

1 **Long-term Immunity of a Microneedle Array Patch of SARS-CoV-2 S1**
2 **Protein Subunit Vaccine Irradiated by Gamma Rays in Mice**

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32

33 **Abstract**

34 COVID-19 vaccines effectively prevent symptomatic infection and severe disease, including
35 hospitalization and death. However, unequal vaccine distribution during the pandemic, especially
36 in low- and middle-income countries, has led to the emergence of vaccine-resistant strains. This
37 underscores the need for alternative, safe, and thermostable vaccine platforms, such as dissolved
38 microneedle array patches (MAP) delivering a subunit vaccine, which eliminate the need for cold
39 chain and trained healthcare personnel. This study demonstrates that the SARS-CoV-2 S1
40 monomer with RS09, a TLR-4 agonist peptide, serves as an optimal protein subunit immunogen.
41 This combination stimulates a stronger S1-specific immune response, resulting in binding to the
42 membrane-bound spike on the cell surface and ACE2-binding inhibition, compared to the
43 monomer S1 alone or trimer S1, regardless of RS09. MAP delivery of the rS1RS09 subunit
44 vaccine elicited higher and longer-lasting immunity compared to conventional intramuscular
45 injection. S1-specific IgG levels remained significantly elevated for up to 70 weeks post-
46 administration. Additionally, different doses of 5, 15, and 45 μg of MAP vaccines induced robust
47 and sustained Th2-prevalent immune responses, suggesting a dose-sparing effect and inducing
48 significantly high neutralizing antibodies against the Wuhan, Delta, and Omicron variants at 15
49 and 45 μg dose. Moreover, gamma irradiation as a terminal sterilization method did not
50 significantly affect immunogenicity, with irradiated vaccines maintaining comparable efficacy to
51 non-irradiated ones. The stability of MAP vaccines was evaluated after long-term storage at room
52 temperature and refrigeration for 19 months, showing minimal protein degradation. Further, after
53 an additional one-month of storage at elevated temperature (42°C), rS1RS09 in both non-
54 irradiated and irradiated MAP degraded less than 3%, while the liquid subunit vaccine degraded
55 over 23%. Overall, these results indicate that gamma irradiation sterilized MAP-rS1RS09
56 vaccines maintain stability during extended storage without refrigeration, supporting their potential
57 for mass production and widespread use in global vaccination efforts.

58

59 1. Introduction

60 The COVID-19 pandemic has left an indelible mark on the world, with over 775 million
61 confirmed cases and nearly 7 million lives lost globally (until April 14, 2024) [1,2]. The United
62 States alone has witnessed more than 1.18 million deaths [3]. Despite the pivotal role of
63 vaccination efforts in combating the pandemic, challenges persist due to the ever-evolving nature
64 of the SARS-CoV-2 virus, which continuously accumulates mutations in its genetic code. This
65 dynamic leads to the emergence of new variants, some of which may evade immunity conferred
66 by previous vaccinations, rising concern of the development of updated booster shots to address
67 the evolving threat.

68 A critical issue during the pandemic is the unequal distribution of vaccines worldwide.
69 Many low- to middle-income countries have struggled to secure adequate vaccine supplies [4,5],
70 promoting the emergence of vaccine-resistant SARS-CoV-2 strains due to high infection rates in
71 unvaccinated regions [6,7]. In response, alternative delivery methods such as dissolved
72 microneedle array patches (MAP) for subunit vaccine administration are being developed and
73 considered. Indeed, the MAP technology has recently been ranked as the highest global
74 innovation priority for achieving equity of vaccine coverage in low- and middle-income countries
75 by a consortium including the Gavi Secretariat, World Health Organization (WHO), Bill & Melinda
76 Gates Foundation, UNICEF and PATH [8]. MAP delivery offers various advantages, including
77 dose sparing, reduced needle-stick injuries, and the potential for painless self-administered
78 injections, thus alleviating needle phobia in patients [9–12]. Moreover, MAP-based vaccines can
79 be pre-formulated and stored stably for extended periods at room temperature, facilitating
80 distribution even in regions with limited cold chain supply networks [13,14]. Additionally, MAP-
81 based intradermal delivery has shown promising results in improving vaccine immunogenicity and
82 safety in a number of vaccines studies of SARS-CoV-2 [15–20] and have been the subject of
83 Phase I/II clinical trials of influenza, Japanese encephalitis, measles and rubella vaccination

84 [12,21–24]. These clinical trials supported that the MAP vaccination is not only safe and well
85 tolerated but also equal to or more effective than the conventional SC or IM injection.

86 Our prior findings have demonstrated the efficacy of MAP-based platforms in eliciting
87 robust, long-lasting, and cross-neutralizing antibodies against SARS-CoV-2 variants, including
88 Omicron and its subvariants [25]. This longevity in antibody response suggests that MAP-based
89 vaccines could offer durable protection against SARS-CoV-2 and its variants. Therefore, the
90 optimization of variant specific MAP boosters becomes essential to enhance vaccine
91 effectiveness against current and future pandemics. To facilitate the clinical use of MAP platforms,
92 terminal sterilization is a crucial consideration, albeit one that can impact the final product's cost
93 [26]. Various terminal sterilization methods have been developed, including dry-heat, steam,
94 chemical sterilization (with ethylene, formaldehyde, or peracetic acid), and ionizing radiation
95 (electron beam [e-beam], gamma ray [γ -ray], and X-ray) [27–29]. However, among them, gamma
96 radiation of a minimum absorbed dose of 25 kGy is regarded as adequate for sterilizing
97 pharmaceutical due to its ability to destroy microorganisms without heat, moisture, or chemical
98 exposure [30]. However, concerns regarding the potential degradation of MAP morphology or the
99 subunit vaccine within the polymer matrix due to high-energy transfer from gamma-ray irradiation
100 must be addressed. First, we selected an ideal subunit vaccine by comparison the
101 immunogenicity of the SARS-CoV-2 S1 protein depending on whether it was in monomeric or
102 trimeric form, and in the presence of RS09, a TLR4 agonist. Following research aims to evaluate
103 the immunogenicity of MAP delivery compared to conventional intramuscular injection, dose
104 sparing effects, and long-term immunity of irradiated and non-irradiated MAP containing SARS-
105 CoV-2 S1 subunit vaccine in BALB/c mice. Additionally, the stability of the SARS-CoV-2 S1
106 subunit vaccine in both irradiated and non-irradiated forms after long-term shelf storage at
107 different temperatures (4°C, room temperature) was also assessed. Our findings suggest that
108 immunogenicity remains comparable between irradiated and non-irradiated MAP, indicating the
109 feasibility of long-term storage. These studies underscore the potential of gamma irradiation as a

110 preferred method for the terminal sterilization of MAP at a commercial scale, enabling widespread
111 distribution to combat the COVID-19 pandemic on a global scale.

112

113 **2. Materials and methods**

114 *2.1. Construction of Recombinant Protein-expressing Plasmids*

115 The coding sequence for the Delta (B.1.617.2) SARS-CoV-2-S1 glycoprotein mutated
116 T19R; T95I; G142D; Del156-157; R158G; L452R; T478K; D614G, encompassing amino acids 1
117 to 661 of full-length from BetaCoV/Wuhan/IPBCAMS-WH-05/2020 (GISAID accession id.
118 EPI_ISL_403928) [25,31] with C-tag (EPEA) flanked with Sal I and Not I sites was codon-
119 optimized using UpGene algorithm for optimal expression in mammalian cells [32,33] and cloned
120 into pAdlox. Similarly, for S1RS09, spanning amino acids 1 to 661 and equipped with the Bam
121 HI–RS09 (APPHALS, TLR4 agonist)–EPEA, synthesis and cloning into pAdlox were performed
122 using the same method [31]. For S1f and S1fRS09, RS09–EPEA in S1RS09 was replaced with
123 bacteriophage T4 fibritin trimerization domain, foldon (f)–EPEA and fRS09–EPEA at Bam HI and
124 Not I sites, respectively. pAd/S was generated by subcloning the codon-optimized SARS-CoV-2-
125 S2 glycoprotein gene, amino acids 662 to 1273 of full-length spike, into the pAd/S1RS09 at BamH
126 I/Not I sites. The plasmid constructs were confirmed by DNA sequencing.

127

128 *2.2. Recombinant Proteins Expression and Purification*

129 The production of SARS-CoV-2 Delta rS1, rS1RS09, rS1f, and rS1fRS09 involved
130 transient expression in Expi293 cells with pAd/S1, pAd/S1RS09, pAd/S1f and pAd/S1fRS09,
131 respectively, utilizing the ExpiFectamie™ 293 Transfection Kit (ThermoFisher) as previously
132 reported [25,31,34]. Subsequently, the recombinant proteins were purified using a
133 CaptureSelect™ C-tagXL Affinity Matrix prepacked column (ThermoFisher), followed the
134 manufacturer's guidelines as previously detailed [25,31,34]. In brief, the C-tagXL column
135 underwent conditioning with 10 column volumes (CV) of equilibrate/wash buffer (20 mM Tris, pH

136 7.4) before sample application. The supernatant was adjusted to 20 mM Tris with 200 mM Tris
137 (pH 7.4) before loading onto a 5-mL prepacked column at a rate of 5 ml/min. The column
138 underwent subsequent washing cycles, alternating between 10 CV of equilibrate/wash buffer, 10
139 CV of strong wash buffer (20 mM Tris, 1 M NaCl, 0.05% Tween-20, pH 7.4), and 5 CV of
140 equilibrate/wash buffer. The recombinant proteins were eluted from the column using an elution
141 buffer (20 mM Tris, 2 M MgCl₂, pH 7.4). The eluted solution was desalted and concentrated with
142 phosphate buffered saline (PBS) in an Amicon Ultra centrifugal filter device with a 50,000
143 molecular weight cutoff (Millipore). The concentration of the purified recombinant proteins was
144 determined by the BCA protein assay kit (Thermo Scientific) with bovine serum albumin (BSA) as
145 a protein standard. The proteins were separated by reducing sodium dodecyl sulfate
146 polyacrylamide gel electrophoresis (SDS-PAGE) and visualized through silver staining.

147

148 *2.3. SDS-PAGE, Silver Staining, and Western Blot*

149 The purified proteins were subjected to SDS-PAGE and visualized through silver staining
150 and western blot. Briefly, after the supernatants were boiled in Laemmli sample buffer containing
151 2% SDS with beta- mercaptoethanol (β -ME), the proteins were separated by Tris-Glycine SDS-
152 PAGE gels and transferred to nitrocellulose membrane. After blocking for 1 hour at room
153 temperature (RT) with 5% non-fat milk in TBS-T, rabbit anti-SARS-CoV spike polyclonal antibody
154 (1:3000) (Sino Biological) was added and incubated overnight at 4°C as primary antibody, and
155 horseradish peroxidase (HRP)-conjugated goat anti-rabbit IgG (1:10000) (Jackson
156 immunoresearch) was added and incubated at RT for 1 hours as secondary antibody. After
157 washing, the signals were visualized using ECL Western blot substrate reagents and iBright 1500
158 (Thermo Fisher).

159

160 *2.4. Preparation of Dissolving Microneedle Array Patches*

161 Dissolving microneedle array patch containing the protein rS1RS09 were fabricated using
162 the DEN (droplet extension technique) method. In this technique, droplets were solidified and
163 formed into cone shape microstructure by air blowing, as previously described [25,35]. The MAP
164 used in this study was provided by Raphas Co., Ltd. (Rep. of Korea) and fabricated to three
165 different forms changing the array number and patch size depending on dosage. Briefly, for MAP-
166 rS1(WU+Beta), 35 base arrays in 1.86 cm X 1.48 cm (2.75 cm²) for 7 µg of patch and 25 base
167 array in 1.6 cm diameter (2.0 cm²) for 5 µg of patch, respectively, were fabricated. For dose
168 sparing of MAP-rS1RS09, 35 base arrays in 1.86 cm X 1.48 cm (2.75 cm²) for 45 µg of patch, 12
169 and 4 base arrays in 1.6 cm diameter (2.0 cm²) for 15 and 5 µg of patches, respectively, were
170 fabricated. All patches were dispensed onto a pattern-mask hydrocolloid adhesive sheet (Hiks
171 C&T) using a customized MPP-1 dispenser (Musashi) with 10% hyaluronic acid (HA, Kikkoman).
172 The dispensed array was then dried overnight at RT. Subsequently, a viscous solution containing
173 the antigens and HA (Contipro Inc) was dispensed onto a different base array. The two dispensed
174 droplets were brought into contact and extended to the target length, and then symmetric air blow
175 was applied at RT to solidify the extended viscous droplets, forming cone-shaped microstructures.
176 The mechanical strength of each MAP was measured using the universal test machine (UTM,
177 ZwickRoell). Briefly, a MAP was mounted on the UTM stage and aligned with the UTM probe. The
178 probe was moved vertically until the microneedle fractured, with the force responsible for breaking
179 the microneedle recorded as its mechanical strength. For final packaging, each MAP was placed
180 in a polyethylene terephthalate (PET) blister pack ; the blister pack was sealed in light-protective
181 foil pouches (thickness: 100 µm) with desiccant. In line with our efforts toward clinical production
182 of MAP-rS1RS09Delta, we sterilized a small group of these MAPs using gamma irradiation to
183 determine clinically relevant sterilizing conditions and any effect on immunogenicity. Gamma-ray
184 irradiation was conducted using a cobalt-60 irradiator (JS-8900, Nordion Inc.) at 15~30 kGy.

