon high-dose SARS-CoV-2 respiratory
39 challenge that translated into rapid elimination of replicating virus in their upper and lower
40 airways and lung parenchyma. SpFN's potent and broad immunogenicity profile and resulting
41 efficacy in NHPs supports its utility as a candidate platform for SARS-like betacoronaviruses.

42

43 **One-Sentence Summary:** A SARS-CoV-2 Spike protein ferritin nanoparticle vaccine, co-
44 formulated with a liposomal adjuvant, elicits broad neutralizing antibody responses that exceed
45 those observed for other major vaccines and rapidly protects against respiratory infection and
46 disease in the upper and lower airways and lung tissue of nonhuman primates.

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58 The coronavirus disease 2019 (Covid-19) pandemic, caused by severe acute respiratory syndrome
59 coronavirus 2 (SARS-CoV-2), has reached a milestone with the emergency use authorization and
60 increasing availability of efficacious vaccines (1). Successes in rapid coronavirus vaccine
61 development, however, have been tempered by the rise of virus variants (2). The accelerating
62 frequency with which variants are emerging raises the prospect that host selective pressures may
63 be driving evolution of mutants to escape vaccine-elicited immunity (3). This concern, coupled
64 with stringent cold-chain requirements for product stability and high unit costs (4, 5), justifies the
65 continued development of cost-effective, thermo-stable vaccines that match current ones in safety
66 and efficacy, but provide broader coverage against a wide range of circulating variants and
67 evolving strains, as well as novel species that may arise from zoonotic reservoirs in the future.

68 Self-assembling protein nanoparticle vaccines offer the advantage of multivalent antigen
69 presentation, a property previously shown to augment immunogenicity over monovalent
70 immunogens (6-8). Ferritin is a naturally occurring, ubiquitous, iron-carrying protein that self-
71 oligomerizes into a 24-unit spherical particle (9). The three-fold axis symmetry of the resulting
72 polymer makes it conducive to conjugation and antigen display of trimeric glycoproteins, such as
73 SARS-CoV-2 Spike (S). Ferritin has been evaluated as a vaccine platform for several pathogens
74 (10-12)—most notably influenza, for which it has demonstrated immune potency and breadth (13,
75 14). As such, ferritin vaccines have advanced to phase 1 clinical trials as a strategy to target
76 multiple influenza strains (15, 16).

77 The prefusion-stabilized form of S is the basis for most major SARS-CoV-2 vaccine
78 candidates (17, 18). Although a correlate of protection from Covid-19 has not been conclusively
79 defined, there is mounting evidence that neutralizing, and some fraction of non-neutralizing,
80 antibodies against S are necessary, if not sufficient, to confer protective immunity (19, 20). The
81 most potent neutralizing antibodies are directed against the S receptor-binding domain (RBD),
82 which mediates attachment to the primary host cell receptor, ACE-2. Prior assessment of a SARS-

83 CoV-2 S ferritin nanoparticle (SpFN) vaccine candidate—co-formulated with a liposomal
84 adjuvant (21)—has demonstrated potent immunogenicity and SARS-CoV-2 protection in mouse
85 models (*unpublished*). These data provide a basis for evaluating SpFN immunogenicity and
86 efficacy against viral replication and pathology in the airways and lungs of nonhuman primates
87 (NHP), a standard model for preclinical evaluation of SARS-CoV-2 vaccines (22).

88

89 **Nanoparticle Vaccine and Study Design**

90 The Spike Ferritin Nanoparticle (SpFN) vaccine was designed as a ferritin-fusion recombinant
91 protein for expression as a nanoparticle. Briefly, the Spike (S) protein sequence was derived from
92 the Wuhan-Hu-1 genome sequence (GenBank accession number: MN908947.3). The
93 S ectodomain was modified to introduce two proline residues (K986P, V987P) and removal of
94 the furin cleavage site (RRAS to GSAS), as previously described (17). To stabilize S trimer
95 formation on the ferritin molecule, the heptad repeat between hinge 1 and 2 (residues 1140 –
96 1161) was mutated to stabilize coiled-coil interactions. An adjuvant—Army Liposomal
97 Formulation QS21 (ALFQ)—was mixed with the SpFN vaccine at room temperature within four
98 hours before administration. ALFQ formulation has been described previously (23). Briefly, it
99 comprises dimyristoyl phosphatidylcholine (DMPC), dimyristoyl phosphatidylglycerol (DMPG),
100 cholesterol (Chol), and synthetic monophosphoryl lipid A (3D-PHAD[®]) (Avanti Polar Lipids,
101 Alabaster, AL) and QS-21 (Desert King, San Diego, CA).

102 In this study, 32 male and female specific-pathogen-free, research-naïve Chinese-origin
103 rhesus macaques (age 3 - 7 years) were distributed—on the basis of age, weight and sex—into 4
104 cohorts of 8 animals (table *S1*). Animals were vaccinated intramuscularly with either 50 or 5 µg
105 of SpFN, formulated with ALFQ, or 1ml of phosphate buffer solution (PBS) in the
106 anterior proximal quadriceps muscle, on alternating sides with each dose in the series.
107 Immunizations were administered twice—4 weeks apart—or once, 4-weeks prior to challenge

108 (fig. *SI*). Animals were challenged with 1×10^6 TCID₅₀ of SARS-CoV-2 (BEI Resources, NIAID,
109 NIH: SARS-Related Coronavirus 2, Isolate USA-WA1/2020, NR-53780 (Lot# 70038893)
110 administered simultaneously by the intratracheal (1.0 ml) and intranasal (0.5 ml per nostril) route.

111

112 **Vaccine Immunogenicity**

113 *Serum antibody responses*

114 We measured longitudinal antibody responses in animals after each vaccination and viral
115 challenge by the Meso Scale Discovery (MSD) electrochemiluminescence platform. Total binding
116 to SARS-CoV-2 prefusion stabilized S protein (S-2P) (17) increased from baseline to an area
117 under the curve (AUC) of 679,213 and 1,646,288 at 4 weeks after two vaccinations with 5 and 50
118 μ g of SpFN, respectively (Fig. 1A). Vaccination with a single 50 μ g dose resulted in a 4-week
119 AUC of 621,605. Binding responses were unchanged in vaccinated groups after viral challenge;
120 whereas, unvaccinated controls had a 200-fold rise. Two-doses of 5 μ g or 50 μ g SpFN elicited
121 reciprocal 50% inhibitory dilution (ID₅₀) neutralizing antibody geometric mean titers (GMT) of
122 22,405 and 52,773, respectively, 2 weeks after second vaccination, and leveled off at 12,171 and
123 22,527 2 weeks later (Fig. 1B). Single dose 50 μ g SpFN elicited a peak GMT of 4063. Authentic
124 virus neutralization activity mirrored group differences seen in the pseudovirus assay, but at
125 somewhat lower values (Fig. 1C).

