

1 **A COVID-19 vaccine candidate using SpyCatcher 2 multimerization of the SARS-CoV-2 spike protein 3 receptor-binding domain induces potent neutralising 4 antibody responses**

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48

49 ABSTRACT

50 There is dire need for an effective and affordable vaccine against SARS-CoV-2 to
51 tackle the ongoing pandemic. In this study, we describe a modular virus-like particle
52 vaccine candidate displaying the SARS-CoV-2 spike glycoprotein receptor-binding
53 domain (RBD) using SpyTag/SpyCatcher technology (RBD-SpyVLP). Low doses of
54 RBD-SpyVLP in a prime-boost regimen induced a strong neutralising antibody
55 response in mice and pigs that was superior to convalescent human sera. We
56 evaluated antibody quality using ACE2 blocking and neutralisation of cell infection by
57 pseudovirus or wild-type SARS-CoV-2. Using competition assays with a monoclonal
58 antibody panel, we showed that RBD-SpyVLP induced a polyclonal antibody response
59 that recognised all key epitopes on the RBD, reducing the likelihood of selecting
60 neutralisation-escape mutants. The induction of potent and polyclonal antibody
61 responses by RBD-SpyVLP provides strong potential to address clinical and logistic
62 challenges of the COVID-19 pandemic. Moreover, RBD-SpyVLP is highly resilient,
63 thermostable and can be lyophilised without losing immunogenicity, to facilitate global
64 distribution and reduce cold-chain dependence.

65 **Keywords**

66 SARS-CoV-2; mAb; SpyTag; VLP; immunity; pre-clinical; coronavirus

67

68 **INTRODUCTION**

69 Coronavirus disease 2019 (COVID-19), caused by a novel coronavirus named severe
70 acute respiratory syndrome coronavirus 2 (SARS-CoV-2), was first reported in Wuhan,
71 China in December 2019 ¹. Since then COVID-19 has spread across the world and
72 was declared a pandemic by the World Health Organisation (WHO) in March 2020. As
73 of August 2020, there have been over 20 million confirmed COVID-19 cases worldwide
74 and around 800,000 deaths ². There are no vaccines or effective treatments for
75 COVID-19 to date; however, as of August 2020, there are 25 vaccine candidates in
76 clinical evaluation and around 140 are in pre-clinical testing ³. Vaccine candidates in
77 current clinical evaluation include inactivated, viral vector (replicating and non-
78 replicating), protein subunit, nucleic acid (DNA and RNA) and virus-like particle (VLP)
79 vaccines with the majority of them focusing on using the full-length SARS-CoV-2 spike
80 glycoprotein (S) as an immunogen.

81

82 SARS-CoV-2 is an enveloped virus carrying a single-stranded positive-sense RNA
83 genome (~30 kb), belonging to the genus *Betacoronavirus* from the *Coronaviridae*
84 family ⁴. The virus RNA encodes four structural proteins including spike (S), envelope
85 (E), membrane (M), and nucleocapsid (N) proteins, 16 non-structural proteins, and
86 nine accessory proteins ⁵. The S glycoprotein consists of an ectodomain (that can be
87 processed into S1 and S2 subunits), a transmembrane domain, and an intracellular
88 domain ⁶. Similar to the Severe Acute Respiratory Syndrome Coronavirus (SARS-
89 CoV), SARS-CoV-2 binds the human angiotensin-converting enzyme 2 (ACE2) via the
90 receptor-binding domain (RBD) within the S1 subunit to facilitate entry into host cells,
91 followed by membrane fusion mediated by the S2 subunit ⁷⁻⁹

92

93 Of the many vaccine platforms, protein subunit vaccines generally have good safety
94 profiles and their production is rapid and easily scalable ¹⁰. Recombinant RBD proteins
95 of SARS-CoV and MERS-CoV have been shown to be immunogenic and induce
96 protective neutralising antibodies in animal models and are therefore considered
97 promising vaccine candidates (reviewed ^{11, 12}). RBD from SARS-CoV- has recently

98 been confirmed to be inducing neutralising antibodies^{13, 14}. Recently published studies,
99 including one from our group, found that the majority of the potent neutralising
100 antibodies isolated from SARS-CoV-2- infected patients bound to the RBD^{15, 16}. We
101 therefore chose to study the immunogenicity of RBD. To improve immunogenicity, we
102 conjugated the RBD onto a virus-like particle (VLP). VLP display of protein antigen
103 has been shown to further enhance immunogenicity by facilitating antigen drainage to
104 lymph nodes, enhancing uptake by antigen-presenting cells and increasing B cell
105 receptor crosslinking^{10, 17}. Moreover, we recently showed that influenza antigens
106 (haemagglutinin (HA) or neuraminidase (NA)) displayed on the VLPs (same VLP used
107 in this study) were highly immunogenic at a low dose (0.1 µg) in mice¹⁸.

108

109 In the present study, we used the SpyTag/SpyCatcher technology for assembly of
110 SARS-CoV-2 RBD on the mi3 VLP, via the formation of an intermolecular isopeptide
111 bond between the RBD and the VLP, as a potential SARS-CoV-2 vaccine^{19, 20} (Figure
112 1A). SpyTag-mediated VLP decoration has been successfully used for the display of
113 diverse antigens from, e.g. *Plasmodium spp.*, influenza A virus, HIV and cancer cells
114 (PD-1L)^{18, 19, 21, 22}. The RBD-SpyVLP vaccine candidate is highly immunogenic in mice
115 and pigs, inducing robust SARS-CoV-2 neutralising antibody responses. The results
116 of our study demonstrate the potential of the RBD-SpyVLPs as an effective and
117 affordable vaccine for COVID-19 with RBD-SpyVLP being resilient, retaining stability
118 and immunogenicity post-lyophilisation, which will greatly facilitate distribution for
119 vaccination by eliminating cold-chain dependence.

120

121 RESULTS

122 1. RBD can be efficiently displayed on the mi3 VLP via SpyTag/SpyCatcher

123 To create the RBD-SpyVLP vaccine candidate, the SpyTag (AHIVMVDAYKPTK)
124 coding sequence was fused between the signal sequence from influenza H7 HA and
125 the N-terminus of the RBD (amino acid 340-538, NITN...GPKK)
126 (A/HongKong/125/2017) (SpyTag-RBD) (see Figure S1 for the full sequence) and the
127 glycoprotein was expressed in mammalian cells (Expi293) prior to purification using
128 Spy&Go affinity chromatography²³. The purified SpyTag-RBD was then conjugated
129 to the SpyCatcher003-mi3 VLP^{18, 20} to generate the SpyTag-RBD:SpyCatcher003-
130 mi3 (RBD-SpyVLP) immunogen (Figure 1A). SpyCatcher003 is a variant of
131 SpyCatcher which was engineered for accelerated reaction with SpyTag²⁴. mi3 is a

132 computationally engineered dodecahedron based on an aldolase from a thermophilic
133 bacterium^{20,25}. The SpyTag-RBD can be efficiently conjugated to the SpyCatcher-mi3
134 VLP, with 93% display efficiency reached after 16 h (Figure 1B). This corresponds to
135 an average of 56 RBDs per VLP. We saw no sign of VLP aggregation following
136 coupling and the RBD-SpyVLP is homogeneous, as shown by a uniform peak of the
137 hydrodynamic radius (R_H) at 20.7 ± 4.2 nm in dynamic light scattering (DLS) (Figure
138 1C). For immunisation, we chose a conjugation ratio that leaves minimal free RBD
139 (1:1 molar ratio), which corresponds to ~64% display efficiency or around 38 RBD per
140 VLP (Figure 1B).

141

142 **2. RBD-SpyVLP is reactive to monoclonal antibodies isolated from recovered
143 patients and is resilient**

144 To confirm the antigenicity of RBD-SpyVLP, we performed a series of binding assays.
145 Binding to RBD-SpyVLP was tested using a panel of novel monoclonal antibodies
146 (mAbs) some of which are strongly neutralising, from COVID-19-infected donors²⁶,
147 that bind to at least three independent epitopes on the RBD. We included the
148 published conformation-specific mAb (CR3022)²⁷, a nanobody-Fc fusion VHH72-Fc
149²⁸, and a dimeric human ACE2-Fc²⁹, for which there are published structures. All
150 tested mAbs and ACE2-Fc bound strongly to the RBD-SpyVLP (Figure 2A), showing
151 that a broad range of epitopes on the RBD-SpyVLP are exposed and correctly folded.
152 An anti-influenza neuraminidase mAb (Flu mAb), used as a negative control, showed
153 no binding to RBD-SpyVLP, confirming the specificity of the assay (Figure 2A).