185

186 *2.5. Animal Immunization*

187 Female BALB/c mice (n=5 animals per group) were bled from the retro-orbital vein before
188 immunization and primed with 45 μ g of either Delta rS1, rS1RS09, rS1f, or rS1fRS09. Mice were
189 bled on week 3 and received a homologous booster of 45 μ g of each protein. Subsequent bleeds
190 were performed on weeks 5, 7, and 18. For the study of long-term immunogenicity, BALB/c mice
191 (5 per group) were primed and boosted at three-week intervals with 45 μ g of rS1RS09 intramuscularly.
192 Serum samples were collected in weeks 0, 3, 5, 7, 9, 12, 16, 20, 28, 40, 71, 90, and 104 after prime
193 immunization.

194 In the experiment with ICR mice, 5 μ g of rS1(Wu+Beta) were injected into the thigh
195 intramuscularly, and MAPs loaded with either 5 or 7 μ g of rS1(Wu+Beta) were applied to the skin
196 of the back region of female ICR mice (n = 5 animals per group). Prior to vaccination, the hair at
197 the vaccination site was removed by shaving and depilatory cream. MAPs were applied to the
198 dehaired back skin of the mice using a handheld spring applicator and held in place by finger
199 pressure for 10 seconds, followed by leaving it on the skin for more than 2 hours. No adverse skin
200 effects, such as skin irritation at the vaccinated region, were observed. At 3 weeks after the
201 primary immunization, mice received booster immunizations with homologous immunogens.
202 Blood samples were collected from the retro-orbital vein of mice every three weeks until week 9.
203 The obtained serum samples were diluted and used to evaluate rS1-specific antibodies by ELISA.

204 MAPs loaded with 5, 15, and 45 μ g of Delta rS1RS09, referred to as MAP-rS1RS09 with
205 or without irradiation, were applied to the skin of the back region of female BALB/c mice (n = 5
206 animals per group). MAP application was performed using the same method as with ICR mice. At
207 3 weeks after the primary immunization, mice were received booster immunizations with
208 homologous immunogens. Blood samples were collected from the retro-orbital vein of mice every
209 three to ten weeks until week 104. The obtained serum samples were diluted and used to evaluate
210 rS1-specific antibodies by ELISA and VNT assay. Since aged mice develop spontaneous
211 leukemias and other tumors, the dedicated veterinarians oversee the animals' physical and

212 psychological health and ruled out any mice having diseases that may influence immune
213 responses. Indeed, one mouse of 15 μg non-irradiated MAP-rS1RS09 group was ruled out at
214 week 71, and two to three mice were ruled out in week 90, and two to four mice in week 104,
215 because they were euthanized due to tumors or were found deceased. Mice were maintained
216 under specific pathogen-free conditions at the University of Pittsburgh, and all experiments were
217 conducted in accordance with animal use guidelines and protocols approved by the University of
218 Pittsburgh's Institutional Animal Care and Use (IACUC) Committee.

219

220 *2.6. Assessment of Serum Humoral Antibodies*

221 Serum humoral antibodies generated against spike protein were assessed using ELISA,
222 as previously described [32,36]. Sera from all mice were collected before vaccination and then
223 every 2 to 10 weeks after immunization. These sera were tested for SARS-CoV-2 S1WU-specific
224 IgG antibodies using conventional ELISA. Furthermore, sera collected at weeks 0, 9, 28, 52, and
225 71 or at weeks 0, 5/6, 20, 40, and 71 after vaccination were also examined for SARS-CoV-2-S1-
226 specific IgG1 and IgG2a antibodies using ELISA. In brief, ELISA plates were coated with 200 ng
227 of recombinant SARS-CoV-2-S1WU protein (Sino Biological) or house purified rS1WU per well
228 overnight at 4°C in carbonate coating buffer (pH 9.5) and then blocked with PBS containing 0.05%
229 Tween 20 (PBS-T) and 2% BSA for one hour. Mouse sera were serially diluted in PBS-T with 1%
230 BSA and incubated overnight. After washing the plates, anti-mouse IgG-horseradish peroxidase
231 (HRP) (1:10000, Jackson ImmunoResearch) was added to each well and incubated for one hour.
232 For detection of IgG1 and IgG2a, HRP-conjugated anti-mouse IgG1 and IgG2a (1:20000, Jackson
233 ImmunoResearch) were added to each well and incubated for 1 hour. The plates were washed
234 three times, developed with 3,3',5,5'-tetramethylbenzidine, and the reaction was stopped. Optical
235 densities (ODs) were read at 450 nm with a SpectraMax iD5 microplate reader (Molecular
236 Devices).

237

238 2.7. Flow Cytometry

239 Two weeks after the booster immunization, pooled sera were obtained from all mice and
240 screened for SARS-CoV-2-S-specific antibodies using fluorescence-activated cell sorter (FACS)
241 analysis of Expi293 cells transfected with either pAd/S or pAd control using ExpiFectamie™ 293
242 as described previously. Briefly, at 36 hours after transfection, the cells were harvested, counted,
243 and washed with PBS. Half a million cells were incubated with 1 μ l of mouse serum from each
244 group or 1 μ l of monoclonal antibody (D003, Sino Biological) for 30 min, followed by staining with
245 a FITC-conjugated anti-mouse secondary IgG antibody (Jackson ImmunoResearch). Data
246 acquisition and analysis were performed using LSRII (BD). The results were calculated as follows:
247 Positive cells (%) = 100 \times (positive cell (%) using each mouse serum / positive cell (%) using
248 monoclonal antibody).

249

250 2.8. ACE2 Blocking Assay

251 Antibodies blocking the binding of SARS-CoV-2 spike including Wuhan, Omicron (BA.1),
252 Omicron sub-variants (BA.2, BA.3, BA.1+R346K, BA.1+L452R), Delta lineage (AY.4), Alpha
253 (B.1.1.7), Beta (B.1.351), and France (B.1.640.2) to ACE2 were detected with a V-PLEX SARS-
254 CoV-2 Panel 25 (ACE2) Kit (Meso Scale Discovery (MSD)) according to the manufacturer's
255 instructions. The assay plate was blocked for 30 min and washed. Serum samples were diluted
256 (1:25 or 1:100) and 25 μ l were transferred to each well. The plate was then incubated at RT for
257 60 min with shaking at 700 rpm, followed by the addition of SULFO-TAG conjugated ACE2, and
258 continued incubation with shaking for 60 min. The plate was washed, 150 μ l MSD GOLD Read
259 Buffer B was added to each well, and the plate was read using the QuickPlex SQ 120 Imager.
260 Electrochemiluminescent values (ECL) were generated for each sample. The lower values than
261 pre-immunized sera were adjusted with the values at week 0. Results were calculated as %
262 inhibition compared to the negative control for the ACE2 inhibition assay, and % inhibition is
263 calculated as follows: % neutralization = 100 \times (1 - (sample signal/negative control signal)).

264

265 2.9. SARS-CoV-2 microneutralization assay

266 Neutralizing antibody titers against SARS-CoV-2 were defined according to the following
267 protocol (59, 60). Briefly, 50 µl of sample from each mouse, starting from 1:10 in a twofold dilution,
268 were added in two wells of a flat bottom tissue culture microtiter plate (COSTAR), mixed with an
269 equal volume of 100 TCID₅₀ of a SARS-CoV-2 Wuhan, Delta (B.1.617.2), or Omicron (BA.1) strain
270 isolated from symptomatic patients, previously titrated. After 1 hour incubation at 33°C in 5% CO₂,
271 3 x 10⁴ VERO E6 cells were added to each well. After 72 h of incubation at 33°C 5% CO₂, wells
272 were stained with Gram's crystal violet solution plus 5% formaldehyde 40% m/v (Carlo ErbaSpA,
273 Arese, Italy) for 30 min. After washing, wells were scored to evaluate the degree of cytopathic
274 effect (CPE) compared to the virus control. Neutralizing titer was the maximum dilution with a
275 reduction of 90% of CPE. A positive titer was equal to or greater than 1:10. The GMT of VNT₉₀
276 endpoint titer was calculated with 5 as a negative shown <1:10. Sera from mice before vaccine
277 administration were always included in VNT assay as a negative control.

278

279 2.10. MAP Storage and Reconstitution for Analysis

280 The packed MAPs, stored in light-protective pouches with desiccant, were kept in a
281 refrigerator at 4 °C or in a drawer at room temperature (RT) for one week or nineteen months. To
282 assess long-term stability, the packed MAPs stored at 4 °C for nineteen months were further
283 stored for one month in a refrigerator at 4 °C, in a drawer at RT, and in temperature-controlled
284 incubator at 42 °C to represent storage conditions in a hot climate. MAPs were removed from the
285 pouches and placed in each well of a 6-well plate with 2 ml of PBS. Reconstitution was achieved
286 by incubating for 30 min at RT with shaking at 700 rpm. The resulting solution containing rS1RS09
287 and microneedle matrix excipients was transferred to a centrifuge tube and stored at -80 °C until
288 further analysis. The stability assay was performed using SDS-PAGE and silver staining, followed
289 by analysis using ImageJ v1.47 software (National Institutes of Health, Bethesda, MD, USA) to

290 determine the relative intensity of each protein band for comparison. The results are presented
291 as the percentage of degradation proteins relative to the smear ratio, which is calculated by
292 dividing the integrated density of the sample band area by that of all protein bands in the
293 preparation lane. The percentage of recombinant protein degradation in the reconstructed MAP
294 or in the buffer was calculated as follows: % protein degradation = $100 \times [($ smear ratio of negative
295 control – smear ratio of sample MAP)/smear ratio of negative control]. The negative control
296 comprised non-irradiated MAP stored at 4°C or recombinant proteins stored at -20°C for 19
297 months. To compare the stability of rS1RS09 in the buffer solution, recombinant proteins (rP)
298 stored for 19 months at -20°C were 10-fold diluted in PBS or in the preservative buffer (10mM
299 Tris, 75mM NaCl, 2mM MgCl₂, 5% Trehalose), divided into two microtubes, and further stored for
300 one month at -20°C or 42°C in a humidity chamber. The percentage of recombinant protein
301 degradation was calculated in the same way as above.

302

303 *2.11. Statistical analysis*

304 Statistical analyses were performed using GraphPad Prism v10 (San Diego, CA). Antibody
305 endpoint titers and neutralization data were analyzed by Kruskal-Wallis test, followed by Dunn's
306 multiple comparisons. A Mann-Whitney U test was used for intergroup statistical comparison.
307 Comparison of intramuscular injection and MAP groups were analyzed by Bartlett's test. Data
308 were presented as the means \pm standard errors of the mean (SEM) or geometric means
309 \pm geometric errors. Significant differences are indicated by * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$.

310

311 **3. Results**

312 *3.1. Purification of Recombinant Proteins*

313 Initially, we compared the immunogenicity of the SARS-CoV-2 S1 protein in monomeric
314 versus trimeric forms, and in the presence of RS09, a TLR4 agonist. To produce four recombinant
315 proteins of the Delta SARS-CoV-2-S1, both in the absence and presence of the bacteriophage

316 T4 trimer domain foldon (f) and RS09, we generated pAd/S1, pAd/S1RS09, pAd/S1f, and
317 pAd/S1fRS09 by subcloning the codon-optimized Delta SARS-CoV-2-S1 gene with a C-tag
318 (EPEA) into the shuttle vector pAd (GenBank U62024) at the *Sal* I and *Not* I sites (**Fig. 1A**).
319 Variant-specific mutations for SARS-CoV-2 Delta (B.1.617.2) S1 proteins are outlined. For the
320 expression of the four Delta S1 subunit proteins, Expi293 cells were transfected with the plasmids
321 pAd/S1, pAd/S1RS09, pAd/S1f, and pAd/S1fRS09. Five days after transfection, the supernatants
322 of Expi293 cells were collected, and the four recombinant proteins (Delta rS1, rS1RS09, rS1f, and
323 rS1fRS09) were purified using C-tagXL affinity matrix and confirmed by silver staining and
324 Western blot (**Fig. 1B and 1C**). Each S1 recombinant protein was recognized by a polyclonal anti-
325 spike of SARS-CoV-2 antibody at the expected glycosylated monomeric molecular weights of
326 about 110 kDa under denaturing reduced conditions. These results indicates that four
327 recombinant proteins (Delta rS1, rS1RS09, rS1f, and rS1fRS09) were successfully purified for
328 subsequent experiments.