126 We performed functional assessments of antibody responses by measuring the ability of
127 sera to inhibit binding of RBD to the ACE2 receptor. Binding inhibition in the 5 and 50 μ g
128 vaccinated animals exceeded unvaccinated controls by a factor of 224 and 998, respectively (Fig.
129 1D). ACE2 competition in the single 50 μ g dose group was 291 times higher than controls. We
130 compared humoral responses to the vaccine against a panel of convalescent plasma samples with
131 same pseudovirus neutralization assay; we found that two doses of either SpFN dose elicited

132 neutralizing activity that was an order of magnitude higher than that of the convalescent sera
133 ($p < 0.01$) (Fig. 1E).

134 We used orthogonal approaches to assess binding antibody specificities to the Spike S1
135 subunit domains. RBD and S-2P binding, by MSD, recapitulated results of the ACE2 binding
136 inhibition assay (fig. S2). Serum binding to the N-terminal domain (NTD), which may be a
137 marker of additional protection through both neutralizing activity and non-neutralizing functions
138 (24), were 500-fold higher compared to baseline, across vaccine groups (fig. S3). We assessed the
139 strength of RBD binding by biolayer interferometry, finding an increasing antibody on-rate
140 association response throughout follow-up (fig. S4). Given the potential importance of auxiliary
141 antibody functions for protection (25, 26), we assessed a suite of Fc-mediated antibody effector
142 functions, including opsonization, ADCD, ADCP, ADNP and trogocytosis (a measure of antigen
143 transfer) (27). All activity peaked at week 6 and was highest in the two-dose 50 μ g SpFN group
144 (fig. S5).

145

146 *Cellular immune responses*

147 The character of the helper CD4⁺ T cell (Th) response is important for respiratory virus vaccine
148 development, given the theoretical concern and precedent for vaccine-associated enhanced
149 respiratory disease and its association with a Th2-biased response (28). We focused our
150 assessment of cell-mediated immunity on canonical cytokines expressed by Th1 (interferon-
151 gamma (IFN- γ), tumor necrosis factor (TNF), interleukin 2 (IL-2)) and Th2 (IL-4, IL-13) CD4 T
152 cells. Robust Th1 responses were observed 4 weeks after second vaccination in all vaccinated
153 groups, except one animal in the 50 μ g single-dose group (Fig. 2A). Th1 responses were
154 polyfunctional and variable but consistently high at week 8, ranging from 0.2% to 17%. Th2 and
155 CD8⁺ T cell responses were minimal or undetectable (Fig. 2B, fig. S6A), though Th1 and Th2
156 responses correlated strongly ($r = 0.72$, $p < 0.0001$).

157 We interrogated key indicators of an engaged memory response such as IL-21, a
158 cytokine, secreted by follicular helper CD4⁺ T cells (Tfh), that regulates the evolution of memory
159 B cells (29). Five of eight animals dosed twice with 50 µg SpFN had IL-21 responses, as did
160 seven of eight animals given 5 µg of vaccine (Fig. 2C). We also examined levels of CD40L: a
161 broad T cell activation marker, expressed on the surfaces of CD4⁺ T cells and Tfh cells, that
162 promotes B cell maturation through antibody isotype switching.(29) All but one animal receiving
163 two-doses of SpFN had detectable CD40L⁺ responses (Fig. 2D), indicating an engaged memory
164 response.

165

166 **Protection against high-dose SARS-CoV-2 respiratory challenge**

167 *Virologic Efficacy*

168 Rhesus macaques generally exhibit mild disease that does not recapitulate the severe pneumonia
169 observed in many people with COVID-19 (22). Protective efficacy, therefore, was assessed
170 virologically and pathologically. The primary virologic endpoint was subgenomic mRNA
171 copies/ml—an indicator of viral replication—in the upper (nasopharyngeal (NP) swabs, saliva)
172 and lower airways (bronchoalveolar lavage (BAL) fluid) of vaccinated compared to control
173 animals. On the second day after simultaneous respiratory challenge, sgmRNA levels in the BAL
174 fluid of control animals peaked at a mean of 10⁶ copies/ml (Fig. 3A). In contrast, none of the eight
175 animals that received two doses of 50 µg SpFN had detectable sgmRNA at day 2. By day 4
176 sgmRNA was undetectable in the BAL fluid of all animals of the 5 µg vaccine group and all but
177 one animal that received single SpFN.

178 Whereas sgmRNA levels reached a mean of 10⁷ copies/ml in the NP swabs of control
179 animals at day 2 after challenge, they were undetectable in six of eight animals that received two
180 doses of 50 µg SpFN (Fig. 3B). All animals in that group had undetectable virus from day 4
181 onward, while virus persisted in the control animals through day 10. Five of eight controls had

182 high levels of sgmRNA in saliva on day 2 post-challenge, whereas virus was undetectable in all
183 animals of the two-dose 50 µg SpFN group (Fig. 3C). Total viral load in the BAL fluid, NP swabs
184 and saliva followed trends similar to those for sgmRNA (fig. S7).

185

186 ***Pathologic Efficacy***

187 Unvaccinated control animals developed histopathologic evidence of multifocal, moderate
188 interstitial pneumonia at 7 days after challenge (Fig. 5A). The pneumonia was characterized by
189 type II pneumocyte hyperplasia, alveolar septal thickening, edema and necrotic debris, pulmonary
190 macrophage infiltration and vasculitis of smaller caliber blood vessels. None of the vaccinated
191 animals, however, had evidence of interstitial pneumonia. Immunohistochemistry revealed viral
192 antigen in alveolar pneumocytes and pulmonary macrophages in at least one lung section of every
193 control animal (Fig. 5E). No viral antigen was detected in any vaccinated animals (Fig. 5F-H).