154

155 We then tested the stability of RBD-SpyVLP, to determine its resilience and likely
156 sensitivity to failures in the cold-chain³⁰. The unconjugated SpyCatcher003-mi3 VLP
157 had previously been shown to be highly thermostable as a platform for antigen display
158¹⁸. For conjugated RBD-SpyVLP, we tested its solubility following storage for two
159 weeks at -80, -20, 4 or 25 °C in Tris Buffered Saline (TBS). We then centrifuged out
160 any aggregates and analysed soluble protein by SDS-PAGE with Coomassie staining.
161 We found no significant change in the soluble fraction following storage at 4 °C (n=3,
162 Kruskal-Wallis and Dunn's post-hoc test, p>0.05), with only a 12% decrease after
163 storage for two weeks at 25 °C (Figure 2B) and no degradation was observed at 25 °C
164 (Figure S2A). We further analysed the integrity of the sample with ELISA against the
165 conformation-dependent CR3022 mAb and observed no loss of antigenicity under

166 these storage conditions (Figure 2C). We next assessed the resilience of RBD-
167 SpyVLP to freezing, challenging RBD-SpyVLP with multiple rounds of freeze-thaw.
168 Even after five rounds of freeze-thaw, there was no significant loss of soluble RBD-
169 SpyVLP (Figure 2D) or CR3022 recognition (Figure 2E) (n=3, Kruskal-Wallis and
170 Dunn's post-hoc test, p>0.05) and no degradation was observed (Figure S2B). After
171 reconstitution following lyophilisation, we saw a minimal change in soluble protein for
172 RBD-SpyVLP (91.5±3.8% of the initial value (mean±SD) (Figure 2F), which was not
173 statistically significant (n=3, Mann-Whitney U test, p>0.05). There was also no
174 difference in terms of binding of RBD-SpyVLP to a panel of mAbs or ACE2-Fc
175 recognising non-overlapping footprints on the RBD (Figure 2G). Overall, RBD-SpyVLP
176 showed a high level of resilience.

177

178 **3. RBD-SpyVLP induces a strong ACE2-blocking and neutralising antibody
179 response in mouse models**

180 We first evaluated the immunogenicity of RBD-SpyVLP in mouse models. C57BL/6
181 mice (n=6) were immunised intramuscularly (IM) with purified RBD alone (0.1 µg or
182 0.5 µg), RBD-SpyVLP (0.1 µg or 0.5 µg equivalents of the RBD component) or VLP
183 alone, all adjuvanted with AddaVax. AddaVax is a squalene-based oil-in-water nano-
184 emulsion adjuvant, a pre-clinical equivalent to the licensed MF59 adjuvant³¹. Mice
185 were then boosted with the same dose of immunogen two weeks later and sera were
186 collected at three weeks post-boost. Both the 0.1 µg and 0.5 µg RBD-only groups
187 showed levels of antibody against RBD or spike glycoprotein only slightly above
188 background, detected using ELISA (serum reciprocal endpoint titre (EPT): 1:94 and
189 1:68, respectively), and showed no difference compared to the VLP-only group (Figure
190 3A and B). Mice immunised with 0.1 µg and 0.5 µg RBD-SpyVLP groups showed high
191 levels of antibody to RBD (EPT: 0.1 µg: 1:16,117, p<0.001 and 0.5 µg: 1:7,300, p<0.01)
192 (Figure 3A) and to spike (EPT: 0.1 µg: 1:1,647, p<0.05 and 0.5 µg: 1:1,212) compared
193 to the VLP group (Figure 3B).

194

195 We then tested RBD-SpyVLP in a second mouse strain (BALB/c) with the same
196 dosage regimen to confirm the immunogenicity (Figure 3A and B). In BALB/c, the 0.1
197 µg and 0.5 µg RBD-SpyVLP groups showed higher levels of RBD-specific antibody
198 (EPT: 0.1 µg: 1:8,406, p<0.01 and 0.5 µg: 1:16,636, p<0.05) (Figure 3A) and spike-

199 glycoprotein specific antibody (EPT: 0.1 µg: 1:9,574, p<0.001 and 0.5 µg: 1:18,556,
200 p<0.05) (Figure 3B) compared to the VLP group.

201

202 The ability of immunised mouse sera to block recombinant soluble ACE2 binding to
203 immobilised RBD was then assessed. Sera from the 0.1 µg and 0.5 µg RBD-SpyVLP
204 immunised C57BL/6 mice had significantly higher ACE2 blocking (IC₅₀: 0.1 µg: 1:132,
205 p<0.05 and 0.5 µg: 1:253; p<0.05) activity compared to the VLP-immunised mice
206 (Figure 3C). C57BL/6 mice immunised with RBD-only showed no detectable ACE2
207 blocking activity, consistent with the ELISA results (Figure 3A & B). Similarly, both 0.1
208 µg and 0.5 µg RBD-SpyVLP-immunised BALB/c mice had significantly higher ACE2
209 blocking (IC₅₀: 0.1 µg: 1:200, p<0.001 and 0.5 µg: 1:560, respectively) compared to the
210 VLP immunised group (Figure 3C). In both mouse strains, 0.1 µg and 0.5 µg RBD-
211 SpyVLP immunised groups had higher antibody titres against RBD (around 5-10-fold;
212 C57BL/6, p<0.001 and p<0.0001; BALB/c, p<0.001 and p<0.0001) and ACE2 blocking
213 (around 10-50-fold; C57BL/6, p<0.0001 and p<0.0001; BALB/c, p<0.0001 and
214 p<0.0001) compared to plasma donated by patients convalescing from COVID-19
215 disease (n=28) (Figure 3B and C). Sera from BALB/c mice immunised with 0.1 µg and
216 0.5 µg RBD-SpyVLP had comparable spike glycoprotein-specific antibody responses
217 compared to convalescent humans whereas sera from C57BL/6 mice immunised with
218 either dose had higher spike glycoprotein-specific antibody responses compared to
219 convalescent humans (around 2-fold, 0.1 µg, p<0.05 and 0.5 µg, p<0.001) (Figure 3B).

220

221 The antibody response in mice was assessed for neutralisation potency using a live
222 SARS-CoV-2 virus (hCoV-19/England/02/2020, EPI_ISL407073) neutralisation assay
223 (VNT) based on virus plaque reduction. Sera from C57BL/6 mice immunised with
224 either dose of unconjugated RBD showed low level neutralising titres (ND₅₀ 1:59 and
225 1:32) compared to the RBD-SpyVLP group, again consistent with ELISA and ACE2
226 blocking activity (Figure 3D). Both 0.1 µg and 0.5 µg RBD-SpyVLP immunised groups
227 exhibited neutralising titres in both C57BL/6 (ND₅₀: 1:450 to 1: 2,095) and BALB/c
228 mice (ND₅₀: 1:230 to 1:1,405) (Figure 3D). Consistent neutralising activity in sera from
229 C57BL/6 mice was found using a VNT in an independent laboratory with the hCoV-
230 19/VIC01/2020 isolate (GenBank MT007544) (Table S1).

231

232 We tested the immunogenicity of RBD-SpyVLP post-lyophilisation and after five
233 freeze-thaw cycles. C57BL/6 mice (n=6) were immunised IM with 0.5 µg RBD-SpyVLP
234 (pre-lyophilised, post-lyophilised or 5× freeze-thaw) and sera were harvested post
235 prime and post boost and tested for binding in a RBD ELISA. There was no difference
236 between the immune responses in all three groups tested (Figure 3E) (Kruskal-Wallis,
237 followed by Dunn's post-hoc test, p>0.05), showing that lyophilisation and freeze-thaw
238 had not compromised the immunogenicity of the RBD-SpyVLP.

239

240 **4. RBD-SpyVLP is highly immunogenic and induces strong neutralising
241 antibody response in pigs**

242 We tested RBD-SpyVLP for its immunogenicity in pigs as a large, genetically outbred
243 animal model. Pigs have previously been reported to be a reliable model to study
244 vaccines for use in humans because of their highly similar physiologies and immune
245 systems^{32, 33}. The pig model has been used recently to test an adenovirus vector
246 vaccine candidate against SARS-CoV-2 (ChAdOx1 nCoV19)³⁴. We immunised pigs
247 (n=3) with a dose of RBD-SpyVLP that we intend to use in humans (5 µg) or with a
248 10-fold higher dose (50 µg) to study dose-response. A third group (n=3) receiving 100
249 µg of purified trimeric spike glycoprotein (spike) was included as a control. Pigs were
250 immunised IM twice, 28-days apart, with 5 µg or 50 µg RBD-SpyVLP or spike, all
251 adjuvanted with AddaVax.

252

253 As early as day 7, pigs immunised with 50 µg RBD-SpyVLP showed a detectable anti-
254 RBD antibody response, whereas no antibody response was detected at day 7 for 5
255 µg RBD-SpyVLP or 100 µg spike groups (Figure 4A) (p<0.05, Kruskal-Wallis test).
256 The antibody response in all three groups increased gradually to day 14 when the
257 response reached a plateau. The group receiving 50 µg RBD-SpyVLP group showed
258 a trend of slightly higher antibody response than the other two groups until the day of
259 boost (day 28) (Figure 4A). The antibody response in all three groups increased by
260 around 100-fold one week after boosting and remained high until at least day 56. There
261 was no dose-response difference between the 5 µg and 50 µg RBD-SpyVLP groups
262 (p>0.5, Kruskal-Wallis test) for day 56 data (Figure 4A).

263

264 ACE2 blocking activity in the pig serum was measured at day 21, day 35 and day 56.
265 No ACE2 blocking was detected at day 21, prior to the booster dose. ACE2 blocking

266 was detected post-boost at day 35 (IC_{50} 1:10 to 1:50) and reduced at day 56 (IC_{50} 1:5
267 to 1:25) in all three groups (Figure 4B). ACE2 blocking activities in all three groups
268 were significantly higher than for sera from convalescent humans (5 μ g RBD-SpyVLP,
269 $p<0.01$; 50 μ g RBD-SpyVLP, $p<0.001$; 100 μ g, $p<0.05$), by around ~3-6-fold ($p<0.05$
270 and $p<0.001$, respectively, Kruskal-Wallis test on day 56 data).