329

330 *3.2. Identification of S1RS09 as an optimal immunogen*

331 To compare the immunogenicity of four Delta S1 proteins, BALB/c mice (6–8 weeks old,
332 n = 5) were immunized intramuscularly (IM) with 45 μ g of either Delta rS1, rS1RS09, rS1f, or
333 rS1fRS09 in a prime-boost regimen, spaced three weeks apart. Blood samples were collected
334 before the initial immunization and at weeks 3, 5, 7, and 18, and then examined for the presence
335 of antibodies specific to SARS-CoV-2-S1 using ELISA (**Fig. 2A and 2B**). As shown in Fig. 2B,
336 statistically significant S1-specific IgG EPT was induced in the mice immunized with either S1 or
337 S1RS09 at weeks 5, 7 and 18, while the mice vaccinated with trimers, rS1f or rS1fRS09 showed
338 lower S1-specific IgG EPT. Interestingly, more statistically significant S1-specific IgG EPT was
339 induced in the mice immunized with S1RS09 all time points (**Fig. 2B**). These findings suggest
340 that SARS-CoV-2 S1 monomer in the presence of RS09 (rS1RS09) is more immunogenic than
341 the trimer S1 subunit protein regardless of RS09.

342 We also examined the S1-specific IgG isotypes switch, IgG1 and IgG2a following
343 immunization with the four Delta S1 protein subunits, indicating a prevalent and/or balanced Th2-
344 or Th1-biased immune response, respectively (**Fig. 2C and D**). Sera collected at weeks 0, 7, and
345 18 post-prime were subjected to isotype-specific ELISA. The S1-specific IgG1 showed a
346 significant increase in the mice immunized with either S1 or S1RS09 at week 7 and maintained
347 significantly high levels until week 18 comparison to pre-immunized sera, although the geometric
348 mean titer (GMT) diverged slightly lower in the S1 group and slightly higher in the S1RS09 group
349 (**Fig. 2C**). In case of S1-specific IgG2a, only S1-immunized mice showed a significant increase
350 at week 7, whereas significant increases were detected in the mice immunized either S1 or S1f
351 at week 18. Interestingly, the GMT of IgG2a increased slightly at week 18 in all groups compared
352 to those at week 7 (**Fig. 2D**). Th2-prevalent responses were observed at all time points based on
353 the ratio of IgG1 to IgG2a antibody subclasses in all the groups at week 7, with slightly higher
354 ratios in the S1RS09 group. It was observed that the S1RS09 group induced more toward Th2-
355 biased immune response by higher IgG1 and similar IgG2a at week 18, while the three other
356 groups switched slightly toward a Th1 response or remained steady (**Fig. 2E**). The ratio of
357 IgG1/IgG2a at week 18 was approximately 10-fold higher when the vaccine was administered
358 with S1RS09 compared to the S1 group.

359 Next, we examined whether these antibodies could bind to membrane-bound full-length
360 spike protein by measuring the reactivity on Expi293 cells transfected with pAd/S (S1Delta +
361 S2Wuhan) or pAd using flow cytometry. The mice immunized with either Delta rS1, rS1RS09,
362 rS1f, or rS1fRS09 developed membrane-bound S-specific antibodies, while no specific antibody
363 response was detected in pre-immunized sera (**Fig. 2F**). The mean of percentage of positive cells
364 was slightly higher in the rS1RS09-immunized mouse group compared to other groups. No
365 antibodies were detected on cells transfected with pAd (data not shown).

366 To assess the capacity of these vaccines to potentially neutralize RBD-ACE2 binding, we
367 evaluated the ability of antibodies in the serum at week 7 to inhibit the binding between ACE2 and

368 the trimeric spike protein of SARS-CoV-2 variants, representing a sensitive measure of
369 neutralizing activity. We used V-PLEX SARS-CoV-2 (ACE2) Kit Panel 18, which included Wuhan,
370 Alpha (B.1.1.7), Beta (B.1.351), Gamma (P.1), Delta (B.1.617.2), Zeta (P.2), Kappa (B.1.617.1),
371 New York (B.1.516.1), India (B.1.617 and B.1.617.3) variants. The ability of antibodies to
372 neutralize the interaction between spike protein of SARS-CoV-2 variants and ACE2 was
373 examined in all animals immunized either rS1, rS1RS09, rS1f, or rS1fRS09 at weeks 0 and 7
374 post-prime at a dilution of 1:20. The ACE2 inhibitory activities of the sera from the mice immunized
375 with Delta rS1, rS1RS09, rS1f, or rS1fRS09 against all variants were on average $50.9\% \pm 5.64$,
376 $57.1\% \pm 12.21$, $32.4\% \pm 7.42$, and $35.4\% \pm 5.53$ at weeks 7, respectively, with $21.2\% \pm 7.15$ at
377 week 0. Overall, the median percent inhibition of the sera from rS1RS09-immunized groups was
378 the highest against all variants among the groups, except against Beta and Gamma, which
379 showed the highest inhibition in rS1-immunized group. Interestingly, the inhibitions against
380 Wuhan, Alpha, Kappa, and India (B.1.617.3) spike proteins were significantly different compared
381 to week 0 in rS1-immunized mice. The ACE2-binding inhibitions of rS1RS09-immunized mice
382 were significantly increased for all tested SARS-CoV-2 variants, except Beta and Gamma when
383 compared to week 0 (**Fig. 3**). They demonstrated moderate ACE2-binding inhibition from rS1f-
384 and rS1fRS09-immunized sera, which were not statistically significant against all variants when
385 compared to week 0. These findings suggest that SARS-CoV-2 S1 monomer in the presence of
386 RS09 (rS1RS09) is more immunogenic than the trimer S1 subunit protein, regardless of RS09,
387 considering it as an ideal vaccine candidate.

388

389 *3.3. Enhanced Immunity of MAP delivery compared to conventional intramuscular injection*

390 In a previous investigation, we demonstrated the effectiveness of a prime-boost SARS-
391 CoV-2 S1 subunit MAP vaccine in eliciting specific antibody responses in mice [25]. We also
392 compared the antibody responses induced by MAP delivered S1 subunit vaccine to those elicited
393 by conventional intramuscular (IM) injections. Eight-week-old ICR mice were immunized with 5

394 μg of rS1(Wu+Beta) (Group 1) intramuscularly (IM) into the thigh, while MAPs loaded with either
395 5 or 7 μg of rS1(Wu+Beta) (Group 2 and 3, respectively) were inoculated intradermally (ID) three
396 weeks apart (**Fig. 4A and 4B**). We measured S1-specific total IgG antibody endpoint titer (EPT)
397 at weeks 3, 6, and 9 after the initial vaccination by ELISA. S1-specific antibodies in mice
398 immunized with MAP-S1(Wu+Beta) (Group 2 and 3) showed significantly elevated IgG levels at
399 weeks 6 and 9 following the booster vaccination compared to pre-immunized controls, whereas
400 antibody levels in sera from IM-immunized mice (Group 1) were not significantly elevated
401 compared to pre-immunized sera and declined slightly at week 9 (**Fig. 4C**). Furthermore, we
402 examined the S1-specific IgG1 and IgG2a titers at week 6 to determine if there were significant
403 differences due to the vaccine dose and route of immunization. IgG1 endpoint titers of Group 2
404 and Group 3 were statistically significant when compared to pre-immunized controls, while only
405 Group 3 IgG2a endpoint titers were statistically significant (**Fig. 4D and 4E**). However, similar
406 Th2-prevalent responses were observed regardless vaccine dose and route of immunization
407 based on the ratio of IgG1 to IgG2a antibodies in all the groups (**Fig. 4F**). These findings suggest
408 that the delivery of protein subunit vaccine via MAP is superior to conventional intramuscular
409 injection, although similar Th2-prevalent responses are observed.

410

411 *3.4. Long-lasting Immunity of MAP delivery compared to conventional intramuscular injection*

412 In a previous study, we demonstrated that a prime-boost SARS-CoV-2 S1 subunit MAP
413 vaccine was more effective in triggering specific antibody responses in mice compared to
414 intramuscular injection. To further evaluate the differences due to the vaccine dose and longevity,
415 we generated and purified large-scale of the S1RS09 recombinant proteins and fabricated MAPs
416 loaded 45 μg of rS1RS09 (MAP-rS1RS09) using a hyaluronic acid-based droplet extension
417 technique (DEN), which offers the benefits of being faster, scalable, and cost-effective (**Fig. 5A**).
418 The DEN MAP features a rectangle design with dimensions of 2 cm x 1.18 cm, housing a grid of
419 7x5 microneedle arrays, each measuring approximately 700 μm in length. The MAP consists of a

420 double layered form with a base layer measuring 900 μm in base width and $200 \pm 20 \mu\text{m}$ in height,
421 and an upper layer measuring $500 \pm 20 \mu\text{m}$ in length. Each MAP occupied an area size of 2.36
422 cm^2 , with 7×5 arrays of approximately $700 \pm 50 \mu\text{m}$ total length (**Fig. 5B and 5C**). The mechanical
423 strength (fracture force) of the MAP was measured to be more than 0.87 N, which is the required
424 strength to penetrate the skin.

425 To compare the long-term immunogenicity in the two different routes, BALB/c mice (5 per
426 group) were primed and boosted at three-week intervals with either 45 μg of rS1RS09
427 intramuscularly or MAP-rS1RS09 intradermally. Serum samples were collected at weeks 0, 3, 5/6,
428 7, 9, 12, 16, 20, 28, 40, and 70 after prime immunization (**Fig. 5D**). Serum samples were serially
429 diluted to determine SARS-CoV-2-S1-specific IgG titers using ELISA. All vaccinated groups had
430 significantly high and similar geometric mean S1 IgG antibody endpoint titer (EPT) at weeks 5
431 and 6 when compared to week 0, illustrating the superior immunogenicity conferred by boost
432 immunization. The IgG geometric mean in mice immunized with MAP-rS1RS09 reached a steady
433 geometric mean S1 IgG EPT through week 40, as high as the peak at week 6, and waned slightly
434 thereafter. In contrast, mice immunized with rS1RS09 IM peaked at week 5, similar to the MAP
435 group, but the immune response steadily waned through week 70, with significant difference
436 between the two groups ($p < 0.01$, Bartlett's test) (**Fig. 5E**). S1-specific antibodies in mice
437 immunized with MAP-rS1RS09 remained significantly elevated compared to pre-immunized sera
438 for at least 70 weeks, whereas those in IM-immunized mice were statistically significant until week
439 9 and no longer statistically significant after that time point (**Fig. 5E**).

440 Furthermore, we examined the S1-specific IgG1 and IgG2a titers at weeks 5/6, 20, 40,
441 and 70 to determine if there were significant differences due to the route of immunization and
442 longevity. IgG1 and IgG2a EPT of the MAP immunized group were statistically significant
443 compared to pre-immunized controls at some tested time points (**Fig. 5F and 5G**). Interestingly,
444 the mean EPT of IgG1 in MAP ID group increased significantly at later time points, at weeks 40
445 and 70, while those in the IM group decreased. The mean EPT of IgG2a in the MAP ID group

446 were increased steadily until week 40 and dropped at week 70, while the IM group showed steady
447 mean titers. Based on the ratio of IgG1 to IgG2a antibodies, similar Th2-prevalent responses were
448 observed regardless of vaccine dose and route of immunization at all time points, except for a
449 slight shift towards to a Th2-biased immune response at week 70 in the MAP ID group due to the
450 steady IgG1 and slight drop in IgG2a (**Fig. 5H**). These results suggest that MAP intradermal
451 delivery induces potent and long-lasting SARS-CoV-2 S1-specific antibody responses compared
452 to subunit intramuscular injection, while showing similar Th2-prevalent responses.

453 454 *3.5. Dose sparing and gamma irradiation for sterilization*

455 Next, we investigated the immunogenicity of MAP-rS1RS09 through dose sparing and
456 gamma irradiation (25 kGy) sterilization as part of our efforts toward clinical-grade manufacturing
457 of MAP subunit vaccines. Gamma irradiation has been previously employed as a terminal
458 sterilization method for MAP-SARS-CoV-2 subunit vaccines [37]. For dose sparing, we produced
459 5, 15, and 45 μg of MAP-rS1RS09 by adjusting the number of microneedle arrays, and some of
460 these MAPs were sterilized by gamma irradiation (**Fig. 6A**). BALB/c mice (5 per group) were
461 primed and boosted at three-week intervals with either non-irradiated (-) or irradiated (+) 5, 15, or
462 45 μg of MAP-rS1RS09 intradermally (**Fig. 6B**). Serum samples were collected at weeks 0, 3, 6,
463 9, 12, 16, 28, 52, 71, 90, and 104 after prime immunization, and the S1-specific IgG EPT was
464 examined by ELISA. All vaccinated groups exhibited higher S1-specific IgG EPT at week 6
465 compared to week 3, with only 15 and 45 μg MAP-rS1RS09 groups showing significant increases.
466 This significant increase was sustained even at week 71 post-prime vaccination when compared
467 to pre-immunized sera, indicating a robust and sustained immunogenic response, consistent with
468 the promising outcomes observed in the earlier experiment (**Fig. 6C**). Interestingly, the MAP-
469 rS1RS09 group maintained a steady geometric mean S1 IgG EPT through week 28, reaching
470 levels as high as the peak at week 6, with minimal waning of the immune response through week
471 104 post-prime. The mean IgG EPT at the final week 104 showed dose-dependence, although

472 statistical significance was weak due to the low number of animals. Notably, the immunogenicity
473 of gamma irradiation-sterilized MAP vaccines was comparable to that of non-irradiated MAP
474 vaccines in all vaccine groups (**Fig. 6D**). These findings support the feasibility of gamma
475 irradiation as a terminal sterilization approach for our clinical MAP-rS1RS09 vaccines.