194

195 **Breadth of Immune Response**

196 We assessed the serum antibody responses elicited by the SpFN vaccine against two
197 predominantly circulating SARS-CoV-2 variants of concern (VOC): B.1.1.7 and B.1.351. Serum
198 binding assessment by biolayer interferometry to the variant forms of SARS-CoV-2 RBD showed
199 no change in binding to B.1.1.7 (N501Y mutation) but a 25% reduction in binding to B.1.351
200 (K417N, E484K, N501Y mutations) in the two-dose 50 µg SpFN group, a decrement that was not
201 statistically significant (fig. S8). We next assessed the serum neutralizing activity elicited by the
202 SpFN vaccine against the two VOCs. Sera from all vaccinated NHPs elicited potent neutralizing
203 activity against both variants in two orthogonal virus neutralization assays. Neutralization
204 capacity of the authentic B.1.1.7 virus variant trended higher than against wild-type WA-1 across
205 all vaccine groups (Fig. 5A-C), and was significantly so in the two-dose 50 µg group ($p=0.02$)
206 (Fig. 5A). Neutralizing activity against the authentic B.1.351 virus variant, however, was

207 diminished slightly (Fig. 5A-C), but this difference did not meet statistical significance in the
208 two-dose 50 μ g group (Fig. 5A) and was only marginally significant in the two-dose 5 μ g group
209 ($p=0.05$) (Fig. 5B). Neutralizing activity against the B.1.1.7 pseudovirus in the orthogonal
210 pseudovirus assay revealed statistically equivalent ID₅₀ GMTs to the WA-1 wild-type
211 pseudovirus (Fig. 5D-F). Reductions in neutralizing activity against the B.1.351 pseudovirus,
212 however, were slightly more pronounced in the pseudovirus as compared to the authentic virus
213 assay. For example, the reciprocal ID₅₀ GMT dropped five-fold in the two-dose 50 μ g group but
214 was still high at a value 10,209 (Fig. 5D). The absolute neutralizing antibody titers were generally
215 elevated after two doses of vaccine, irrespective of the virus variant against which they were
216 measured (Fig. 5G).

217 We expanded the assessment of the breadth of SpFN immunogenicity by interrogating the
218 neutralizing and non-neutralizing antibody and cellular immune responses against SARS-CoV-1,
219 Binding of vaccinee sera to SARS-CoV-1 RBD, as measured by biolayer interferometry, was
220 absent in controls but was relatively potent in vaccinated animals—binding at half the strength of
221 that observed to SARS-CoV-2 RBD (fig. S3, S4, Fig. 6A). Antibody-dependent cellular
222 phagocytosis (ADCP) activity also increased remarkably in all vaccine groups, reaching a score
223 that was 100-fold higher two-weeks after last 50 μ g SpFN vaccination than baseline or
224 unvaccinated controls (Fig. 6B). Two vaccinations with high-dose SpFN also yielded significant
225 plaque reduction neutralizing activity against authentic SARS-CoV-1 with a reciprocal ID₅₀
226 GMT of 390 (Fig. 6C) that was significantly above background ($p=0.007$). An orthogonal
227 pseudovirus neutralization assay exhibited significant background activity in PBS controls. We
228 minimized background by analyzing neutralization activity at a 90% inhibitory dilution (ID₉₀)
229 and found that two doses of 50 μ g SpFN induced neutralizing activity against SARS-CoV-1 that
230 was 6-fold higher (GMT 667) than controls (Fig. 6D). CD4⁺ T cell responses to SARS-CoV-1 S
231 peptides, though lower in absolute percentage than to SARS-CoV-2, were still robust and strongly

232 Th1-biased (Fig. 6E,F). The CD8⁺ T cell response, in contrast, was minimal (fig. S9A-C). In
233 aggregate, the immunogenicity profile of SpFN to SARS-CoV-1, though lower in magnitude,
234 recapitulated the quality of response observed to SARS-CoV-2.

235

236 **Discussion**

237 The recent success in the rapid development of safe and efficacious SARS-CoV-2 vaccines has
238 been tempered by the emergence of virus variants to which vaccine-induced immunity has shown
239 diminished potency or efficacy (30-34). There remains a need, therefore, for next-generation
240 vaccines that target the broadening antigenic diversity of SARS-CoV-2 and related coronaviruses.
241 The major vaccines that have progressed to human efficacy trials all present SARS-CoV-2 S
242 based on the genetic sequence of the Wuhan-Hu-1 isolate. All of these vaccines have
243 demonstrated protective efficacy in NHPs against respiratory challenge with the closely matched
244 USA-WA1/2020 (25, 35-39). These earlier animal studies, however, did not evaluate the
245 neutralization capacity of serum against other coronavirus species. In our study of an adjuvanted
246 SpFN vaccine, we recapitulate or surpass the protective efficacy against SARS-CoV-2 infection
247 seen in other studies, but with a greater reduction in replicating virus against a more potent
248 challenge than has been used previously. Additionally, we demonstrate that SpFN elicited
249 antibodies neutralize SARS-CoV-2 ten times more potently than most vaccines and that
250 neutralizing activity is either higher or equivalent against two major VOCs (B.1.1.7, B.1.351) in
251 an authentic virus neutralization assay and equivalent or mildly diminished in a pseudovirus
252 neutralization assay. Finally, SpFN demonstrates neutralization capacity of SARS-CoV-1—a
253 separate species that has 26% and 36% sequence divergence in the S protein and S1 subunit,
254 respectively (40)—above thresholds associated with protection in animals studies (41, 42).

255 SARS-CoV-2 vaccine efficacy studies in NHPs generally compare elicited antibody
256 responses to those from recovered Covid-19 patients. We found neutralizing activity in the two-

257 dose 50 µg SpFN group to be ten-fold higher than that in recovering patients. However, given the
258 preponderance of animal data generated across the vaccine landscape and the lack of
259 standardization across convalescent serum panels, we deemed it more relevant to focus
260 immunogenicity comparisons to published data from other vaccines evaluated in NHP studies.
261 We found that SARS-CoV-2 antibody responses in animals vaccinated with high-dose SpFN were
262 significantly higher than those generated by high doses of leading genetic vaccines (35, 39). That
263 differential increased when SpFN was compared to recombinant adenovirus vector vaccines (25,
264 38). The platform closest to SpFN in composition and design is the adjuvanted S-2P rosette,
265 NVX-CoV2373. In a study of cynomolgus macaques, NVX-CoV2373 induced a SARS-CoV-2
266 antibody responses that exceeded other vaccines but was lower than that generated by SpFN.
267 Direct quantitative comparisons of NHP immunogenicity and efficacy studies, however, can be
268 difficult to interpret, as doses and platforms vary across studies, and immunologic and virologic
269 endpoints are not measured by validated or identical assays. We have made attempts to overcome
270 this limitation by analyzing specimens with orthogonal assays harmonized to consensus
271 platforms. Additionally, our pseudovirus neutralization assay demonstrated equivalence to others
272 in a multi-site concordance survey of reference laboratories.