271

272 Neutralising antibody responses in the pigs were assessed against both SARS-CoV-
273 2 pseudotyped lentivirus and live SARS-CoV-2 virus. All three groups exhibited similar
274 neutralisation titres against pseudovirus (ND_{50} : ~1:500 to 1:4,000) after boost which
275 remained until at least day 56, with no difference between the three groups ($p>0.05$,
276 Kruskal-Wallis test on day 56) (Figure 4C). Similarly, no difference in neutralising
277 activity was detected in all three groups against live virus at day 21 (Figure 4D).
278 Neutralisation was detected on both day 35 and 56 in all groups, with ND_{50} levels
279 between 1:6,000 to 1:11,000 on day 56 and no significant difference between the three
280 groups ($p>0.05$, Kruskal-Wallis test on day 56 data) (Figure 4D). Nasal and oral
281 secretions were also tested for the presence of neutralising antibodies: here RBD-
282 specific IgG, but notably not IgA, was detected in all three groups with no significant
283 difference between groups post-boost for both nasal and oral secretions ($p>0.05$,
284 Kruskal-Wallis followed by Dunn's Post-hoc) (Figure 4E & F).

285

286 We assessed the level of RBD-specific B cells in peripheral blood. The predominance
287 of an IgG response following the booster immunisation was confirmed in all three
288 groups by assessment of RBD tetramer labelling of IgM, IgG and IgA B cells in
289 peripheral blood and IgG ELISpot assay (Figure S3A & B). In each case, RBD-specific
290 IgG B cells peaked in peripheral blood shortly after the boost (Figure S3A). We also
291 performed longitudinal analysis of CD4 $^{+}$ and CD8 $^{+}$ T cell responses following
292 immunisation. Intracellular cytokine staining of S-peptide-stimulated peripheral blood
293 mononuclear cells (PBMCs) demonstrated a T cell IFN- γ response, slightly larger in
294 the CD4 $^{+}$ than CD8 $^{+}$ T cell pool, which peaked 7 days after prime and boost
295 immunisations (Figure S3C).

296

297

298 **5. RBD-SpyVLP induces polyclonal antibody responses against the RBD in
299 mice and pigs**

300 A concern regarding RBD-based vaccines is whether the immune response will be
301 focussed to a single site on the antigen, because of the relatively small size of the RBD,
302 potentially leading to a narrow response that would be sensitive to immune escape^{35, 36}.
303 A recent report showed that passive immunisation with a single mAb led to escape
304 mutants of SARS-CoV-2, whereas a cocktail of two neutralising antibodies to independent
305 epitopes prevented emergence of neutralisation-escape mutations, demonstrating the
306 importance of a polyclonal antibody response³⁷. We assessed sera from mice and pigs
307 immunised with RBD-SpyVLP for antibody responses that target multiple RBD epitopes
308 using a competition ELISA against four different mAbs: FI-3A, FD-11A, EY6A and S309,
309 that target three non-overlapping epitopes on the RBD with FD-11A and S309 showing
310 overlap as defined by competition ELISA (see diagram in Figure 2A)^{26, 38, 39}. BALB/c mice
311 immunised with RBD-SpyVLP showed competition against all four mAbs tested (Figure
312 5A). BALB/c mice immunised with VLP-only showed no competition ($p<0.05$, Mann
313 Whitney U-Test) (Figure 5A). A similar pattern was observed in the C57BL/6 mice (Figure
314 S5). Comparison of preimmune and day 42 post- RBD-SpyVLP-immunisation sera from
315 pigs showed a trend of partial competition against all four mAb, but this was not
316 statistically significant because of inter-animal variation (Figure 5B). When responses of
317 individual pigs were compared to their pre-immune sera 2 out of 3 animals in each dosing
318 group showed significant competition for the antibodies FI-3A and S309 that defined
319 independent neutralising epitopes compared to their preimmune sera (Figure S5). These
320 results show that RBD-SpyVLP does not have an immunodominant epitope and does not
321 induce a highly focussed antibody response, making it a vaccine candidate that is likely
322 to resist the generation of neutralisation-escape mutants.

323

324

325 **DISCUSSION**

326 In the current study we investigated an RBD-based VLP vaccine candidate for COVID-
327 19 based on SpyTag/SpyCatcher technology which was used to assemble RBDs into
328 the mi3 VLP via the formation of an irreversible isopeptide bond¹⁹. We showed RBD-
329 SpyVLPs to be strongly immunogenic in mice and pigs, inducing high titre neutralising
330 antibody responses against wild type SARS-CoV-2 virus. This study confirms that the
331 RBD is the key immunogenic domain for eliciting neutralising monoclonal antibodies
332 against SARS-CoV-2, in line with studies showing that highly neutralising antibodies
333 isolated from convalescent patients bind to the RBD^{16, 26, 38, 40-46}. We showed that

334 RBD-SpyVLPs are recognised by a panel of mAbs isolated from convalescent patients
335 ²⁶ binding to various epitopes on the RBD (Figure 2A). This distributed reactivity shows
336 that all of the epitopes that could potentially induce protective antibodies to RBD are
337 present in RBD-SpyVLPs.

338

339 We detected negligible antibody responses in mice vaccinated with equivalent doses
340 (0.1 µg or 0.5 µg) of purified RBD alone, but strong responses to the RBD when
341 displayed on the VLP (Figure 3). Previous studies showed that RBD from SARS-CoV
342 and SARS-CoV-2 can induce neutralising antibodies in animal models but typically
343 after administration of much higher doses (e.g. ~50 to 100 µg) and with frequent
344 dosing ^{13, 14, 47}. On the other hand, we showed that high titre neutralising antibody
345 responses can be detected in two strains of mice immunised with relatively low doses
346 of RBD-SpyVLP (to ND₅₀ ~500 to 2,000). These results confirm the enhanced
347 immunogenicity of RBD when displayed on SpyVLPs. Sera from mice immunised with
348 both 0.1 µg or 0.5 µg of RBD-SpyVLP exhibited high levels of antibody against SARS-
349 CoV-2 RBD and full-length spike glycoprotein and ACE2 blocking activity (Figure 3B,
350 C and D). All of these responses were higher than the levels found in plasma from
351 convalescent humans. Together, these observations suggest that RBD-SpyVLP
352 vaccination could potentially elicit protective antibody responses against SARS-CoV-
353 2 in humans.

354

355 RBD-SpyVLP vaccination also induces high titre neutralising antibody responses in
356 pigs (ND₅₀ ~1:11,000) with a dose that we aim to use for subsequent human trials (5
357 µg) (Figure 4D). At a dose around 2-fold less (based on molar ratio), 5 µg of RBD-
358 SpyVLP induced similar neutralisation titres compared to 100 µg of spike glycoprotein,
359 showing the excellent immunogenicity of RBD-SpyVLP vaccination. Transudate RBD-
360 specific IgG from serum can be detected in the oral and nasal cavity (Figure 4E & F).
361 Surprisingly, no increase in antibody titre was observed in pigs that received a higher
362 dose of antigen (50 µg RBD-SpyVLP). There was a trend that the 50 µg RBD-SpyVLP
363 group generated a more rapid and higher response post-prime but the antibody
364 response between the 5 µg and 50 µg RBD-SpyVLP groups were identical post-boost.
365 This suggests a threshold effect.

366

367 Since RBD-SpyVLPs induce antibody responses that target multiple epitopes on the
368 RBD the chance of selecting neutralisation-escape mutants should be greatly reduced.
369 Circulating SARS-CoV-2 stains are constantly mutating and the likelihood of
370 persistence of the virus in the human population is high^{48, 49}.

371

372 We observed differences in the levels of ACE2 blocking in sera from immunised mice
373 and pigs (Figure 3C and 4B). Sera taken from mice immunised with RBD-SpyVLP had
374 at least one order of magnitude higher ACE2 blocking activity compared to serum from
375 pigs, despite neutralisation titres being comparable (Figure 3D and 4D). This suggests
376 that mice and pigs may produce distinct antibody responses against the vaccine
377 candidate, although they were equally potent in neutralising live viruses. Surprisingly,
378 ACE2 blocking in both mild and critical/severe convalescent humans who had natural
379 infection were also low compared to the sera from immunised mice (Figure 3D).
380 Nevertheless, the RBD-SpyVLP induces strong neutralising antibody responses in
381 both mice and pigs. The potential for a vaccine based on the RBD is further
382 emphasised by the sterile immunity induced in non-human primates (*Macaca mulatta*)
383 by two doses of 20 to 40 µg of unconjugated RBD in Al(OH)₃ adjuvant¹⁴, and the
384 successful induction of high titre neutralising antibody responses with an elegant self-
385 assembling RBD-virus like nanoparticle⁵⁰.

386

387 A recently published report on an inactivated SARS-CoV-2 vaccine candidate showed
388 that vaccinated non-human primates (NHP) with a serum neutralising titre (ND₅₀) of
389 <1:100 were still protected against wild type SARS-CoV-2 challenge with no weight
390 loss and no detectable lung pathology⁵¹. In our study, RBD-SpyVLP-vaccinated mice
391 and pigs had ND₅₀ at least an order of magnitude higher than 1:100, which would be
392 expected to provide protection. Sera from pigs immunised with RBD-SpyVLP had
393 similar, if not higher, neutralisation titres against wild type SARS-CoV-2 compared to
394 pigs immunised by adenoviral vector (ChAdOx1 nCoV-19; both vaccines given as 2
395 doses in the same timeframe)³⁴. A single dose of ChAdOx1 nCoV-19 has been shown
396 to be protective against viral pneumonia and lung pathology following SARS-CoV-2
397 challenge (ND₅₀ ~1:20) in NHP but not virus shedding in the nasal cavity⁵². The
398 seemingly similar ND₅₀ of the RBD-SpyVLP compared to the ChAdOx1 nCoV-19 in
399 pigs suggests that the RBD-SpyVLP will be as protective as the ChAdOx1 nCoV-19
400 in NHP and most likely in humans if similar neutralisation titres are achieved. A pre-

401 stabilised spike glycoprotein vaccine candidate, NXV-CoV2273 developed by
402 Novavax, showed neutralising antibody responses in humans that were 4-fold higher
403 than convalescent human sera, when given two doses of 5 µg with Matrix M1 adjuvant
404 ⁵³. In light of the higher efficacy of RBD-SpyVLP compared to spike glycoprotein in
405 pigs (Figure 4), we are hopeful that at least similar efficacy of RBD-SpyVLP could be
406 achieved in humans.