476 We also examined the S1-specific IgG1 and IgG2a titers to determine if there were
477 significant differences in the ratio of these subtypes due to vaccine dose and gamma irradiation.
478 Sera collected at weeks 0, 9, 28, 52, 71, 90, and 104 after prime-boost immunization with either
479 non-irradiated (-) or irradiated (+) of 5 or 45 μg of MAP-rS1RS09 were subjected to isotype-
480 specific ELISA. The IgG1 subclasses of the 5 μg MAP-rS1RS09 group showed a significant
481 increase at weeks 9 in the non-irradiated group and at week 52 in the irradiated group after initial
482 immunization compared to pre-immunized sera (**Fig. 7A**). The IgG2a subclasses of the non-
483 irradiated 5 μg MAP-rS1RS09 group increased significantly at weeks 52 and 71, while those of
484 the irradiated MAP group increased at weeks 28, 52 and 71 (**Fig. 7B**). Interestingly, mean titers
485 of IgG1 and IgG2a in both non-irradiated and irradiated groups remained steady at most time
486 points, with reduced titer at week 104, indicating a shift toward a Th2-biased immune response
487 (**Fig. 7C**).

488 The IgG1 subclasses of both the non-irradiated and irradiated 45 μg MAP-rS1RS09
489 groups showed a significant increase at weeks 52 and 71 after initial immunization compared to
490 pre-immunized sera (**Fig. 7D**). The IgG2a subclasses of the non-irradiated 45 μg MAP-rS1RS09
491 group increased significantly at weeks 28 and 52, while those of the irradiated MAP group
492 increased at weeks 9, 28, and 52 (**Fig. 7E**). Interestingly, the mean titer of IgG1 remained steady
493 at all time points, while that of IgG2a decreased slightly at week 104, indicating a shift toward a
494 Th2-biased immune response (**Fig. 7F**), consistent with the findings in **Fig. 5A and B**. Therefore,
495 a high dose of antigen induced a higher IgG1 titer and a significant IgG2a titer earlier than in the
496 low-dose group. In the case of IgG2a, the irradiated groups induced earlier responses than the
497 non-irradiated groups in both high- and low-dose groups, although there were no significant

498 differences among the groups. Thus, these results suggest that Th2-prevalent responses were
499 observed at all time points based on the ratio of IgG1 to IgG2a antibody subclasses, and there
500 were no significant differences due to vaccine dose or gamma irradiation.

501

502 *3.6. Non-irradiated and irradiated MAP-S1RS09 induce virus neutralizing antibodies*

503 We measured the neutralization capacity of antibodies induced by our vaccine against
504 SARS-CoV-2 Wuhan, Delta, and Omicron variants using a microneutralization assay (NT₉₀) (**Fig.**
505 **8**). Robust titers of neutralizing antibodies were found against the Wuhan and Delta variant, with
506 the highest levels in the 15 μg and 45 μg dose groups, showing a statistical difference observed
507 only against the Delta variant. Additionally, the 15 μg and 45 μg dose groups were the only ones
508 to show any detectable neutralization activity against the Omicron variant (**Fig. 8**). Notably, the
509 immunogenicity of gamma irradiation-sterilized MAP-S1RS09 vaccines was comparable to that
510 of non-irradiated MAP vaccines in the 15 μg and 45 μg dose groups against the Wuhan and Delta
511 variant, and in the 45 μg dose group against Omicron. However, sera from one to three out of five
512 mice showed neutralization activity in the 5 μg non-irradiated group against the Wuhan and Delta
513 variant, and in the 15 μg irradiated group against Omicron. Altogether, no significant differences
514 were observed between irradiation and non-irradiation in all the groups, thereby supporting the
515 feasibility of gamma irradiation as a terminal sterilization approach for the clinical use of MAP-
516 rS1RS09 vaccines. These findings demonstrate that administering a minimum dose of 15 μg of
517 MAP can generate detectable neutralizing antibodies against various SARS-CoV-2 variants,
518 including Omicron (BA.1).

519

520 *3.7. Long-term Stability of MAP-S1RS09 stored at Elevated Temperature*

521 To assess the stability of irradiated MAP-rS1RS09, patches packed with desiccant were
522 stored either in the refrigerator or at RT for one week, reconstituted in a buffer by shaking for 30

523 minutes, and compared for protein degradation (**Fig 9A**). rS1RS09 from the irradiated patches
524 showed slight smearing in both temperature conditions in the silver-stained gel after SDS-PAGE
525 (**Fig 9B**). To further assess the effects of extended storage time, we stored the patches for
526 nineteen months at either refrigerator or RT, and investigated protein degradation by measuring
527 the relative smear ratio compared to recombinant proteins stored at -20°C for 19 months in
528 parallel. As shown in **Fig. 9C**, there are no differences of rS1RS09 in MAP stored for nineteen
529 months in both temperature conditions compared to rS1RS09 in MAP stored for one week (**Fig**
530 **9B**) or rS1RS09 stored at -20°C for nineteen months, suggesting excellent stability
531 (**Supplementary Fig. 1A**). The mean percentage of protein degradation based on relative
532 intensity (RI) of non-irradiated and irradiated patches using ImageJ analysis was 0.3% and 1.95%
533 when stored at 4°C for nineteen months, and 1.85% and 3.25% when stored at RT, respectively
534 (**Fig 9D**). The degree of protein degradation in irradiated MAPs was comparable to that in non-
535 irradiated MAPs, with no significant differences (1.40% vs. 1.55%), indicating notable stability at
536 RT as good as at 4°C . Furthermore, to assess the stability of long-term storage, the packed MAPs
537 stored at 4°C for nineteen months were further stored under three different conditions: at 4°C in
538 a refrigerator, at RT on a lab bench, and at 42°C in a temperature-controlled incubator for one
539 month (**Fig. 9A**). There were no significant differences of rS1RS09 in MAPs depending on
540 temperature conditions compared to non-irradiated MAPs stored at 4°C for nineteen months (**Fig.**
541 **9E and Supplementary Fig. 1B**). The percentages of protein degradation in both non-irradiated
542 and irradiated MAPs were on average $0.64\% \pm 0.64$ at 4°C , $1.60\% \pm 0.39$ at RT, and $2.62\% \pm 0.03$
543 at 42°C , respectively. In contrast, storage of rS1RS09 in buffer solution at 42°C for one month
544 led to loss of $24.28 \pm 0.66\%$ of proteins, whereas rS1RS09 in MAP showed a loss of $2.80 \pm 0.15\%$,
545 based on silver-staining followed by ImageJ analysis (**Fig. 9F and 9G**). The findings revealed that
546 the MAP platform storage of rS1RS09 is stable for at least twenty months without refrigeration,

547 indicating the temperature stability of our gamma irradiation sterilized MAP-rS1RS09 vaccines
548 without dependence on the cold chain.

549

550 **4. Discussion**

551 In our prior research, we established that a SARS-CoV-2 S1 subunit MAP vaccine
552 effectively triggered the S1-specific immune system in mice [37] and the second and third boosts
553 elicited robust antibody responses against S1, effectively neutralizing the virus, resulting in
554 enduring high-level, long-lasting antibody responses [25].

555 In this study, we found that the SARS-CoV-2 S1 monomer in the presence of RS09 (rS1RS09) is
556 more immunogenic than the trimeric S1 subunit protein, regardless of RS09, considering it as an
557 ideal vaccine candidate. rS1RS09 induced a high level of IgG titer, capable of binding spike
558 proteins expressed on the cell surface, mimicking natural viral trimeric spike proteins.
559 Furthermore, rS1RS09 induced not only a greater IgG titer but also neutralizing antibodies, as
560 determined by the ACE2 binding inhibition assay (**Fig. 1-3**). RS09, TLR4 agonist peptide motif,
561 was identified as an adjuvant using phage display combinatorial peptide technology, which
562 functionally mimicked lipid polysaccharide (LPS), activated NF- κ B signaling, and induced
563 inflammatory cytokine secretion [38]. Interestingly, low (0.1 and 1 μ g) and high dose (10 and 100
564 μ g) of inhaled LPS in a mouse model of asthma induced Th2 and Th1 responses to allergens,
565 respectively [39,40]. Low-dose LPS induced TNF- α secretion, which is a key upstream molecule
566 of Th2 polarization, while high-dose LPS induced IFN- γ , which is a key role in up-regulating
567 allergen-specific IgG2a, thus restoring Th1 response. Similarly, our present study found that
568 rS1RS09 induced a more toward to Th2-biased immune response compared S1 alone, especially
569 at week 18 post-prime, as evidenced by the increase in IgG1 level (**Fig.2C and 2E**). This
570 phenomenon might be attributed to the low dose of RS09, approximately 0.39 μ g RS09 from the
571 45 μ g of rS1RS09. In a past study study, RS09 reported firstly as an adjuvant was injected at 25

572 μg [38], and 10 μg of RS09 adjuvant with a tetravalent DENV nanoparticle vaccine stimulated
573 high IgG2a titers in BALB/c mice [41]. Likewise, antigen containing triple RS09 motif at the C-
574 terminus can impart TLR4 agonist activity, which is approximately 10.85 μg from the 100 μg of
575 the artificial antigen [42]. This suggests that further evaluation of the molecular context of the
576 TLR4 agonistic motif in rS1RS09 should be required using HEK-Blue cells, derived from HEK293
577 cells carrying human TLR4/MD2/CD14 gene and a SEAP (secreted embryonic alkaline
578 phosphatase gene) reporter construct inducible by NF- κ B signaling [42].

579 Furthermore, the present data showed that the monomeric form of S1 induced a higher
580 IgG titer compared to the trimeric form regardless of the RS09 motif (**Fig. 2B and 2F**). This might
581 be explained by many epitopes being hidden in the trimeric form compared to the monomeric
582 form, similar to HIV-1, where its most conserved epitopes are concealed inside the Env core and
583 are only exposed after CD4 receptor engagement [43]. Indeed, the epitope sites are not hidden
584 in the isolated RBD construct, but in the Spike trimer, and the RBDs must move from a down- to
585 an up- position, and antibody binding is further hindered spatially by the N-terminal domain (NTD)
586 and the S2 subunit [44].

587 We also compared the antibody responses induced by MAP of the S1 subunit antigen
588 immunization to those elicited by conventional intramuscular (IM) injections. Despite of no
589 significant difference between the two groups with a low dose of 5 μg rS1(Wu+Beta), the delivery
590 of protein subunit vaccine though MAP is superior to conventional intramuscular injection,
591 although the actual amount of delivered antigen is lower, with an average of 70% [25] (**Fig.4**).
592 Vaccine-induced protection against SARS-CoV-2 diminishes over time, leading to instances of
593 SARS-CoV-2 reinfection [45–47]. Consequently, the challenge of sustaining long-term immunity
594 is the major obstacles in enhancing and approving COVID-19 vaccines [48]. While it's complex to
595 directly compare human and mouse lifespans, the enduring immune responses observed in this
596 study following additional MAP boosts suggest a potential solution to overcome these hurdles.
597 Interestingly, all groups showed similar Th2-prevalent responses regardless vaccine dose and

598 route of immunization (**Fig. 4F**). In contrast, with a high dose of 45 μg rS1RS09, all vaccinated
599 groups had significantly high and similar geometric mean S1 IgG endpoint titer (EPT) after boost
600 when compared to week 0. The IgG geometric mean of the IM group peaked at week 5 and waned
601 the immune response through week 71, while that of the MAP-rS1RS09 group reached a steady
602 geometric mean S1 IgG EPT through week 71 as high as the peak at week 6, showing a significant
603 difference between two groups (**Fig. 5**). The IgG1 and IgG2a EPT of the MAP group were
604 statistically significant compared to pre-immunized controls and showed similar Th2-biased
605 immune response except later time point, at week 70, leaning more toward to Th2-biased in the
606 MAP group. These results are in line with those of an inactivated influenza vaccine encapsulated in
607 MAP, which induced a much higher IgG1 titer and lower IgG2a titer compared to the intramuscular
608 injection of a liquid subunit vaccine [49].

609 Nevertheless, our research revealed that rS1RS09 encapsulated in MAP induced a
610 superior immune response compared to conventional IM injection (**Fig. 4 and 5**). A previous study
611 had shown that gamma irradiation has been employed as a terminal sterilization method for MAP-
612 SARS-CoV-2 subunit vaccines [37]. As part of our efforts toward clinical-grade manufacturing of
613 MAP subunit vaccines, further investigation of dose sparing and gamma irradiation (25 kGy)
614 sterilization was performed. As shown in Fig 6, 15 and 45 μg of MAP-rS1RS09 groups induced
615 and sustained significant IgG titer even at week 71 post-prime vaccination when compared to pre-
616 immunized sera. The immunogenicity of gamma irradiation sterilized MAP vaccines was
617 comparable to that of non-irradiated MAP vaccines in all vaccine groups (**Fig. 6C**). Thereby, these
618 findings support the feasibility of gamma irradiation as a terminal sterilization approach for the
619 clinical usage of our MAP-rS1RS09 vaccines.