273 Potent neutralizing antibody responses may offer advantages for both vaccine efficacy and
274 durability. Thus far, neutralizing activity has been predictive of efficacy in human trials, as
275 vaccines that generate lower antibody titers have diminished efficacy (34). An open question
276 remains, however, regarding the length of immunity conferred by SARS-CoV-2 vaccines. For
277 those infectious diseases that are contained by neutralizing antibodies, peak titers have been
278 shown to predict durability (43-45). As such, SpFN may offer longer protection than counterparts;
279 though this requires empirical confirmation.

280 Cross-neutralizing activity against SARS-CoV-2 VOCs is largely diminished for other
281 vaccines at an approximately ten-fold reduction. SpFN induced serum cross-neutralizing

282 responses, however, that were not significantly reduced. Additionally, we found serum binding to
283 mutated SARS-CoV-2 RBD was either unaffected or mildly diminished. Cross-neutralizing
284 activity against SARS-CoV-1 has not been reported yet for other vaccines in advanced
285 development. Some early reports of nanoparticle vaccine approaches presenting RBD have begun
286 to show breadth of neutralizing but these studies either have been limited murine immunogenicity
287 and have yet to demonstrate large animal efficacy or do not generate a full repertoire of multi-site
288 directed antibodies given their restriction to one domain of the S glycoprotein (46-48). We found
289 a comprehensive binding and neutralizing antibody response and a balanced cellular immune
290 response against SARS-CoV-1. Although background neutralizing activity was high in one assay,
291 neutralizing potency against SARS-CoV-1 was confirmed in an orthogonal virus neutralization
292 assay. Still, we are testing purified IgG from vaccine serum in both assays to control for the
293 background levels at baseline and in controls.

294 We hypothesize that breadth of immune response across the SARS betacoronaviruses
295 elicited by the SpFN vaccine may be the result of several factors. First, the quantity of the
296 polyclonal antibody response may surpass a threshold that overcomes resistance to neutralization
297 of antigenically distinct virus variants. Second, repetitive, ordered display of antigen on a self-
298 assembling nanoparticle has been shown to drive an expanded germinal center reaction with
299 resultant increases in B cell receptor mutation, affinity maturation and plasma cell differentiation
300 (6-8). Lastly, the adjuvant, ALFQ, may drive some of the breadth through CD4 T cell activation
301 (49, 50), especially given the high Th1 response elicited by the co-formulation. Epitope mapping
302 and adjuvant comparison studies are underway to dissect the immune potency and breadth.

303 The collective immune response elicited by SpFN translated into a robust and rapid
304 reduction in replicating virus in the upper and lower airways of animals and resultant prevention
305 of pulmonary pathology. It is notable that SpFN protected against a viral challenge that was
306 higher than in any other NHP study to date, as replicating virus levels detected in the upper and

307 lower airways of unvaccinated controls reached a mean of 10^6 to 10^7 copies/ml. Despite this
308 higher challenge, SpFN still protected as rapidly as other leading vaccines. SpFN also quickly
309 eliminated sgRNA in NP swabs, which has implications for decreasing viral shedding and
310 transmission. The absence of an antibody titer boost after challenge suggests a potent anamnestic
311 response and lends additional evidence for near-sterilizing immunity, which again may have
312 implications for preventing viral transmission. Altogether, the immunologic potency and breadth
313 and virologic and pathologic efficacy of SpFN in NHPs support its advancement to evaluation in
314 a phase 1 clinical trial.

315

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435 K.M. and M.G.J designed the study. H.A.K., S.V., N.L.M., D.B. provided additional inputs into
436 study design modifications. I.E.N, A.A., K.K.P., C.M.C., C.S., R.E.C, P.V.T., W-H.C., R.S.S.,
437 A.H., E.J.M., C.E.P., W.C.C., M.C., C.S., P.J.L., A.A., K.M.W., M.D., I.S., J.R.C., K.G.L, V.D.,
438 S.M., K.A., R.C., S.J.K. D.M.P., N.K., V.R.P., Y.H., L.L.J., G.D.G. performed immunologic and
439 virologic assays. H.A.E, A.C., M.G.L. led the clinical care of the animals design and
440 interpretation of data. S.P.D., X.Z., E.K.D performed histopathology. K.M., M.G.J., P.V.T., W-
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442 and A.A. designed and provided the adjuvant. M.G.J., H.A.K. J.A.H., M.G.T., C.S. R.J.O.,
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448