407

408 Recently published studies reveal the relatively short-lived antibody response to
409 SARS-CoV-2 in convalescent patients ^{54, 55}. Further work is required to define the
410 significance of these results for the longevity of protective immunity. It is possible that
411 re-infection with the virus would lead to a strong protective secondary antibody
412 response from memory B cells. Here, we have shown that the antibody response in
413 pigs immunised with RBD-SpyVLP persisted for at least 2 months and remained
414 neutralising. Studies on the longevity of the immune responses are to be undertaken
415 in the near future.

416

417 Apart from being highly immunogenic, SpyVLPs provide a versatile modular vaccine
418 platform to facilitate conjugation with other antigens. Should a mutation in the RBD
419 arise in circulating SARS-CoV-2 strains ⁴⁸, a matched RBD-SpyVLP could be
420 manufactured rapidly. In addition, more than one RBD variant can be co-displayed on
421 the VLP, to provide broader protection against various SARS-CoV-2 strains ⁵⁶. The
422 RBD-SpyVLP can also be co-displayed with antigens from other pathogens such as
423 the HA and NA from influenza virus. We have recently shown HA and NA to be highly
424 immunogenic in mice after formulation as a SpyVLP ¹⁸. This assembly could potentially
425 provide protection against both SARS-CoV-2 and influenza viruses. Testing resilience
426 of the vaccine candidate, we found that RBD-SpyVLP is stable at ambient temperature,
427 resistant to freeze-thaw, and can be lyophilised and reconstituted with minimal loss in
428 activity (Figure 2B-G) or immunogenicity (Figure 3F). This resilience may not only
429 simplify vaccine distribution worldwide, especially to countries where cold-chain
430 resources are incomplete, but also reduce the overall vaccine cost by removing cold-
431 chain dependence. We are currently investigating cheaper and more scalable
432 alternatives to produce RBD-SpyVLP to cope with the global demand for a SARS-
433 CoV-2 vaccine. Collectively, our results show that the RBD-SpyVLP is a potent and
434 adaptable vaccine candidate for SARS-CoV-2.

435

436 **MATERIALS AND METHODS**

437 **Expression constructs**

438 The SpyTag-RBD expression construct (Figure S1) consists of influenza H7
439 haemagglutinin (A/HongKong/125/2017) signal-peptide sequence, SpyTag⁵⁷, (GSG)₃
440 spacer and SARS-CoV-2 spike glycoprotein (GenBank: NC045512) (amino acid 340-
441 538). The insert was ordered from GeneArt and subcloned into pcDNA3.1 expression
442 plasmid using unique *Not*I-*Eco*RI sites to create pcDNA3.1-SpyTag-RBD (GenBank
443 and Addgene deposition in progress) (see Figure S1). pET28a-SpyCatcher003-mi3
444 (GenBank and Addgene deposition in progress) was created by replacing SpyCatcher
445 in pET28a-SpyCatcher-mi3 with SpyCatcher003¹⁸. RBD used in the RBD ELISA was
446 expressed from a codon optimised RBD cDNA subcloned into the vector
447 pOPINTTGNeo incorporating a C-terminal His₆ tag (RBD-6H) as previously described
448³⁴. Human ACE2 fused to human IgG1 Fc domain (ACE2-Fc) used in the ACE2
449 competition ELISA was expressed from codon optimised human ACE2 cDNA (amino
450 acid 18 to 615) fused to the Fc region and a C-terminal His₆ tag subcloned into the
451 vector pOPINTTGNeo.

452

453 **Expression and purification of SpyCatcher003-mi3**

454 SpyCatcher003-mi3 was expressed in *E. coli* BL21(DE3) RIPL cells (Agilent) as
455 previously described¹⁸. Heat-shock transformed cells were then plated on LB-Agar
456 plates (50 µg/mL kanamycin) and incubated for 16 h at 37 °C. A single colony was
457 picked and cultured in 10 mL starter LB culture (50 µg/mL kanamycin) for 16 h at 37 °C
458 and shaking at 200 rpm. The preculture was diluted 1:100 into 1 L LB (50 µg/mL
459 kanamycin and 0.8% (w/v) glucose) and cultured at 37 °C, 200 rpm until OD600 ~0.6.
460 Protein expression was induced with isopropyl β-D-1-thiogalactopyranoside (IPTG)
461 (420 µM) and incubated at 22 °C, 200 rpm for a further 16 h. The culture was
462 centrifuged and the pellet was resuspended in 20 mL 25 mM Tris-HCl, 300 mM NaCl,
463 pH 8.5 with 0.1 mg/mL lysozyme, 1 mg/mL cOmplete mini EDTA-free protease
464 inhibitor (Merck) and 1 mM phenylmethanesulfonyl fluoride (PMSF). Cell suspension
465 was incubated at 22 °C for 30 min on a platform shaker and sonicated on ice 4 times
466 for 60 s at 50% duty-cycle using an Ultrasonic Processor (Cole-Parmer). Cell lysate
467 was clarified at 35,000 g for 45 min at 4 °C. The supernatant was filtered through 0.45

468 μ m and 0.22 μ m syringe filters (Starlab) and 170 mg ammonium sulfate was added
469 per mL of lysate. SpyCatcher003-mi3 particles were precipitated by incubating the
470 lysate at 4 °C for 1 h while mixing at 100 rpm. Precipitated particles were pelleted by
471 centrifugation at 30,000 g for 30 min at 4 °C. The collected pellet was resuspended
472 into 8 mL TBS pH 8.5 (25 mM Tris-HCl, 150 mM NaCl). Residual ammonium sulfate
473 was removed by dialysing for 16 h against 500-fold excess of TBS. Dialysed
474 SpyCatcher003-mi3 was concentrated to 4 mg/mL using a Vivaspin 20 100 kDa spin
475 concentrator (Vivaproducts) and centrifuged at 17,000 g for 30 min at 4 °C to pellet
476 any insoluble material. The supernatant was filtered through a 0.22 μ m syringe filter.
477 The purified SpyCatcher003-mi3 was then further purified using size exclusion
478 chromatography (SEC). In brief, 2.5 mL was loaded into a HiPrep Sephacryl S-400
479 HR 16-600 SEC column (GE Healthcare) equilibrated with TBS using an ÄKTA Pure
480 25 system (GE Healthcare). Proteins were separated at 1 mL/min while collecting 1
481 mL elution fractions. The fractions containing the purified particles were identified by
482 SDS-PAGE, pooled, and concentrated using a Vivaspin 20 100 kDa MW cut-off
483 centrifugal concentrator. Endotoxin was removed from the SpyCatcher003-mi3
484 samples using Triton X-114 phase separation as previously described ¹⁸. The
485 concentration of endotoxin-depleted particles was measured using bicinchoninic acid
486 (BCA) assay (Pierce) and particles were stored at -80 °C.

487

488 **Expression and purification of SpyTag-RBD**

489 SpyTag-RBD was expressed in Expi293F cells using ExpiFectamine293 transfection
490 reagent (Thermo Fisher) according to the manufacturer's protocol. Supernatant was
491 harvested between 5 to 7 days post transfection and filtered through a 0.22 μ m filter,
492 before purifying using Spy&Go affinity purification with minor modifications ^{18,23}. Briefly,
493 filtered supernatants were diluted with one third supernatant volume of TP buffer (25
494 mM orthophosphoric acid adjusted to pH 7.0 at 22 °C with Tris base) and adjusted to
495 pH 7. Spy&Go resin in same buffer was mixed with the diluted supernatant and
496 incubated at 4 °C for 1 h with gentle agitation. The mixture was poured into an Econo-
497 Pak column (Bio-Rad) and allowed to empty by gravity. The resin was washed with 2
498 \times 10 column volumes of TP buffer and SpyTag-RBD was eluted with 2.5 M imidazole
499 in TP buffer adjusted to pH 7.0 at room temperature (RT). One column volume of
500 elution buffer was added to the resin at a time and incubated for 5 min before collecting
501 each fraction. Elution fractions were analysed using SDS-PAGE with Coomassie

502 staining and the fractions containing SpyTag-RBD were pooled and dialysed against
503 10 mM Tris-HCl pH 8.0 with 200 mM NaCl. The sample was concentrated using
504 Vivaspin-20 10 kDa and further purified via SEC using ÄKTA Pure 25 (GE Life
505 Sciences) equipped with Superdex 75pg 16-600 column (GE Life Sciences), run at 1
506 mL/min. The dialysis buffer was used as the mobile phase. The final yield of purified
507 SpyTag-RBD was around 100 mg/L. The heterogeneity in the SpyTag-RBD band on
508 SDS-PAGE is expected from the presence of different glycoforms on the N-linked
509 glycosylation sites in the construct (Figure S1).