620 We also examined the S1-specific IgG1 and IgG2a titers to determine if there were
621 significant differences in the ratio of these subtypes due to vaccine dose and gamma irradiation
622 by comparison with 45 μg and 5 μg groups. Critically, a 45 μg dose of antigen induced a higher
623 IgG1 titer and a significant IgG2a titer earlier than in a 5 μg dose group, resulting in a forward

624 Th2-biased response (**Fig.7**). Furthermore, the irradiated groups induced earlier responses than
625 non-irradiated groups in both high and low dose groups, although there were no significant
626 differences among the groups, consistent with previous reports that Malaria SPf66 synthetic
627 peptide release rate from poly (lactic-co-glycolic acid) (PLGA) microspheres was slightly faster
628 after gamma-irradiation, although no apparent effect on SPf66 integrity and formulation
629 properties, such as morphology, size, and loading [50]. Subcutaneous administration of irradiated
630 and non-irradiated microspheres into mice induced a similar immune response (IgG, IgG1, IgG2a
631 levels) [50]

632 Notably, gamma irradiation-sterilized MAP vaccines induced comparable neutralizing
633 antibodies against three SARS-CoV-2 variants to that of non-irradiated MAP vaccines, with no
634 statistical significance among groups, conforming that the immunogenic activity of the antigen is
635 maintained after gamma-irradiation (**Fig. 8**). Critically, sera from one and three out of 5 mice
636 showed neutralization activity in the non-irradiated 5 μ g group against the Wuhan and Delta
637 variants, respectively, whereas the irradiated 5 μ g group was under the limit of detection. In
638 contrast, sera from two out of 5 mice showed neutralization activity in the irradiated 15 μ g group
639 against Omicron variant, whereas the non-irradiated 15 μ g group was under the limit of detection,
640 indicating few differences in antigenicity by gamma-irradiation. Several previous studies have
641 reported that irradiation would cause breakage and polymerization of peptide chains, and
642 glycosylation modification of proteins [51,52]. Irradiation treatment effects on the protein structure
643 and digestion characteristics of seed-watermelon (*Citrullus lanatus* var.) kernel protein depending
644 on doses of irradiation. After the irradiation treatment, the kernel protein was unfolded by
645 decrease in alpha-helix and beta-sheet structure, and an increase in coil structure, resulting in
646 the exposure of hydrophobic groups [53]. Moreover, gamma irradiation induces protein changes
647 that enhance immunogenicity for snake venoms, *Coccidian* parasites, and *Toxoplasma gondii*
648 protein extracts [54,55]. Therefore, these findings support that the little change in antigenicity by
649 gamma irradiation as a terminal sterilization approach may be beneficial for neutralizing a broad

650 spectrum of emerging variants. Under detectable neutralizing activity in low-dose MAP vaccine
651 can be overcome through additional boost or combinations with adequate adjuvants such as
652 Poly(I:C), monophosphoryl lipid A, cholera toxin, zymosan [15,56–58].

653 We also demonstrated that the MAP platform storage of rS1RS09 is stable for at least
654 twenty months without refrigeration, indicating the temperature stability of our gamma irradiation-
655 sterilized MAP-rS1RS09 vaccines without dependence on the cold chain (**Fig. 9**). Furthermore,
656 an additional one-month storage at high temperature showed that superior benefits of the MAP
657 platform compared to the solution (**Fig. 9F and 9G**). Similarly, recent studies have shown that
658 MAP for administration of inactivated influenza or inactivated polio vaccine improved thermal
659 stability compared with conventional liquid vaccines, which could enable the distribution of
660 vaccines with less reliance on cold chain storage [49,59]. Moreover, researchers found that
661 hyaluronic acid (HA) plays an antioxidant role not only in patches but also in the solution during
662 the manufacturing process and prevents reductions in AA2G content by irradiation [60]. Furthering
663 this, gamma irradiation affected the mechanical properties and architecture of the needles of
664 carboxymethyl cellulose (CMC), but not HA-based patch [61]. They also suggested that e-beam
665 (40 kGy), which provides lower penetration and higher doses into sterilizing materials, is suitable
666 for the terminal sterilization of HA-based DMN patches, because gamma-ray (20 and 30 kGy)
667 irradiation significantly degraded the encapsulated AA2G, while e-beam maintained AA2G activity
668 [60]. However, AA2G is a vitamin C derivative, which is expected to be easily oxidized by the
669 radicals generated from irradiation. From the perspective of immunogenicity, gamma irradiation
670 of antigens can mimic the promotion of slow release by generating smaller solubility protein
671 aggregates like an adjuvant or oxidizing the antigens, as similarly reported for neutrophil
672 myeloperoxidase in acute inflammation [62–64]. Gamma irradiation-inactivated (RI) Respiratory
673 Syncytial Virus (RSV) vaccine exacerbates pulmonary inflammation by switching from prefusion
674 to postfusion F protein. RI-RSV caused more severe ERD than did formalin-inactivated (FI)-RSV,
675 and these immune responses are likely due to pre-F to post-F conformational changes by

676 radiation-induced redox oxygen species (ROS) [65]. Moreover, gamma irradiation of antigens
677 also induced a better cellular immune response [66]. In addition, irradiated proteins undergo one
678 of the chemical changes, such as cross-linking, which can support the chemical bonding of HA to
679 antigens, stimulating strong and enduring humoral responses like an adjuvant [67]. Indeed, in this
680 study, the GMT of IgG in all irradiated groups was observed to be slightly higher than those
681 induced in the non-irradiated groups at most time points (**Fig.6B**).

682 This study had two limitations, which will be addressed in future research. These include
683 conducting a SARS-CoV-2 challenge study to evaluate the protective effectiveness of non-
684 irradiated and irradiated MAP vaccination and assessing the immunogenicity of non-irradiated
685 and irradiated MAP-S1RS09Delta stored for the long-term at different temperatures compared to
686 those stored for the short-term. Interestingly, a study has reported that IgG titers were similar in
687 mice vaccinated with MAP encapsuled inactivated influenza vaccine stored for one month at 4°C,
688 25°C, and 45°C compared to MAP vaccine stored for one day at 4°C, leading to complete protection
689 against the virus challenge with no significant weight loss [49].

690 In summary, our research revealed that delivery with MAP platforms for subunit vaccine
691 is superior to conventional IM injection in terms of immunogenicity and long-term stability. We
692 investigated the impact of gamma irradiation as a terminal sterilization method for MAP platforms
693 targeting SARS-CoV-2 S1 antigens, resulting in the development of comparable antibody
694 responses. These findings suggest the potential for further research into utilizing irradiated MAP
695 to deliver S1 subunit vaccines for emerging SARS-CoV-2 variants. Such vaccines could serve as
696 safe boosters to enhance cross-neutralizing antibody responses. This approach holds promise
697 as a viable platform for facilitating widespread global vaccination efforts.

698

699 **5. Conclusions**

700 This study presents the first assessment of long-term immunogenicity and stability of
701 gamma-irradiation-sterilized dissolving microarray patches (MAP) for the SARS-CoV-2 subunit

702 vaccine. rS1RS09 was identified as an ideal immunogen after investigating the immunogenicity
703 of four recombinant Delta S1 proteins (rS1, rS1RS09, rS1f, and rS1fRS09) in monomeric or
704 trimeric forms in BALB/c mice. Moreover, MAP delivery of rS1RS09 elicited strong and sustained
705 antibody responses, outperforming conventional intramuscular injections. Both irradiated and
706 non-irradiated MAP vaccines showed comparable immunogenicity, demonstrating a robust, long-
707 lasting immunity and a dose-sparing effect. The enhanced stability of the subunit vaccine in MAPs
708 during extended storage outside the cold chain supports their potential for global distribution and
709 holds promise as an effective, stable, and scalable SARS-CoV-2 vaccine candidate. Overall,
710 gamma irradiation as a terminal sterilization method is feasible for mass production and future
711 pandemic preparedness.

712

713 **Declaration of competing interest** the authors declare that they have competing interests
714 regarding the research presented in this manuscript. Specifically, AG and EK are co-founders of
715 GAPHAS PHARMACEUTICAL INC., a private startup company that could potentially benefit from
716 the findings of this research. AG, EK, and MSK have equity in GAPHAS PHARMACEUTICAL
717 INC. However, the authors have taken measures to ensure that the research is conducted
718 objectively and that the data and conclusions presented in this manuscript are not influenced by
719 their competing interests. The study was designed, conducted, and analyzed independently of
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721

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1001
1002

1003 **FIGURE LEGENDS**

1004 **Fig. 1.** Construction and expression of recombinant SARS-CoV-2-S1Delta proteins. (A) A shuttle
1005 vector carrying the codon-optimized four constructs of SARS-CoV-2-S1Delta gene encoding N-
1006 terminal 1-661 with c-tag (EPEA, glutamic acid-proline-glutamic acid-alanine) was designated as
1007 shown in the diagram. Amino acid changes in the SARS-CoV-2-S1 region of in this study are
1008 shown. ITR: inverted terminal repeat; CMVp, cytomegalovirus promoter; RBD: receptor binding
1009 domain. (B) Purified proteins, rS1RS09 (lane1), rS1f (lane2), and rS1fRS09 (lane3), isolated by
1010 c-tag affinity purification were separated by SDS-PAGE and visualized by silver staining.
1011 Molecular weight marker (MW marker) is indicated on the left. (C) Detection of the Purified SARS-
1012 CoV-2-S1 proteins, rS1RS09 (lane1), rS1f (lane2), and rS1fRS09 (lane3), respectively, by
1013 western blot using rabbit anti spike of SARS-CoV Wuhan polyclonal antibody.

1014

1015 **Fig. 2.** Comparison in mouse immunized with four SARS-CoV-2 rS1Delta protein subunit
1016 vaccines. (A) Schedule of immunization and blood sampling for IgG end point titration. Balb/c
1017 mice (5 per group) were immunized with 45 μ g of rS1, rS1RS09, rS1f, and rS1fRS09 proteins of
1018 SARS-CoV-2 Delta then administered intramuscularly at week 0 and 3. Syringes indicated the
1019 timing of immunizations, and the red drops denote times at which blood was drawn. (B) Sera were
1020 diluted, and SARS-CoV-2-S1-specific antibodies were quantified by ELISA to determine the IgG
1021 endpoint titer. The IgG titers at each time points were showed in each mouse. The bars represent
1022 geometric mean with geometric SD. (C and D) Sera at weeks 0, 5, and 18 were diluted, and
1023 SARS-CoV-2-S1WU-specific IgG1 (C) an IgG2a (D) were quantified by ELISA to determine each
1024 IgG subclasses endpoint titer. The titers at each time points were showed for each mouse. The
1025 bars represent geometric mean. (E) S1-specific IgG1/IgG2a ratios of individual mice at weeks 7
1026 and 18 as mean values with SEM. (F) Flow cytometry assay of Expi293 cells expressing S1Delta-
1027 S2WU at the cell surface. At 36 hrs post-transfection with pAd/ S1Delta-S2WU, binding to SARS-
1028 CoV-2-S at the Expi293 cell surface was analyzed by incubation with mice sera obtained at week

1029 7 after immunization followed by staining with FITC-conjugated anti-mouse IgG. Groups were
1030 compared by the Kruskal-Wallis test at each time point, followed by Dunn's multiple comparisons.
1031 Significant differences are indicated by * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, n.s., not significant.

1032

1033 **Fig. 3.** Percent ACE binding inhibition of neutralizing antibodies against SARS-CoV-2 variants.
1034 Antibodies in sera capable of neutralizing the interaction between SARS-CoV-2 Wuhan, Alpha
1035 (B.1.1.7), Beta (B.1.351), Gamma (P.1), Delta (B.1.617.2), Zeta (P.2), Kappa (B.1.617.1), New
1036 York (B.1.516.1), India (B.1.617 and B.1.617.3) variants spike and ACE2 were examined at week
1037 0 (peach), and at week 7 post-prime in all animals immunized with rS1 (green), rS1RS09 (blue),
1038 rS1f (pink), and rS1fRS09 (purple) of SARS-CoV-2 Delta. Serum samples were diluted in 1:20
1039 before adding the V-PLEX plates. Box and whisker plots represent the median and upper and
1040 lower quartile (box) with min and max (whiskers). Groups were compared by Kruskal-Wallis test
1041 at each variant, followed by Dunn's multiple comparisons. Significant differences are indicated by
1042 *; $p < 0.05$ and **; $p < 0.01$. Asterisks represent statistical differences compared with pre-
1043 immunized sera (W0).