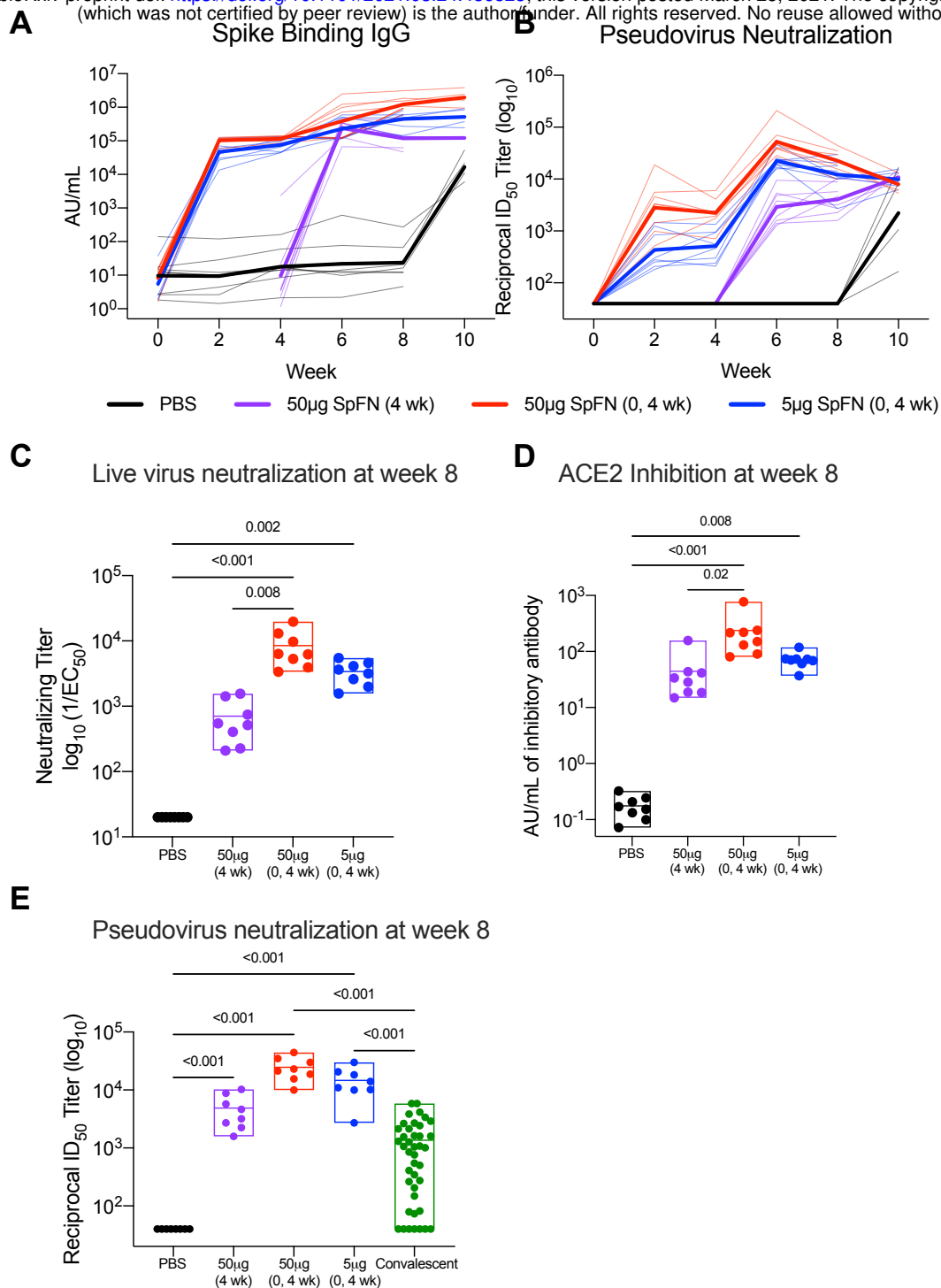
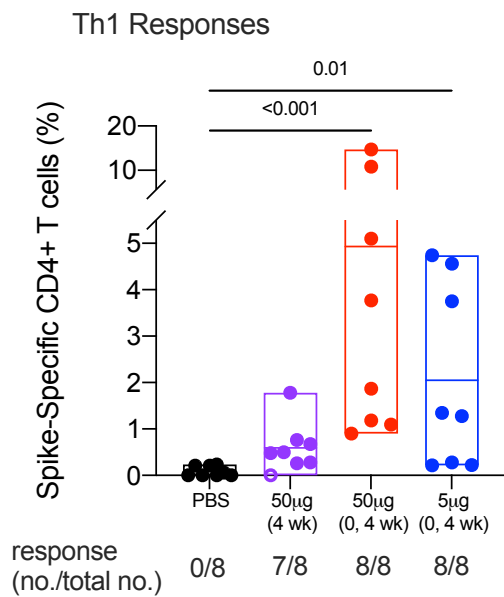


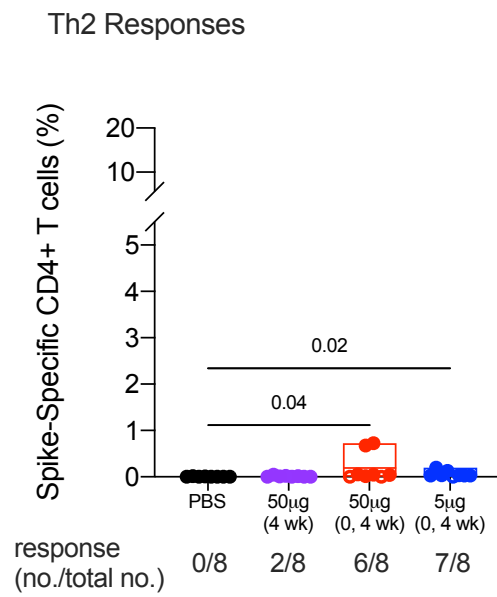
Fig. 1. SpFN elicited binding and neutralizing antibody responses elicited in rhesus macaques.

(A, B) Animals were vaccinated with 5 or 50 μ g of SpFN at weeks 0 and 4 or 50 μ g of SpFN at week 4 only. Control animals were given phosphate-buffered saline (PBS) instead. Serum specimens were assessed for (A) SARS-CoV-2 Spike-specific IgG by the MSD electrochemiluminescent platform and (B) SARS-CoV-2 pseudovirus neutralization every 2 weeks following vaccination and 1 to 2 weeks following viral challenge. Data are depicted as the area under the curve (AUC) of IgG binding and virus neutralization reciprocal 50% inhibitory dilution (ID_{50}), respectively. Thick lines indicate geometric means within each group and thin lines represent individual animals. (C) Authentic virus neutralization was also assessed at 4 weeks after last vaccination. (D) Inhibition of angiotensin-converting enzyme 2 (ACE2) receptor binding to the receptor-binding domain (RBD) at 4 weeks after last vaccination was measured on the MSD platform and reported in arbitrary units (AU)/ml. (E) Pseudoneutralization activity was compared to a panel of human convalescent sera (N=41 samples). In the box plots, horizontal lines indicate the mean and the top and bottom reflect the minimum and maximum. Symbols represent individual animals and overlap with one another for equal values where constrained. Significance was assessed using a Kruskal-Wallis test followed by a Dunn's post-test.