510

511 **RBD-SpyVLP conjugation**

512 SpyTag-RBD at 2 μ M, 3 μ M, 4 μ M or 6 μ M were conjugated at 4 °C for 16 h with 2 μ M
513 SpyCatcher003-mi3 (VLP:RBD ratio 1:1, 1:1.5, 1:2 and 1:3) in TBS pH 8.0. Possible
514 aggregates were then removed by centrifugation at 16,900 g for 30 min at 4 °C.
515 Samples of the supernatant were mixed with reducing 6 \times loading dye (0.23 M Tris-
516 HCl, pH 6.8, 24% (v/v) glycerol, 120 μ M bromophenol blue, 0.23 M SDS, 0.2 M
517 dithiothreitol) and resolved on 12%, 14% or 16% SDS-PAGE using XCell SureLock
518 system (Thermo Fisher). Gels were then stained with InstantBlue Coomassie
519 (Expedion) and imaged using ChemiDoc XRS imager (Bio-Rad). The intensities of
520 bands on each lane were quantified using ImageLab (version 5.2) software (Bio-Rad)
521 and Fiji distribution of ImageJ (version 1.51n). Conjugation efficiency (as % of
522 unconjugated SpyVLP left) was calculated as (band density of unconjugated SpyVLP
523 left in the conjugation reaction/band density of SpyVLP only control (2 μ M)).

524

525 **RBD-SpyVLP thermostability and lyophilisation tests**

526 30 μ L of RBD-SpyVLP stored in thin-walled PCR tubes were subjected to freeze-thaw
527 cycles (one to five cycles) by storing the tubes in a -80 °C freezer for 15 min or until
528 the whole tube had frozen over, followed by incubation at RT for 10 min. For the
529 storage temperature study, 30 μ L of RBD-SpyVLP stored in thin-walled PCR tubes
530 were incubated at -80 °C, -20 °C, 4 °C or RT (25 °C) for 14 days. The RBD-SpyVLP
531 samples were then resolved on 4-12% Tris-Bis SDS-PAGE (Thermo Fisher) and
532 analysed by densitometry following Quick Coomassie (Generon) staining. All samples
533 were analysed in triplicate and plotted as mean \pm 1 standard deviation (SD). The
534 sample stored at -80 °C for two weeks, which had been through only 1 freeze-thaw
535 cycle, was defined as 100% soluble. For lyophilisation, 100 μ L of RBD-SpyVLP (125.5

536 µg/mL) in TBS pH 8.0 prepared in a Protein LoBind microcentrifuge tube (Fisher
537 Scientific) was snap-frozen in liquid nitrogen for 30 min. A BenchTop 2K freeze-dryer
538 (VisTis) was used for 24 h at 0.14 mbar and -72.5 °C to freeze-dry the sample.
539 Lyophilised sample was reconstituted in the same original volume (100 µL) of MilliQ
540 water and centrifuged at 16,900 g for 30 min to remove aggregates, before analysis
541 with SDS-PAGE or ELISA. For testing of RBD-SpyVLP on ELISA, 50 µL of RBD-
542 SpyVLP samples diluted in PBS (137 mM NaCl, 2.7 mM KCl, 10 mM Na₂HPO₄, 1.7
543 mM KH₂PO₄, pH 7.4) to 0.5 µg/mL were coated on NUNC plates at 4 °C overnight,
544 washed with PBS, and blocked with 300 µL of 5% skimmed milk in PBS for 1 h at RT.
545 Plates were then washed and incubated with 50 µL CR3022 (10 µg/mL) antibody (for
546 freeze-thaw and storage temperature study) or a panel of mAbs as indicated in the
547 graph (5 µg/mL) (for the lyophilisation study) diluted in PBS/0.1% BSA for 1 h at RT.
548 Plates were washed and incubated with horse radish peroxidase (HRP) conjugated
549 goat-anti-human IgG antibody (Dako, P0447) (diluted 1:1,600 in PBS/0.1% BSA) for
550 1 h at RT. Plates were then washed and developed with 50 µL of POD 3,3',5,5'-
551 tetramethylbenzidine (TMB) substrate (Roche) for 5 min and stopped with 50 µL of 1
552 M H₂SO₄. Absorbance was measured on a Clariostar plate reader (BMG Labtech). To
553 test the reactivity of RBD-SpyVLP against a panel of anti-SARS-CoV-2 RBD
554 antibodies, 50 µL of RBD-SpyVLP samples diluted in PBS to 0.5 µg/mL were coated
555 on NUNC plates at 4 °C overnight. Plates were then washed and blocked with 300 µL
556 of 5% (w/v) skimmed milk in PBS for 1 h at RT. Plates were washed and incubated
557 with antibodies diluted in PBS with 0.1% (w/v) BSA in a 2-fold dilution series in 50 µL
558 for 1 h. Second layer antibody was added as described above and plates were
559 developed as above.

560

561 **Dynamic light scattering (DLS)**

562 Samples were centrifuged for 30 min at 16,900 g at 4 °C to pellet possible aggregates.
563 Before each measurement, the quartz cuvette was incubated in the instrument for 5
564 min to stabilise the sample temperature. Samples were measured at 125-250 µg/mL
565 total protein concentration. 30 µL of sample was measured at 20 °C using an
566 Omnisizer (Victotek) with 20 scans of 10 s each. The settings were 50% laser intensity,
567 15% maximum baseline drift, and 20% spike tolerance. The intensity of the size
568 distribution was normalised to the peak value and plotted in GraphPad Prism 8
569 (GraphPad Software).

570

571 **Mouse immunisation and sampling**

572 To prepare the RBD-SpyVLP for vaccination at 125 µg/mL (based on SpyTag-RBD
573 concentration), 5 µM SpyTag-RBD was conjugated with 3.33 µM or 5 µM of
574 SpyCatcher003-mi3 in TBS pH 8.0 at 4 °C for 16 h. The reaction was centrifuged for
575 30 min at 17,000 g at 4 °C to remove potential aggregates. RBD-SpyVLP was
576 aliquoted and stored at -80 °C. Matching non-conjugated SpyTag-RBD or
577 SpyCatcher003-mi3 VLP were diluted in the same buffers and incubated and
578 centrifuged in the same way. RBD-SpyVLP (125 µg/mL) was diluted to 4 µg/mL (0.1
579 µg dose) or 20 µg/mL (0.5 µg dose) in the same buffer freshly before immunisation.
580 Before IM immunisation, RBD-SpyVLP was mixed 1:1 (25 µL + 25 µL) with AddaVax
581 adjuvant (Invivogen). Mouse experiments were performed according to the UK
582 Animals (Scientific Procedures) Act Project Licence (PBA43A2E4) and approved by
583 the University of Oxford Local Ethical Review Body. Female C57BL/6 or BALB/c mice
584 (~ 5 weeks old at the time of first immunisation) were obtained from BMS Oxford or
585 Envigo. Mice were housed in accordance with the UK Home Office ethical and welfare
586 guidelines and fed on standard chow and water ad libitum. Isoflurane (Abbott) lightly
587 anaesthetised mice were immunised on day 0 and day 14 IM with 50 µL of RBD-
588 SpyVLP at 0.1 µg or 0.5 µg or equivalent dose of unconjugated SpyTag-RBD or
589 SpyCatcher003-mi3 VLP. Sera samples were obtained on day 42 via cardiac puncture
590 of humanely sacrificed mice. The collected whole blood in Microtainer SST tubes (BD)
591 was allowed to clot at RT for 1 h before spinning down at 10,000 g for 5 min. The
592 clarified sera were heat-inactivated at 56 °C for 30 min before storing at -20 °C.

593

594 **RBD ELISA (Mouse and human sera)**

595 RBD-6H was expressed in Expi293 according to the manufacturer's protocol and
596 purified using HisTrap HP column (Cytivia) and desalted using Zeba Spin Desalting
597 Column (Thermo Fisher). To detect anti-RBD antibody in the immunised mouse sera,
598 50 µL purified RBD-6H (amino acids 330 to 532) (2 µg/mL) diluted in PBS was coated
599 on NUNC plates at 4 °C overnight. Plates were then washed with PBS and blocked
600 with 300 µL of 5% skimmed milk in PBS for 1 h at RT. In round-bottom 96-well plates,
601 heat-inactivated mouse sera (starting dilution 1 in 40) was diluted in PBS/0.1% BSA
602 in a 2-fold serial dilution in duplicate. 50 µL of the diluted sera was then transferred to
603 the NUNC plates for 1 h at RT. Plates were then washed with PBS and 50 µL of