1044

1045 **Fig. 4.** Immune responses induced by different vaccine routes of administration in ICR mice. (A)
1046 Immunogens and delivery routes of each group (B) Experimental schedule representing the
1047 immunization timeline. ICR mice (5 per group) were prime and boosted three weeks intervals with
1048 either 5 μ g of rS1RS09Delta intramuscularly (IM) or MAP-rS1RS09Delta intradermally (ID). The
1049 red drops represented bleeding. (C) Reciprocal serum endpoint dilutions of SARS-CoV-2-S1-
1050 specific antibodies were measured by ELISA to determine the IgG endpoint titers at weeks 0, 3,
1051 6, and 9. Sera at weeks 0 and 6 were diluted, and SARS-CoV-2-S1WU-specific IgG1 (D) an IgG2a
1052 (E) were quantified by ELISA to determine each IgG subclasses endpoint titer. The bars represent
1053 geometric mean. (F) S1-specific IgG1/IgG2a ratios of individual mice at weeks 6 as mean values
1054 with SEM. IM and MAP groups were compared for statistically significant differences using

1055 Kruskal-Wallis test at each variant, followed by Dunn's multiple comparisons. *: p value,0.05; **: p,0.01.

1057

1058 **Fig. 5. Immune responses induced by different vaccine routes of administration of high**
1059 **dose.** (A) Silver stained SDS-PAGE of purified SARS-CoV-2 Delta rS1RS09 protein (B)
1060 Fabrication of dissolving microneedle patch, MAP-rS1RS09 was composed of 35 microneedles.
1061 (C) Each MN was approximately 700 μm in length, occupying a 2.36 cm^2 area size of patch, 7 \times 5
1062 arrays of 700 μm -long microneedles with 45 μg of rS1RS09Delta (D) Experimental schedule
1063 representing the immunization timeline. Balb/c mice (5 per group) were prime and boosted three
1064 weeks intervals with either 45 μg of rS1RS09Delta intramuscularly or MAP-rS1RS09Delta
1065 intracutaneously. The red drops represented bleeding. (E) Reciprocal serum endpoint dilutions of
1066 SARS-CoV-2-S1-specific antibodies were measured by ELISA to determine the IgG endpoint
1067 titers until week 90. Sera at weeks 0, 5, and 18 were diluted, and SARS-CoV-2-S1WU-specific
1068 IgG1 (F) an IgG2a (G) were quantified by ELISA to determine each IgG subclasses endpoint titer.
1069 The titers at each time points were showed for each mouse. The bars represent geometric mean.
1070 (H) S1-specific IgG1/IgG2a ratios of individual mice at weeks 7 and 18 as mean values with SEM.
1071 IM and MAP groups were compared for statistically significant differences using non-parametric
1072 Mann-Whitney-U-test compared with pre-immunized sera. *p < 0.05, **p < 0.01, ***p < 0.001, n.s.,
1073 not significant.

1074

1075 **Fig. 6. Dose sparing and gamma irradiation responses.** (A) Fabrication of dissolving
1076 microneedle patch, MAP-rS1RS09 was composed of 35 microneedles. (B) Experimental
1077 schedule representing the immunization timeline. Balb/c mice (5 per group) were prime and
1078 boosted three weeks intervals with either non-irradiated (-) or irradiated (+) of 5, 15, 45 μg of MAP-
1079 rS1RS09Delta. The red drops represented bleeding. (C) Reciprocal serum endpoint dilutions of
1080 SARS-CoV-2-S1-specific antibodies were measured by ELISA to determine the IgG endpoint

1081 titers at weeks 0, 3, 6, 16, 28, 52, 71, 90, and 104. (D) The SARS-CoV-2-S1-specific IgG titers at
1082 all time points until week 104 post-prime (BALB/c (N = 5) except at week 90 (N = 2-3) and at week
1083 104 (N = 1-3)).

1084

1085 **Fig. 7. IgG subclasses.** Sera were collected from the mice immunized 5 μ g of MAP-rS1RS09
1086 with and without irradiation (A, B, and C) or 45ug of MAP-rS1RS09 with and without irradiation
1087 (D, E, and F) at weeks 0, 9, 28, 52, 71, 90 and 104. SARS-CoV-2-S1WU-specific IgG1 (A and D)
1088 an IgG2a (B and E) were quantified by ELISA to determine each IgG subclasses endpoint titer.
1089 The titers at each time points were showed for each mouse. The bars represent geometric mean.
1090 (C and F) S1-specific IgG1/IgG2a ratios of individual mice at each time points as mean values
1091 with SEM. Groups were compared by the Kruskal-Wallis test at all time points, followed by Dunn's
1092 multiple comparisons. Significant differences are indicated by * $p < 0.05$, ** $p < 0.01$.

1093

1094 **Fig.8. Neutralization of SARS-CoV-2 variants.** Serum from mice immunized with SARSCoV-2
1095 S1 via MAP intradermal delivery was assessed using a microneutralization assay (VNT₉₀) for
1096 neutralization against SARS-CoV-2 variants Wuhan, Delta variant (B.1.617.2), and Omicron
1097 variant (BA.1). Serum titers that resulted in 90% reduction in cytopathic effect compared to the
1098 virus control were reported. Data are representative of the geometric mean with error bars
1099 representing geometric standard deviation. Each group was compared to the pre-immunized sera
1100 using the Mann-Whitney test. Significant differences are indicated by ** $p < 0.01$, * $p < 0.05$.

1101

1102 **Fig. 9. Stability persistence of irradiated MAP** (A) Experimental schedule representing storage
1103 temperature and the reconstruction timeline of MAP. Silver staining of the recombinant proteins
1104 reconstructed from the non-irradiated (–) and irradiated (+) MAP stored at either 4°C or RT for 1
1105 week (B) and 19 months (C). (D) Percentage of protein degradation in the MAP from silver staining
1106 in Fig. 9C, measured by ImageJ. The recombinant protein stored at – 20°C for 19 months was

1107 set to 0%. (E) Percentage of protein degradation for an additional one month at different
1108 temperatures with the MAP stored for 19 months at 4°C, measured by ImageJ. The recombinant
1109 protein stored at –20°C for 19 months was set to 0%. (F) Silver staining of the recombinant
1110 proteins (rP) stored at –20°C for 20 months (lane 1), in PBS (lane 2) or in the preservative buffer
1111 (lane 3) stored at 42°C for 1 month after 19 months at –20°C, and rP reconstituted from the non-
1112 irradiated (–) MAP stored at 42°C for 1 month after 19 months at 4°C (lane 4). (G) Percentage of
1113 protein degradation of rS1RS09 in the buffer solution or MAP from silver staining in Fig. 9F,
1114 measured by ImageJ.

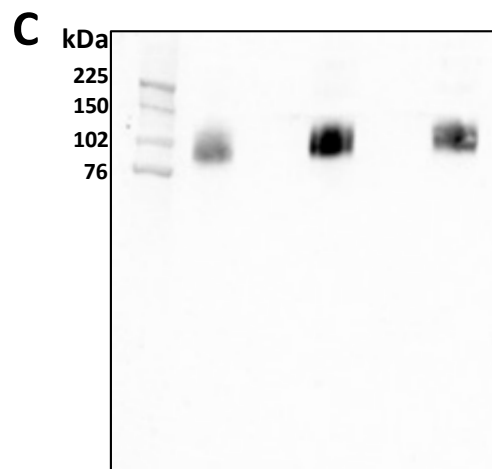
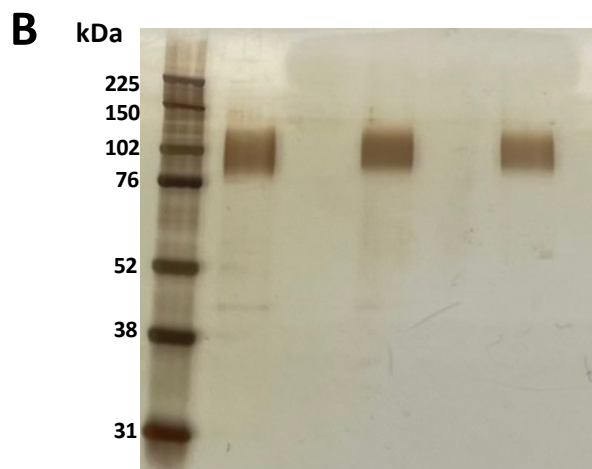
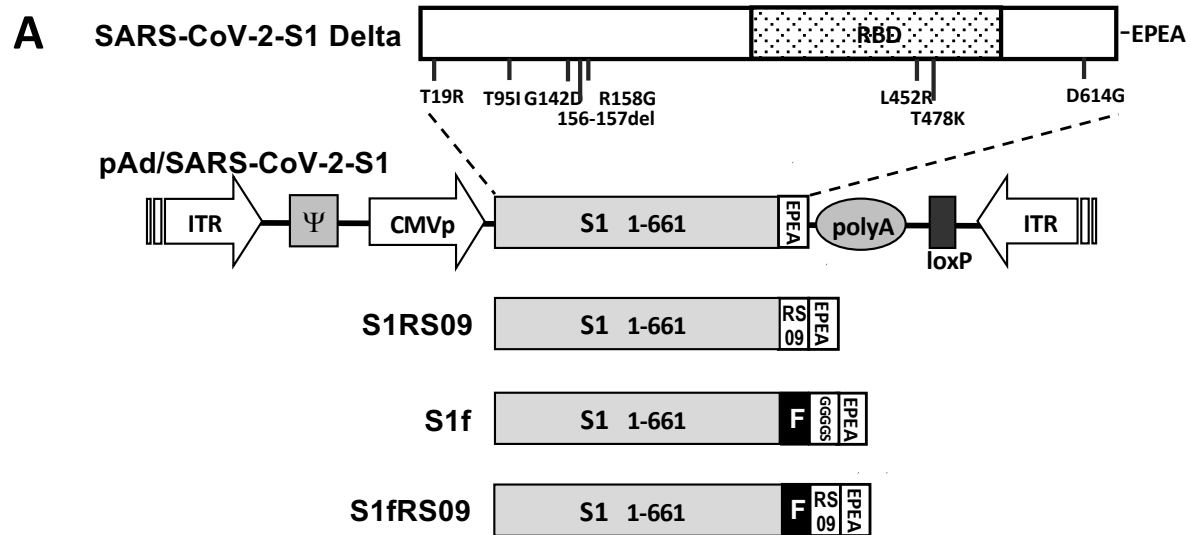