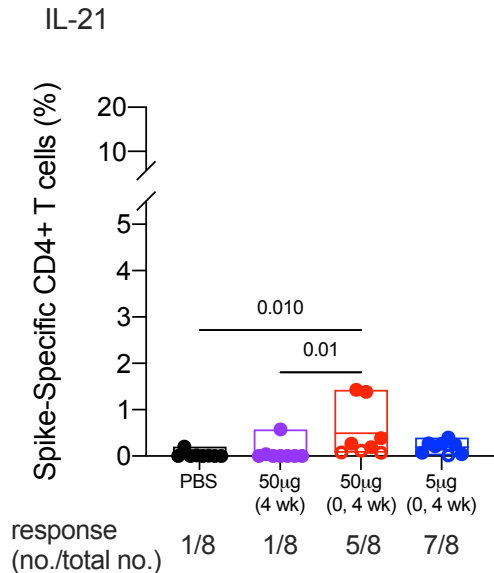
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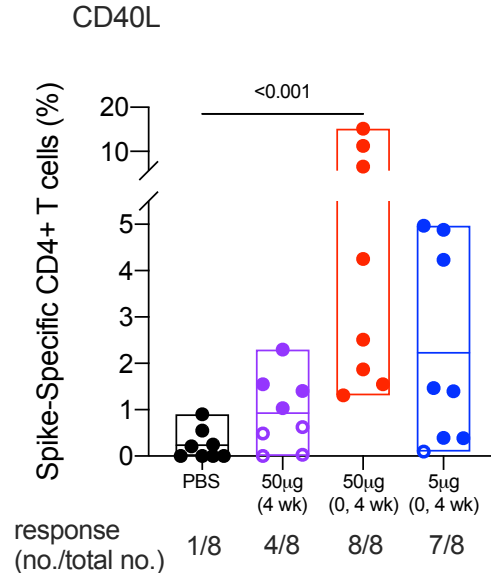


Fig. 2. SpFN vaccine elicited SARS-CoV-2 Spike-specific CD4+ T cell responses in rhesus macaques. T cell responses were assessed by SARS-CoV-2 S peptide pool stimulation and intracellular cytokine staining of peripheral blood mononuclear cells collected at 4 weeks after last vaccination. S-specific memory CD4+ T cells expressing the indicated markers are shown as follows: **(A)** Th1 cytokines (IFN γ , TNF and IL-2); **(B)** Th2 cytokines (IL-4 and IL-13); **(C)** IL-21; and **(D)** CD40L. Boolean combinations of cytokine positive memory CD4+ T cells were summed. Probable positive responses, defined as >3 times the group background at baseline, are depicted as closed symbols. Positivity rates within each group are shown below each graph as a fraction. In the box plots, horizontal lines indicate the mean and the top and bottom reflect the minimum and maximum. Significance was assessed using a Kruskal-Wallis test followed by a Dunn's post-test.

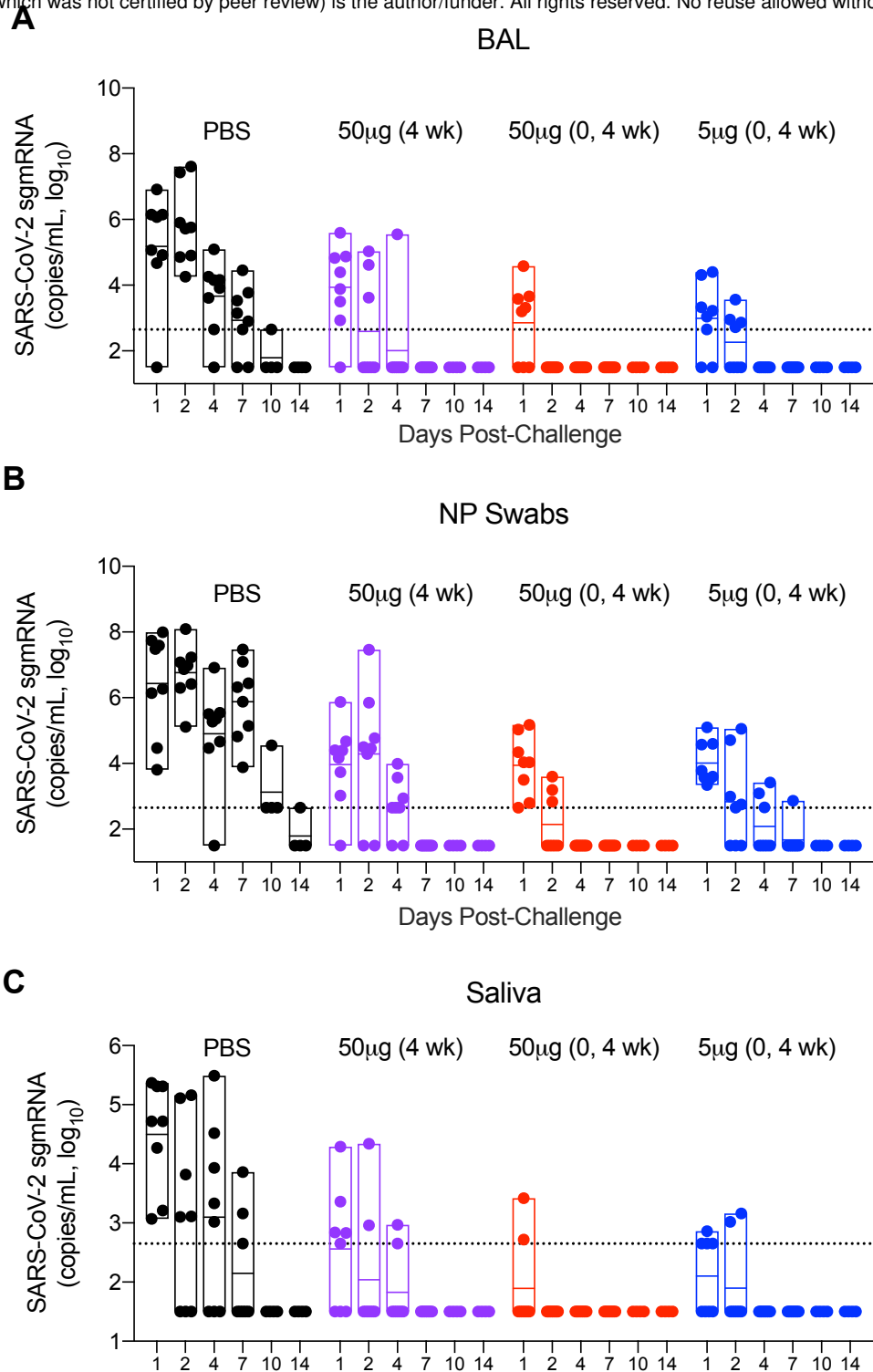


Fig. 3. Viral replication in the lower and upper airways after SpFN vaccination and subsequent SARS-CoV-2 respiratory challenge. Subgenomic messenger RNA (sgmRNA) copies per milliliter were measured in the: (A) bronchoalveolar lavage fluid, (B) nasopharyngeal swabs and (C) saliva of vaccinated and control animals for two weeks following intranasal and intratracheal SARS-CoV-2 (USA-WA1/2020) challenge of vaccinated and control animals. Specimens were collected on 1, 2, 4, 7, 10 and 14 days post-challenge. Dotted lines demarcate assay lower limits of linear performance range (Log₁₀ of 2.65 corresponding to 450 copies/ml). In the box plots, horizontal lines indicate the mean and the top and bottom reflect the minimum and maximum.