604 secondary HRP goat anti-mouse antibody (Dako P0417) diluted 1:800 in PBS/0.1%
605 BSA was added to the wells for 1 h at RT. Plates were washed and developed as
606 described above. Serum RBD-specific antibody response was expressed as endpoint
607 titre (EPT). EPT is defined as the reciprocal of the highest serum dilution that gives a
608 positive signal (blank+10 SD) determined using a five-parameter logistic equation
609 calculated using GraphPad Prism 8. RBD antibody response in the convalescent
610 plasma was tested on a cell-based RBD ELISA, since human plasma gives high
611 background on NUNC plates in our hands. MDCK-RBD cells were generated as
612 previously described^{26, 39, 58}. Briefly, MDCK-SIAT1 cells (ECACC 05071502)⁵⁹ were
613 stably transfected using lentiviral vector to express SARS-CoV-2 RBD (amino acid
614 340-538, NITN...GPKK) fused at the C-terminus to the transmembrane domain of
615 haemagglutinin H7 (A/HongKong/125/2017) (EPI977395)
616 (KLSSGYKDVLWFSFGASCFILLAIVMGLVFICVKNGNMRCTICI) for surface
617 expression. MDCK-RBD was seeded at 3×10^4 cells/well in flat-bottom 96-well plates
618 and incubated overnight at 37 °C and 5% CO₂ prior to the assays. Human plasma was
619 incubated with MDCK-SIAT1 cells for 1 h at RT prior to the assays to remove
620 background binding on MDCK-SIAT1 cells. Plates seeded with MDCK-RBD were
621 washed with PBS and 50 µL of pre-absorbed human sera diluted in a 2-fold dilution
622 series (starting dilution 1 in 5) were added to the cells for 1 h at RT. The same set of
623 sera were added in parallel plates seeded with MDCK-SIAT1 to obtain background
624 binding on MDCK-SIAT1. Plates were washed and 50 µL of a secondary Alexa Fluor
625 647 goat anti-human antibody (Life Technologies A21455) (1:500 in PBS with 0.1%
626 (w/v) BSA) were added for 1 h at RT. Plates were washed and 100 µL of PBS/1%
627 formalin was added. Fluorescence signal was then read on a Clariostar plate reader.
628 Background signal obtained on the parallel MDCK-SIAT1 plates were subtracted. EPT
629 was determined as described above. Convalescent plasma samples were collected
630 from March to May 2020 at John Radcliffe Hospital, Oxford. Patients' severity was
631 determined according to the WHO guidance (described in⁶⁰).

632 633 **Spike glycoprotein ELISA (Mouse and human sera)**

634 A cell-based ELISA as described previously²⁶ was used to determine the anti-spike
635 glycoprotein antibody response in the mouse sera and convalescent plasma. Briefly,
636 MDCK-Spike was produced by stably transfecting parental MDCK-SIAT1 cells with
637 full-length SARS-CoV-2 spike glycoprotein cDNA using a lentiviral vector. MDCK-

638 Spike cells (3×10^4 cells/well) was seeded in 96-well plates and incubated overnight
639 at 37 °C. Mouse sera was diluted as above and 50 µL was transferred to the washed
640 plates seeded with MDCK-Spike cells for 1 h at RT. Human plasma was pre-incubated
641 with MDCK-SIAT1 cells as described above, before dilution and adding to MDCK-
642 Spike for 1 h at RT. Parallel plates seeded with MDCK-SIAT1 for background
643 subtraction was done as described above for human plasma. For mouse sera, 50 µL
644 of a secondary Alexa Fluor 647 goat-anti mouse antibody (1:500) (Life Technologies
645 A21235) was then added for 1 h at RT. For human sera, 50 µL of a secondary Alexa
646 Fluor 647 goat-anti human antibody (1:500) (Life Technologies A21455) was used.
647 Plates were then washed with PBS and 100 µL of PBS/1% formalin was added to each
648 well. Fluorescence signal was read on a Clariostar plate reader and the EPT titre was
649 calculated as described above.

650

651 **ACE2 competition ELISA**

652 ACE2 competition ELISA was done with RBD-SpyVLP immobilised on NUNC plates
653 as previously described with slight modifications ²⁶. ACE2-Fc was expressed in
654 Expi293 and purified using MabSelect SuRe column (Cytivia) and salts were removed
655 using Zeba Spin Desalting Column (Thermo Fisher) into PBS. ACE2-Fc was
656 chemically biotinylated using EZ-Link Sulfo-NHS-LC-Biotinylation Kit (Thermo Fisher)
657 according to manufacturer's protocol. 25 ng/well of RBD-SpyVLP was coated on the
658 ELISA plate at 4 °C overnight. Plates were then washed with PBS and blocked with
659 300 µL of 5% (w/v) skimmed milk in PBS for 1 h at RT. Heat-inactivated mouse or pig
660 sera or human plasma samples were titrated in duplicate as half-log₁₀ 8-point serial
661 dilution, starting at 1 in 5 in 30 µL with PBS with 0.1% (w/v) BSA. 30 µL of biotinylated
662 ACE2-Fc at 0.2 nM (40 ng/mL) (or 0.4 nM for human plasma) was added to the
663 samples. 50 µL of the biotinylated ACE2-Fc:sample mixture was transferred to the
664 RBD-SpyVLP coated plates and incubated for 1 h at RT. A secondary Streptavidin-
665 HRP (S911, Life Technologies) diluted to 1:1,600 in PBS/0.1% BSA was added to the
666 PBS-washed plates and incubated for 1 h at RT. Plates were then washed and
667 developed as above. Biotinylated ACE2-Fc without sera or plasma was used to obtain
668 the maximum signal and wells with PBS/BSA buffer only were used to determine the
669 minimum signal. Graphs were plotted as % binding of biotinylated ACE2 to RBD.
670 Binding % = (X - Min) / (Max - Min) * 100 where X = measurement of the sera or
671 plasma, Min = buffer only, Max = biotinylated ACE2-Fc alone. ACE2 blocking activity

672 of the sera or plasma was expressed as IC₅₀ determined using non-linear regression
673 curve fit using GraphPad Prism 8.

674

675 **Authentic SARS-CoV-2 virus neutralisation assay (PRNT)**

676 96 well plates containing a confluent monolayer of Vero-E6 cells were incubated with
677 10-20 plaque forming units (PFU) of SARS CoV-2 (hCoV-19/England/02/2020,
678 EPI_ISL_407073, kindly provided by Public Health England) and two-fold serial
679 dilution of heat-inactivated mouse sera for 3 h at 37 °C, 5% CO₂, in triplicate per serum
680 sample. Inoculum was then removed, and cells were overlaid with virus growth
681 medium containing Avicel microcrystalline cellulose (final concentration of 1.2%)
682 (Sigma-Aldrich). The plates were then incubated at 37 °C, 5% CO₂. At 24 h post-
683 infection, cells were fixed with 4% paraformaldehyde and permeabilised with 0.2% (v/v)
684 Triton-X-100 in PBS stained to visualise virus plaques, as described previously for the
685 neutralisation of influenza viruses ⁶¹, but using a rabbit polyclonal anti-NSP8 antibody
686 (Antibodies Online; ABIN233792) and anti-rabbit-HRP conjugate (Bio-Rad) and
687 detected using HRP on a TMB based substrate. Virus plaques were quantified and
688 ND₅₀ for sera was calculated using LabView software as described previously ⁶¹.

689

690 **Authentic SARS-CoV-2 plaque reduction neutralization assay (PRNT)**

691 SARS-CoV-2 (hCoV-19-Australia/VIC01/2020, GenBank MT007544) ⁶² was diluted to
692 a concentration of around 1,000 PFU/mL and 75 µL was mixed with an equal volume
693 of minimal essential medium (MEM) (Life Technologies) containing 1% (v/v) foetal
694 bovine serum (FBS) (Life Technologies) and 25 mM HEPES buffer (Sigma-Aldrich)
695 with doubling pooled mouse sera dilutions (starting dilution 1:40) in a 96-well V
696 bottomed plate. The plate was then incubated at 37 °C in a humidified incubator for 1
697 h before the virus-antibody mixture was transferred 24-well plates containing confluent
698 monolayers of Vero E6 cells (ECACC 85020206) cultured in MEM containing 10% (v/v)
699 FBS. The plates were incubated for 1 h at 37 °C and overlaid with MEM containing
700 1.5% carboxymethylcellulose (Sigma-Aldrich), 4% (v/v) FBS and 25mM HEPES buffer.
701 Plates were incubated at 37 °C for five days prior to fixation with 20% formalin in PBS
702 overnight. Plates were then washed with tap water and stained with 0.2% crystal violet
703 solution (Sigma-Aldrich) and plaques were visualised and counted. A mid-point probit
704 analysis (written in R programming language for statistical computing and graphics)

705 was used to determine the dilution of antibody required to reduce numbers of SARS-
706 CoV-2 virus plaques by 50% (ND₅₀) compared with the virus-only control (n = 5). A
707 human MERS convalescent serum known to neutralise SARS-CoV-2 (National
708 Institute for Biological Standards and Control, UK) was included in each run as assay
709 control.

710

711 **mAb competition assays**

712 mAb competition ELISA was done with RBD-SpyVLP immobilised on NUNC plates as
713 described above with slight modifications. Plates were coated and blocked as above.
714 Heat-inactivated mouse or pig sera or human plasma samples were titrated in
715 duplicate as half-log 8-point serial dilution, starting at 1 in 5 in 30 µL with PBS with
716 0.1% (w/v) BSA or tested at 1:20 in quadruplicates. 30 µL of chemically biotinylated
717 mAb (FI-3A, FD-11A, EY6A or S309) titrated to determine the lowest binding
718 concentration at top plateau (all mAbs produced in house)^{26, 38, 39} was added to the
719 sera. Biotinylation was conducted as described above. 50 µL of the biotinylated
720 mAb:sera mixture was transferred to the RBD-SpyVLP coated plates and incubated
721 for 1 h at RT. A secondary Streptavidin-HRP (Life Technologies, S911) diluted to
722 1:1,600 in PBS/0.1% BSA was added to the PBS-washed plates and incubated for 1
723 h at RT. Plates were then washed and developed as above. Biotinylated mAb without
724 sera or plasma was used to obtain the maximum signal and wells with PBS with 0.1%
725 (w/v) BSA buffer only were used to determine the minimum signal. Graphs were
726 plotted as % binding of biotinylated mAb to RBD-SpyVLP. Binding % = (X - Min) / (Max
727 - Min) * 100 where X = measurement of the sera or plasma, Min = buffer only, Max =
728 biotinylated mAb alone.