Fig. 1

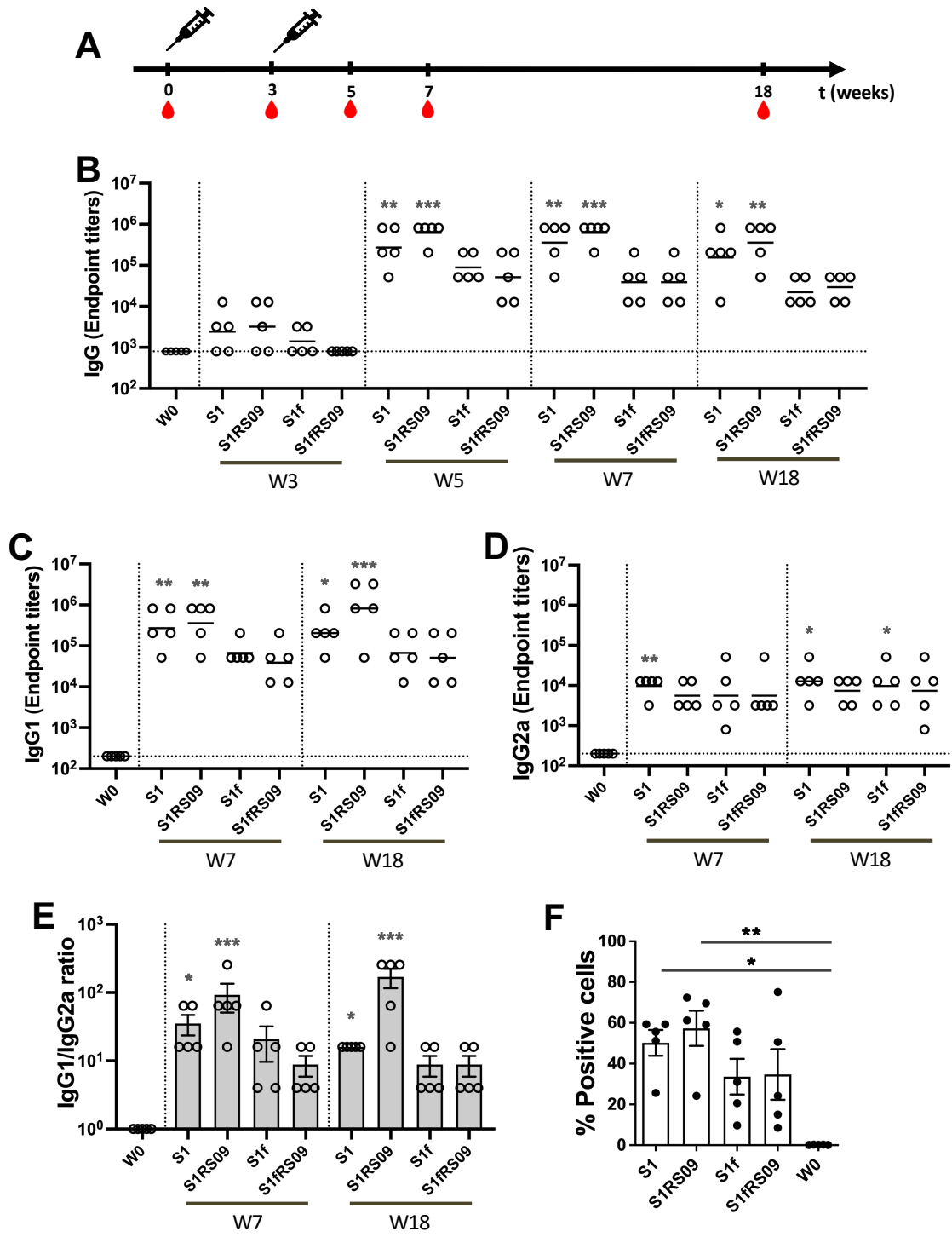


Fig. 2

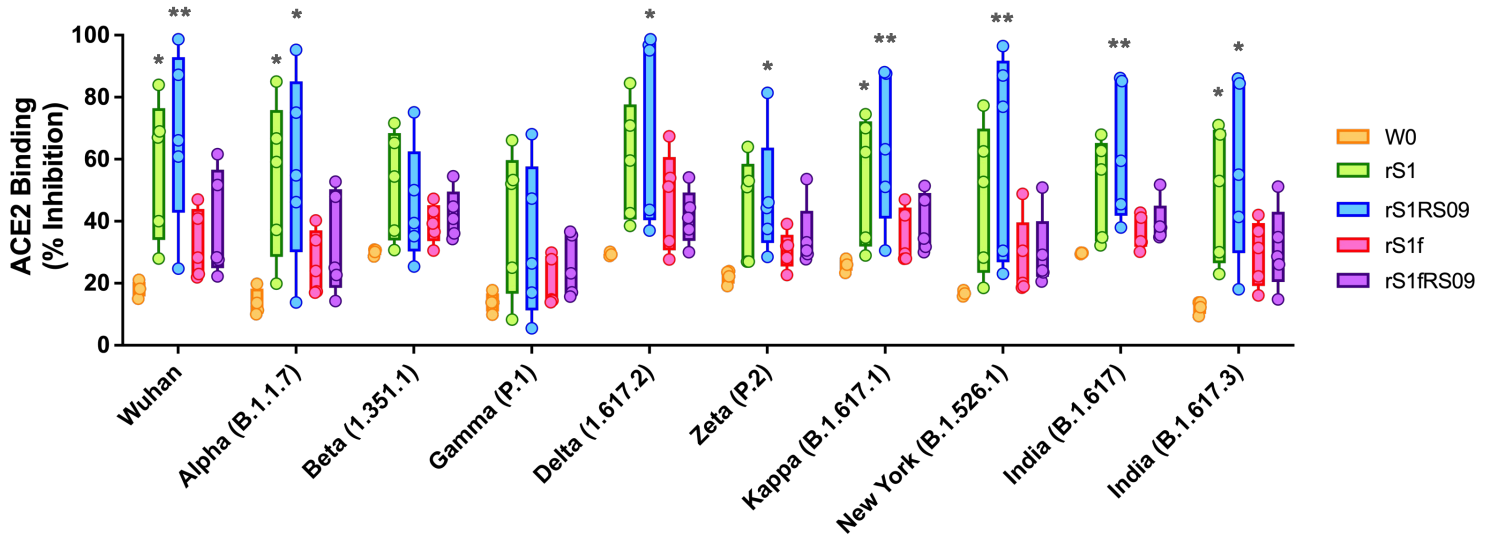


Fig. 3

A

	Prime (W0)	Boost (W3)	No. of Array	Patch size	Dose	Route
G1	rS1(WU+Beta)	rS1(WU+Beta)	N/A	N/A	5 μ g	IM
G2	MAP-S1(WU+Beta)	MAP-S1(WU+Beta)	35	2 cm X 1.18 cm (2.36 cm ²)	5 μ g	ID
G3	MAP-S1(WU+Beta)	MAP-S1(WU+Beta)	25	1.5 cm diameter (1.7 cm ²)	7 μ g	ID

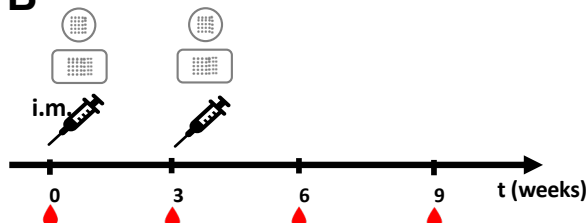
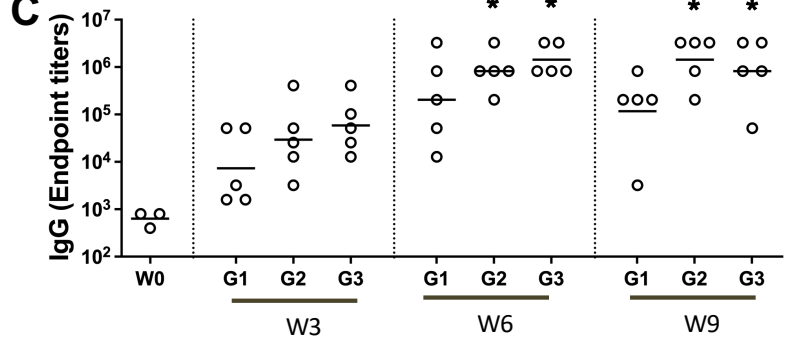
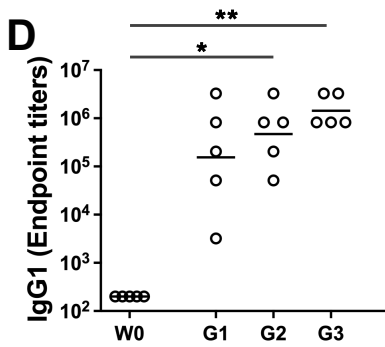
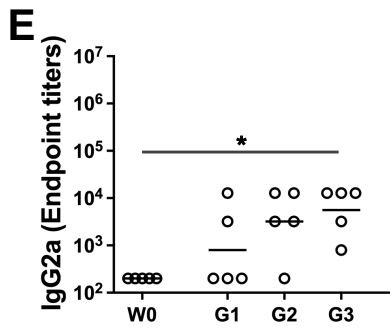
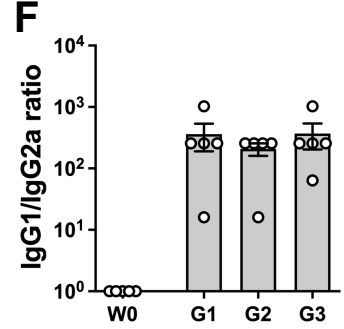
B**C****D****E****F**