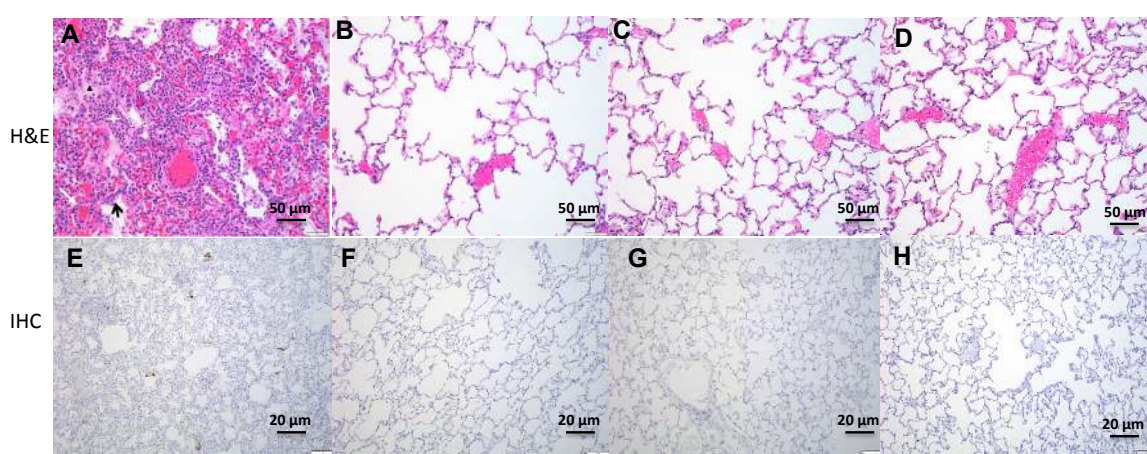


Fig. 4. Histopathology and virus detection in the lungs of SpFN vaccinated and unvaccinated control rhesus macaques following SARS-CoV-2 respiratory challenge. At 7 days post-challenge paraffin-embedded lung parenchymal tissue sections, were (A-D) stained with hematoxylin and eosin (H&E) and (E-H) for immunohistochemistry (IHC). (E) Viral antigen is seen in brown aggregates. Representative images are presented at two magnifications.

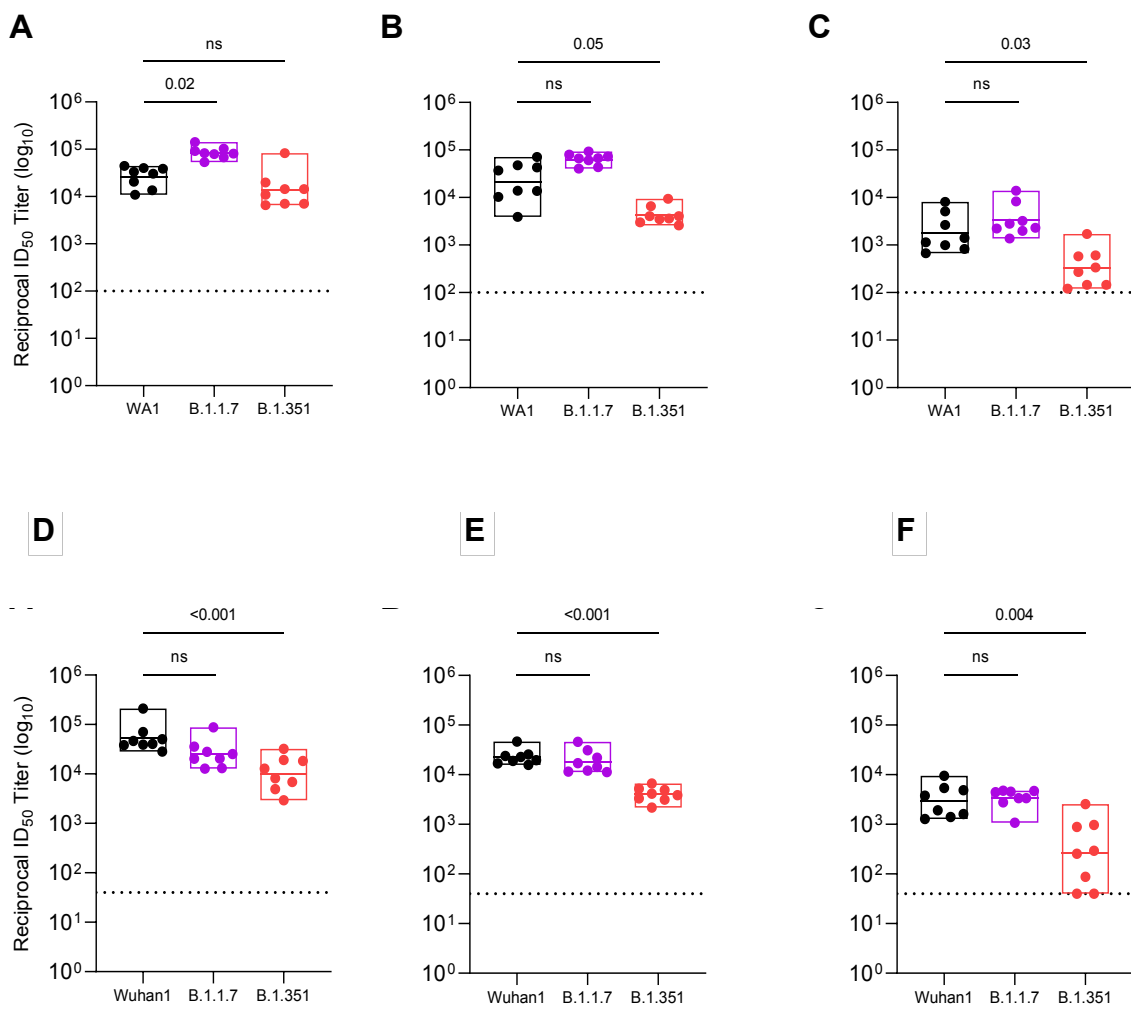
50 µg (0, 4 wk)

5 µg (0, 4 wk)

50 µg (4 wk)