729

730 **Pig immunisation and sampling**

731 Pig studies were performed in accordance with the UK Animals (Scientific Procedures)
732 Act 1986 and with approval from the local Animal Welfare and Ethical Review Body
733 (AWERB) (Project Licence PP1804248). RBD-SpyVLP for pig immunisation was
734 prepared as above. Nine weaned, Large White-Landrace-Hampshire cross-bred pigs
735 of 8–10 weeks of age from a commercial rearing unit were randomly allocated to three
736 treatment groups (5 µg RBD-SpyVLP, 50 µg RBD-SpyVLP or 100 µg spike
737 glycoprotein) (n = 3). RBD-SpyVLP was diluted to 5 µg/mL or 50 µg/mL in 25 mM Tris-
738 HCl, pH 8.0, 150 mM NaCl and mixed with an equal amount of AddaVax (Invivogen)

739 (1 mL + 1 mL) prior to immunisation. The spike glycoprotein was a soluble trimeric
740 spike with the pre-fusion stabilisation substitutions (K983P, V984P, furin cleavage site
741 removed and inclusion of a C-terminal T4-foldon domain for trimerization)⁶³. The spike
742 glycoprotein was expressed in Expi293 cells and purified as previously described³⁴.
743 Briefly, supernatant containing soluble spike glycoprotein was purified using
744 immobilised metal affinity followed by gel filtration in TBS (pH 7.4). Pigs were dosed
745 via IM injection into the brachiocephalic muscle with 2 mL of RBD-SpyVLP (5 µg or 50
746 µg) or spike (100 µg) at day 0 and day 28. Blood samples were taken on a weekly
747 basis at 0, 7, 14, 21, 28, 35, 42 and 56 days post-immunisation (DPI) by venepuncture
748 of the external jugular vein: 8 mL/pig in BD SST vacutainer tubes (Fisher Scientific)
749 for serum collection and 40 mL/pig in BD heparin vacutainer tubes (Fisher Scientific)
750 for PBMC isolation. Additional heparin blood samples were collected on 31 and 33
751 DPI to track the plasma cell response to boost. Sera samples were stored at -20 °C
752 and heat-inactivated at 56 °C for 2 h before use in pVNT or VNT assays. Oral and
753 nasal swabs were collected weekly and placed in 500 µL Media 199 (Thermo Fisher)
754 supplemented with 0.0025% Nystatin (Merck), 0.01% Penicillin-Streptomycin (Gibco),
755 0.025% 1M HEPES solution (Gibco), 0.005% (w/v) sodium bicarbonate (Merck) and
756 0.067% (w/v) BSA (Merck) (VTM). Swabs were centrifuged at 700 × g for 5 min before
757 aspirating the liquid and storing with the swab at -20 °C. Prior to assessment of
758 antibodies, swabs and VTM were loaded in Spin-X Centrifuge 0.45 µM columns
759 (Fisher Scientific) and fluid collected by centrifugation at 21,000 × g for 5 min.
760

761 **RBD ELISA (Pig sera and swab fluids)**

762 An ELISA to analyse anti-RBD antibody response in pig sera was performed as
763 previously described³⁴. Briefly, 50 µL of 2 µg/mL purified RBD-6H as described above
764 was coated on flat-bottomed 96-well plates (Immuron 4 HBX; Thermo Fisher Scientific)
765 overnight at 4 °C. Plates were washed with TBS (pH 7.4) with 0.1% (v/v) Tween-20
766 and blocked with 100 µL of PBS containing 3% skimmed milk for 1 h at RT. Pig sera
767 samples were diluted in PBS with 1% (w/v) skimmed milk and 0.1% (v/v) Tween-20 in
768 a 2-fold serial dilution starting at 1:10 dilution and 100 µL of the diluted sera was added
769 to the coated plates for 1 h at RT. A conjugated secondary goat anti-pig IgG HRP
770 (Abcam, Cambridge, UK) at 1:10,000 dilution in PBS with 1% (w/v) skimmed milk and
771 0.1% (v/v) Tween-20 was added for 1 h at RT. Plates were washed and 100 µL TMB
772 (One Component Horse Radish Peroxidase Microwell Substrate, BioFX, Cambridge

773 Bioscience) was added to each well and the plates were incubated for 7 min at RT.
774 100 μ L BioFX 450nmStop Reagent (Cambridge Bioscience) was then added and
775 absorbance was determined using a microplate reader. End-point antibody titres
776 (mean of duplicates) were defined as following: the \log_{10} OD was plotted against the
777 \log_{10} sample dilution and a regression analysis of the linear part of this curve allowed
778 calculation of the endpoint titre with an absorbance of twice the mean absorbance of
779 pre-immunised sera. RBD-specific antibody titres in oral and nasal swab fluids were
780 determined by ELISA as detailed above except that the conjugated secondary
781 antibody was replaced with either goat anti-porcine IgG HRP (Bio-Rad Antibodies) at
782 1:20,000 dilution in PBS with 1% (w/v) skimmed milk and 0.1% (v/v) Tween-20 or goat
783 anti-porcine IgA HRP (Bio-Rad Antibodies) at 1:20,000 dilution in the same diluent.

784

785 **Virus neutralization assay (VNT) (pig sera)**

786 VNT on pig sera was done as described previously ³⁴. Briefly, Vero E6 cells were
787 seeded in 96-well flat-bottom plates (1×10^5 cells/mL) and incubated at 37 °C
788 overnight prior to the assays. Two-fold serial dilutions of sera (starting dilution 1 in 5)
789 in quadruplet were prepared in 96-well round-bottom plates using Dulbecco's Modified
790 Eagle Medium (DMEM) with 1% (v/v) FBS and 1% Antibiotic-Antimycotic (Gibco). 75
791 μ L of the diluted sera was mixed with an equal volume of media containing 64 PFU of
792 SARS-CoV-2 virus (hCoV-19/England/02/2020, EPI_ISL407073) and incubated for 1
793 h at 37 °C. Media in the wells seeded with Vero E6 was replaced with 100 μ L DMEM
794 with 10% (v/v) FBS and 1% Antibiotic-Antimycotic (Gibco) and 100 μ L of the sera-virus
795 mixture was added into the wells. The plates were incubated for six days at 37 °C.
796 Cytopathic effect (CPE) was monitored on a brightfield microscopy, and by fixation
797 using formaldehyde (VWR, Leighton Buzzard, UK) and staining using 0.1% Toluidine
798 Blue (Sigma-Aldrich). CPE was scored by researchers who were blinded to the identity
799 of the samples. No sera or no virus controls were run in parallel on each plate.
800 Neutralisation titres (ND₅₀) were expressed as the reciprocal of the serum dilution that
801 prevented CPE in 50% of the wells.

802

803 **Pseudovirus neutralisation test (pVNT) (pig sera)**

804 Lentiviral-based SARS-CoV-2 pseudoviruses were generated as described previously
805 ³⁴. Briefly, HEK293T cells were seeded at a density of 7.5×10^5 in 6-well plates before
806 transfection with the following plasmids: 500 ng of SARS-CoV-2 spike, 600 ng p8.91

807 (HIV-1 gag-pol), 600 ng CSFLW (lentiviral genome plasmid encoding a firefly
808 luciferase transgene)³⁴ using 10 µL polyethylene imine (PEI) (1 µg/mL) In Opti-MEM
809 media (Thermo Fisher). The media was replaced with 3 mL DMEM with 10% FBS and
810 incubated at 37 °C. Supernatant containing pseudovirus was harvested at 48 h and
811 72 h post transfection. Collected supernatant was centrifuged at 1,300 × g for 10 min
812 at 4 °C to remove debris. To perform the assay, 2 × 10⁴ HEK293T target cells
813 transfected with 500 ng of a human ACE2 expression plasmid (Addgene) were seeded
814 in a white flat-bottom 96-well plate one day prior to the assays. Pig sera were diluted
815 with a four-fold serial dilution in serum-free media (starting dilution 1 in 20) and 50 µL
816 was added to a 96-well plate in quadruplicate. Pseudovirus pre-titrated to give 1 × 10⁶
817 relative light unit (RLU) in 50 µL DMEM with 10% FBS was added to the sera and
818 incubated at 37 °C for 1 h. The pseudovirus:sera mix was then transferred to the target
819 cells and incubated at 37 °C for 72 h. Firefly luciferase activity was measured using
820 BrightGlo luciferase reagent on a GloMax-Multi+ Detection System (Promega).
821 Pseudovirus neutralisation titres (ND₅₀) were expressed as the reciprocal of the serum
822 dilution that inhibited luciferase signal in 50% of the wells.