Fig. 4

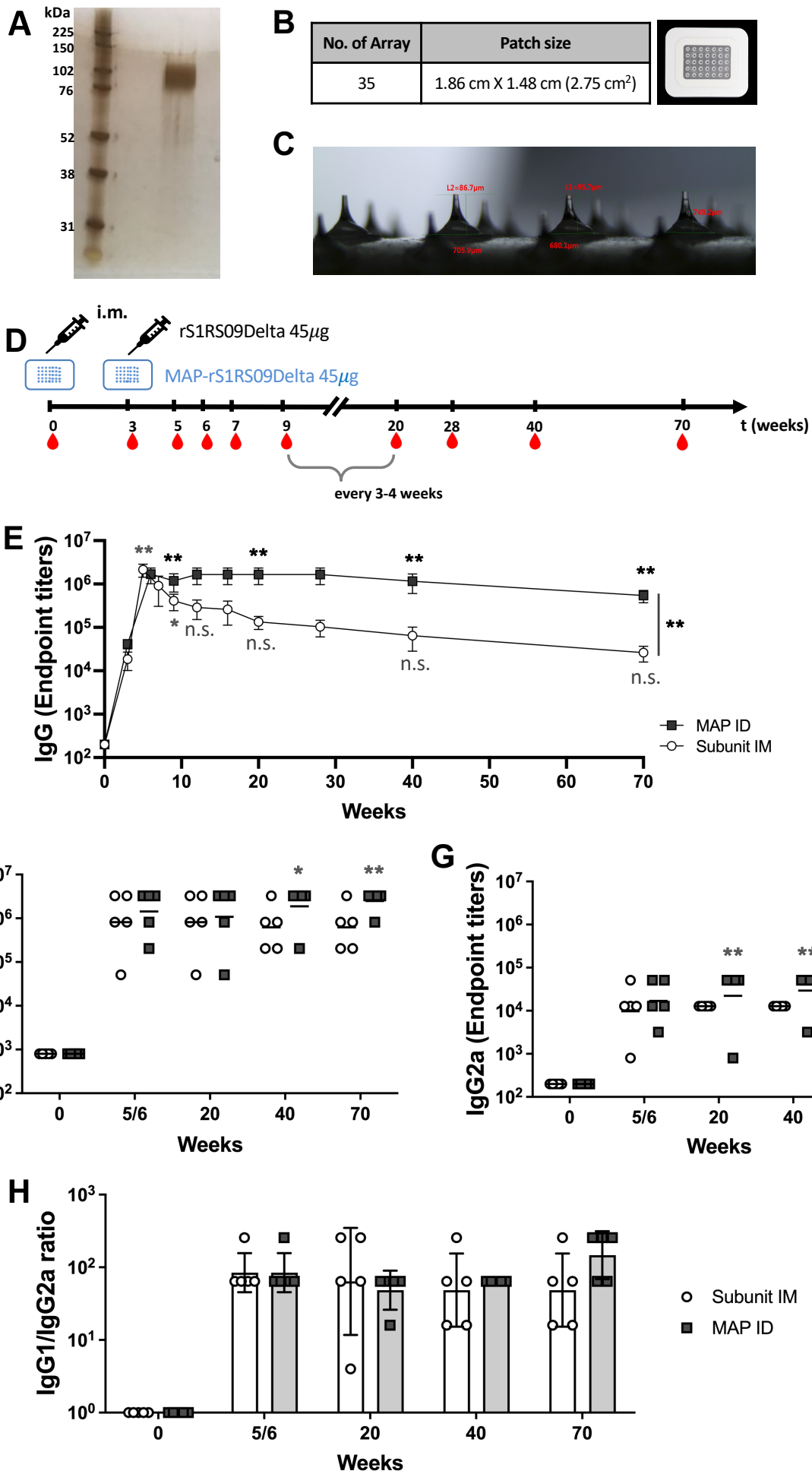


Fig. 5

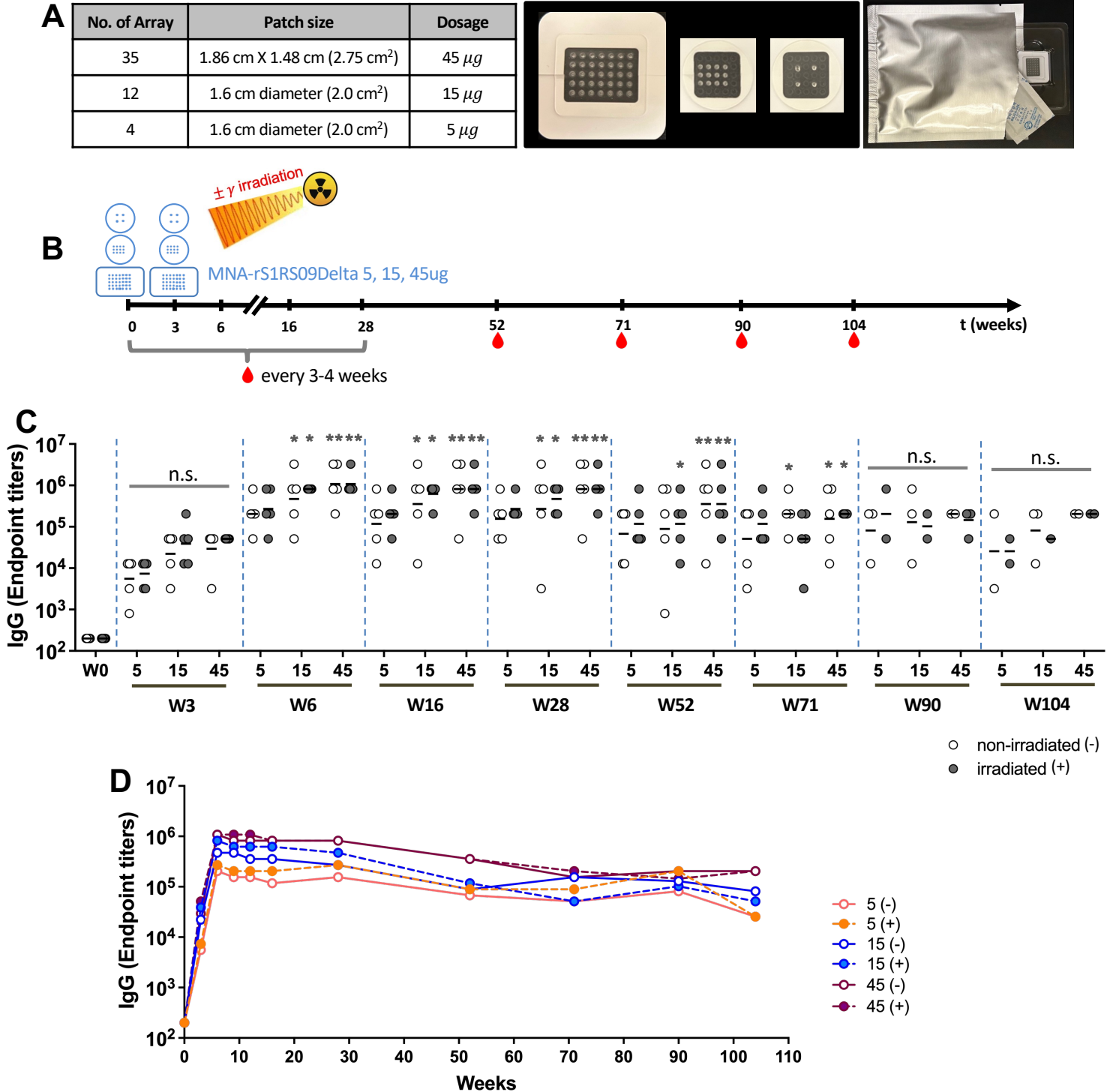


Fig. 6

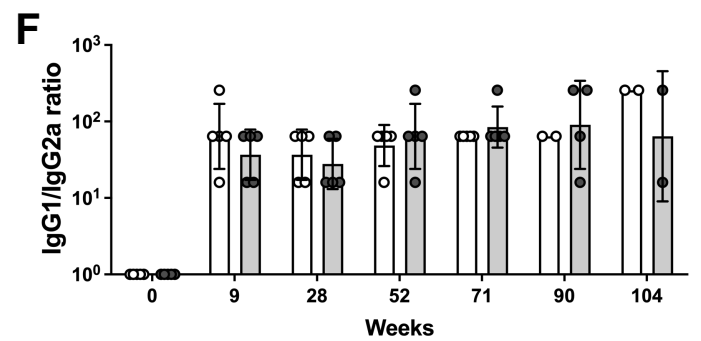
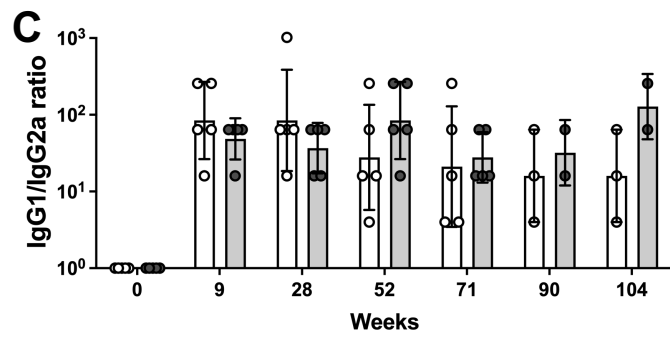
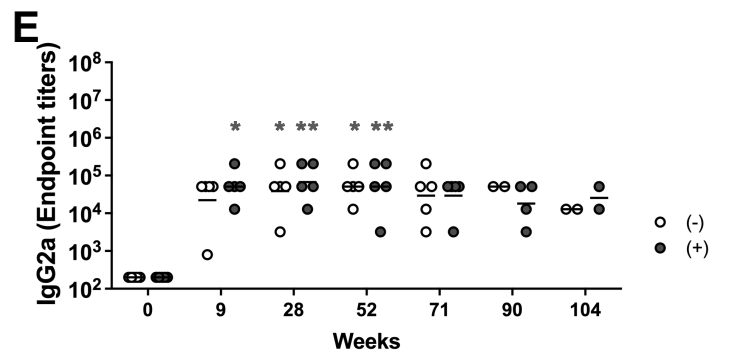
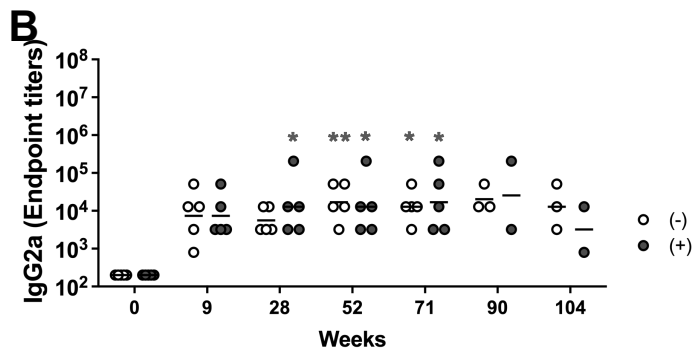
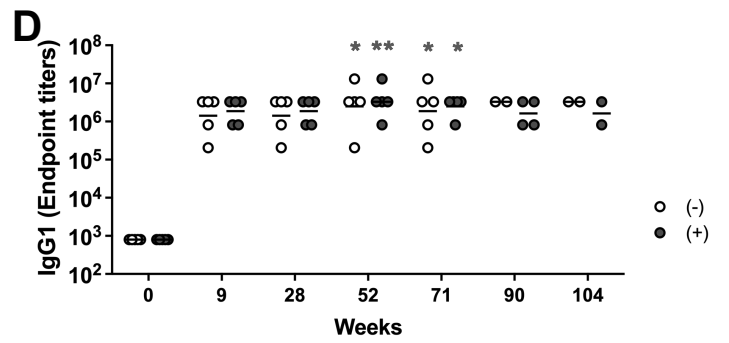
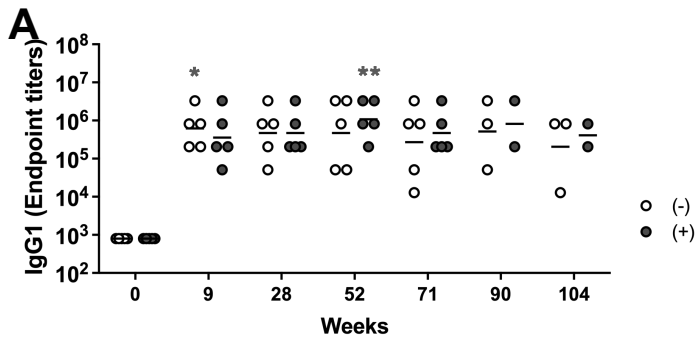


Fig. 7

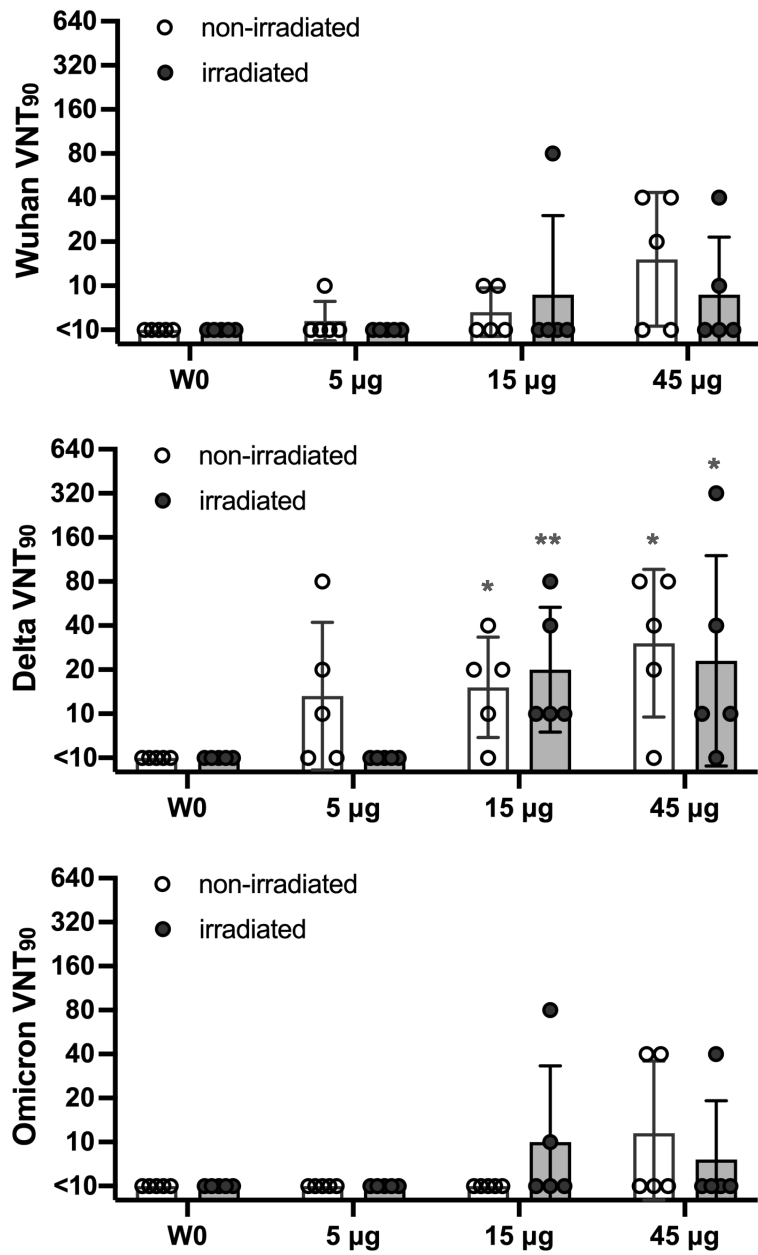


Fig. 8

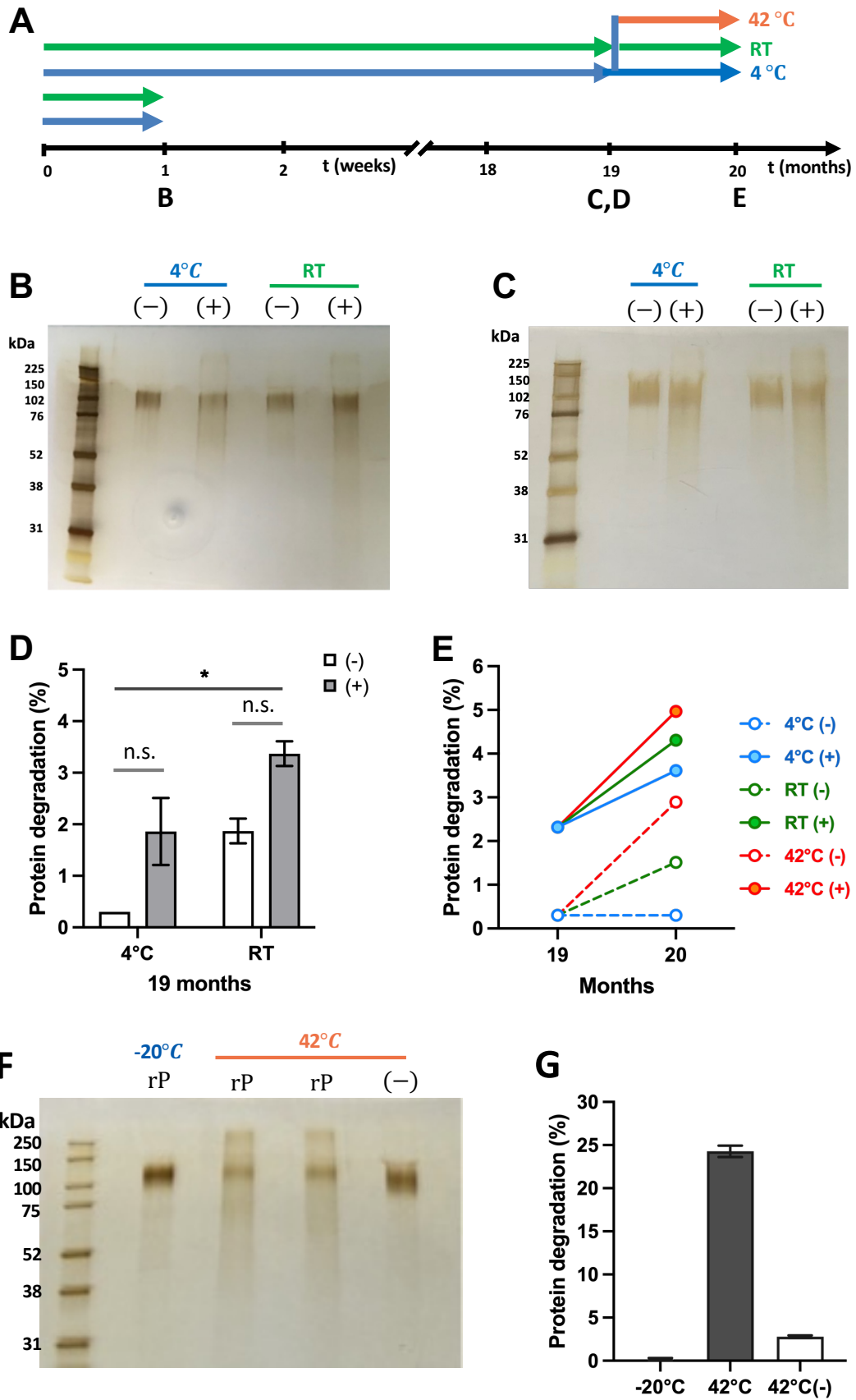


Fig. 9