Authentic Virus

Pseudovirus



G

Immunization	Neutralization assay	ID50 GMT			ID50 GMT fold Δ from WA1	
		WA1	B.1.1.7	B.1.351	B.1.1.7	B.1.351
50 µg (0, 4 wk)	Authentic virus	26122	84333	13614	+3.2**	-1.9*
5 µg (0, 4 wk)	Authentic virus	21281	63096	4207	+3.0**	-5.1*
50 µg (4 wk)	Authentic virus	1770	3304	330	+1.9*	-5.4**
50 µg (0, 4 wk)	Pseudovirus	52723	25003	10209	-2.1*	-5.2****
5 µg (0, 4 wk)	Pseudovirus	22387	18197	3990	-1.2*	-5.6****
50 µg (4 wk)	Pseudovirus	2917	3334	264	-1.1*	-11.1***

*non-significant; **p<0.05; ***p<0.01; ****p<0.001

Fig. 5. Pseudovirus and authentic virus neutralizing antibody responses elicited by SpFN vaccination in rhesus macaques against SARS-CoV-2 variants B.1.1.7 and B.1.351, as compared to responses against SARS-CoV-2 WA-1 authentic virus and Wuhan-1 pseudovirus. (A-C) Authentic virus and (D-G) pseudovirus neutralizing antibody responses were measured 2 weeks after last vaccination with either two doses of 50 µg (A, D) or 5 µg (B, E) or one dose of 50 µg (C, F). Reciprocal ID50 GMT fold-change from wild-type neutralization (WA-1 or Wuhan-1) was assessed (G), with statistical significance set at a p-value of < 0.05. Statistical comparisons were done by Kruskal-Wallis test followed by a Dunn's post-test. In the box plots, horizontal lines indicate the mean and the top and bottom reflect the minimum and maximum.

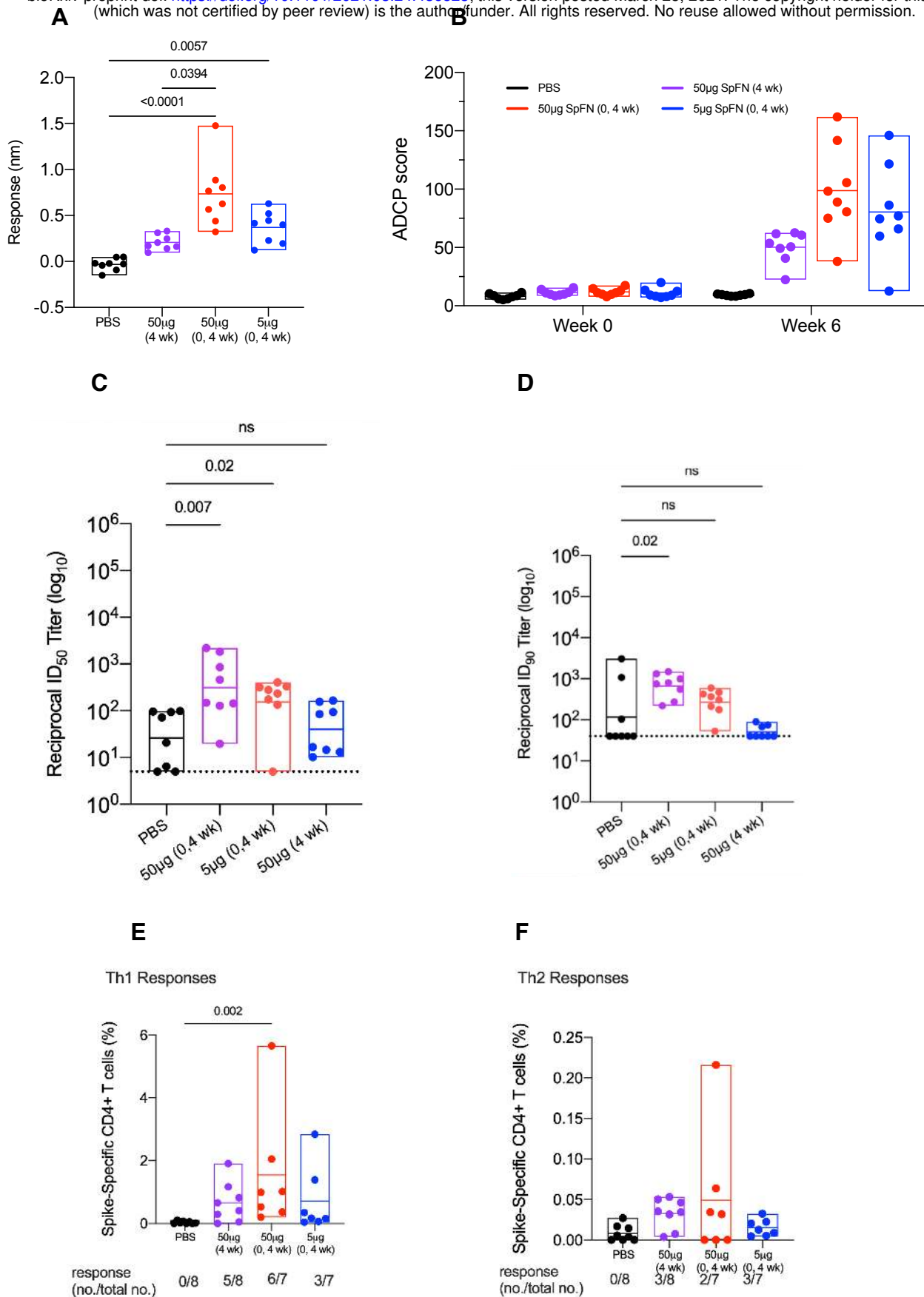


Fig. 6. Humoral and Cellular Immune Responses to SARS-CoV-1 Elicited by SpFN in Rhesus Macaques.

Serum binding responses to SARS-1 RBD by biolayer interferometry, (B) antibody dependent cellular phagocytosis, (C) authentic SARS-CoV-1 (Urbani) neutralization (ID₅₀), (D) pseudo-SARS-CoV-1 (Urbani) neutralization (ID₉₀), (E) SARS-CoV-1 (Urbani) Spike-specific CD4+ Th1 and (F) Th2 response were measured 2 weeks after last vaccination. Significance was assessed with a Kruskal-Wallis test followed by a Dunn's post-test. In the box plots, horizontal lines indicate the mean and the top and bottom reflect the minimum and maximum.