823

824 **Intracellular cytokine staining assay (pig PBMC)**

825 Assessment of intracellular cytokine expression following stimulation of PBMCs with
826 synthetic peptides representing SARS-CoV-2 S protein was conducted as described
827 previously³⁴. In brief, PBMCs were isolated from heparinized blood by density gradient
828 centrifugation and suspended at 1 × 10⁷ cells/mL in RPMI-1640 medium, GlutaMAX
829 supplement, HEPES (Gibco) supplemented with 10% (v/v) heat-inactivated FBS (New
830 Zealand origin, Life Science Production), 1% Penicillin-Streptomycin and 0.1% 2-
831 mercaptoethanol (50 mM; Gibco) (cRPMI). 50 µL PBMC were added per well to 96-
832 well round bottom plates and stimulated in triplicate wells with SARS-CoV-2 S peptide
833 pools at a final concentration of 1 µg/mL peptide. Unstimulated cells in triplicate wells
834 were used as a negative control. After 14 h incubation at 37°C, 5% CO₂, cytokine
835 secretion was blocked by addition 1:1,000 BD GolgiPlug (BD Biosciences) and cells
836 were further incubated for 6 h. PBMC were surface-labelled with Zombie NIR fixable
837 viability stain (BioLegend), CD3-FITC mAb (clone BB23-8E6-8C8, BD Biosciences),
838 CD4-PerCP-Cy5.5 mAb (clone 74-12-4, BD Biosciences) and CD8α-PE mAb (clone
839 76-2-11, BD Biosciences). After fixation (Fixation Buffer, BioLegend) and
840 permeabilization (Permeabilization Wash Buffer, BioLegend), cells were stained with:

841 IFN- γ -Alexa Fluor 647 mAb (clone CC302, Bio-Rad Antibodies,) and TNF- α -Brilliant
842 Violet 421 mAb (clone Mab11, BioLegend). Cells were analysed using a BD
843 LSRFortessa flow cytometer (BD Biosciences) and data analysed using FlowJo
844 software (BD Biosciences). Total SARS-CoV-2 S-specific IFN- γ -positive responses for
845 live CD3 $^+$ CD4 $^+$ and CD3 $^+$ CD4 $^-$ CD8 $^+$ T cells are presented after subtraction of the
846 background response detected in the media-stimulated control PBMC samples of
847 each pig, prior to summing together the frequency of S-peptide pools 1-3 specific cells.
848

849 **RBD-tetramer staining assay (pig PBMC)**

850 A biotinylated form of RBD was generated for B-cell tetramer staining assays. An RBD
851 protein with a C-terminal biotin acceptor peptide (RBD-BAP) was expressed from
852 plasmid pOPINTTGNeo in Expi293 cells according to the manufacturer's instructions.
853 Culture supernatants were clarified by centrifugation and purified through a 5 mL
854 HisTrap FF column (GE Healthcare), using the $\ddot{\text{A}}$ KTA Pure chromatography system
855 (Cytiva). Fractions containing RBD-BAP were concentrated and the excess imidazole
856 removed by buffer exchange using an Amicon 10kDa (Merck). RBD-BAP was
857 biotinylated using *E. coli* biotin ligase (BirA). GST-BirA enzyme was expressed,
858 purified and biotinylated as previously described ⁶⁴. Biotinylation reactions were
859 assembled with 100 μ M RBD-BAP, 1 μ M GST-BirA, 5mM magnesium chloride
860 (Ambion), 2mM ATP and 150 μ M D-Biotin (both Merck) and incubated twice for one
861 hour at 30 °C with additional fresh biotin and GST-BirA added in-between incubations.
862 GST-BirA was removed from the reaction with a GST HiTrap column, as above, and
863 RBD-BAP was purified out by dialysis as above. Biotinylation of RBD-BAP was
864 confirmed by streptavidin band shift assay ⁶⁴ and quantified by BCA assay (Pierce).
865 RBD tetramers were assembled by combining biotinylated RBD with streptavidin-
866 Brilliant Violet 421 or streptavidin-Brilliant Violet 650 (both BioLegend) at a molar ratio
867 of 4:1. Negative control 'decoy' tetramers were similarly assembled using biotinylated
868 Nipah virus soluble glycoprotein ⁶⁵ and streptavidin-PerCP Cy5.5 (BioLegend).
869

870 PBMC were stained with SARS-CoV-2 S RBD-tetramers to assess the frequency of
871 circulating specific B cells during the course of the study. For 28, 31, 33 and 35 days
872 post-infection, fresh PBMC were analysed (in triplicate), while for 0, 7, 14, 42 and 56
873 days post-infection, previously cryopreserved PBMC were assessed (in quadruplicate).
874 RBD tetramers were assembled by combining biotinylated RBD with streptavidin-

875 Brilliant Violet 421 or streptavidin-Brilliant Violet 650 (both BioLegend) at a molar ratio
876 of 4:1. Negative control 'decoy' tetramers were similarly assembled using biotinylated
877 Nipah virus soluble glycoprotein⁶⁵ and streptavidin-PerCP Cy5.5 (BioLegend). PBMC
878 were washed in cold PBS and seeded at 1×10^6 cells/well in 96-well round bottom
879 plates and with combinations of RBD and decoy tetramers by incubation for 30 min on
880 ice. After washing, cells were stained with Zombie Aqua, CD3-PE-Cy7 mAb (clone
881 BB23-8E6-8C8, BD Biosciences), CD14-PE Vio 770 mAb (clone REA599, Miltenyi
882 Biotec), IgG-Alexa Fluor-647 mAb (Cohesion Biosciences, Generon,), IgA-FITC
883 polyclonal Ab (BioRad Antibodies) and IgM-PE mAb (clone K52 1C3, BioRad
884 Antibodies; conjugated using Lightning-Link® PE Antibody Labeling Kit, Expedeon,
885 Abcam), for 30 min on ice. After washing and fixation in 4% paraformaldehyde for 30
886 min at 4°C, cells were analysed on a BD LSRFortessa flow cytometer with downstream
887 analysis using FlowJo software. Following exclusion of dead cells, CD3⁺, CD14⁺ and
888 decoy tetramer⁺ cells, the percentage of IgA⁺, IgG⁺ or IgM⁺ cells dual-labelled with
889 both RBD tetramers was assessed.

890
891 **RBD IgG ELISpot assay (pig PBMC)**

892 Sterile 96-well Multiscreen-HA filter plates with a mixed cellulose membrane
893 (MAHAS4510, Millipore) were coated with 100 µL of 15 µg/mL anti-porcine IgG mAb
894 (clone MT421, Mabtech, 2BScientific) diluted in 0.05M carbonate-bicarbonate buffer
895 pH 9.2 (Merck). Coated plates were incubated for a minimum of 18 h at 4 °C. Plates
896 were then washed with PBS and blocked using cRPMI for at least 1 h at 37 °C, 5%
897 CO₂, 95% humidity. Blocking solution was then removed, and PBMC were added at a
898 density of 5×10^5 /well for antigen-specific response or at 5×10^4 /well for wells
899 assigned to total IgG (positive control). The plates were then incubated for 18 h at
900 37 °C, 5% CO₂, 95% humidity. Media was removed and cells were lysed with cold
901 distilled water, followed by three PBS washes as before. To measure total IgG, 50
902 µL/well of biotinylated anti-IgG mAb (clone MT424-biotin, Mabtech) was added at 0.5
903 µg/mL. To assess antigen-specific responses 50 µL/well of biotinylated SARS-CoV-2
904 RBD was added at 2.5 µg/mL. As a negative control, 50 µL/well of biotinylated Nipah
905 G protein was added to the relevant wells at 2.5 µg/mL. All antigens were diluted in
906 PBS with 0.5% (v/v) FCS. An additional set of negative control wells were also
907 prepared by adding 50 µL/well PBS with 0.5% (v/v) FCS. Each condition was tested
908 in triplicate. Plates were incubated for 2 h at RT, before washing five times with PBS.

909 Following this, 50 µL/well of streptavidin-alkaline phosphatase (streptavidin-ALP)
910 enzyme conjugate (Mabtech) (diluted 1:1,000 in PBS with 0.5% (v/v) FCS) was added
911 to each well and plates were incubated for 1 h at RT (protected from light).
912 Streptavidin-ALP was removed, and plates were washed another 5 times with PBS,
913 followed by addition of 50 µL/well BCIP/NBTplus substrate (Mabtech), neat. Plates
914 were left for 30 min, until distinct spots developed. Finally, development was stopped
915 by addition of 150 µL/well of 4 °C distilled water followed by rinsing both the front and
916 back of the plates with copious tap water. Plates were air-dried, before spots were
917 counted using a CTL ImmunoSpot Analyzer (Cellular Technologies).

918

919 **Statistical analysis**

920 All statistical analyses were performed using GraphPad Prism 8 (GraphPad Software).
921 Statistical differences were analysed using either Mann-Whitney U test or Kruskal-
922 Wallis test followed by Dunn's multiple comparisons. A p value <0.05 was deemed
923 statistically significant.

924

925 **CONTRIBUTIONS**

926 **Production & characterisation of RBD-SpyVLP and mouse immunogenicity**
927 **studies:** T.K.T. performed the RBD-SpyVLP conjugation, *in vitro* antigenicity and
928 stability characterisation, and all mouse experiments and analysis. P.R. produced the
929 RBD protein, developed mice and human immunoassays and performed analysis on
930 mouse and pig sera. L.S. performed competition ELISA on mouse and pig sera. R.R.
931 and A.H.K. produced and purified the SpyCatcher003 VLP, purified the RBD protein,
932 and performed DLS and lyophilisation. T.K.T., P.R., M.H. and A.R.T. conceived and
933 designed the experiments. KY.A.H provided the monoclonal antibodies. S.H., R.H.,
934 R.S.D. and J.W.M. designed and performed neutralisation assays on the mouse sera.
935 J.A.T., K.R.B. and M.W.C performed the VNT on the pooled mouse sera. **Pig**
936 **immunogenicity studies:** J.W.P.H., J.C.E., R.K.M, M.P., V.M., K.M., C.C., R.W.,
937 A.G., M.A., V.M., and S.P.G conducted the pig immunogenicity study, processed
938 samples and conducted the T cell and B cell analyses. N.Z., C.C., M.T., and D.B.
939 conducted the pVNT assays with pig sera. I.D., H.S., A.Z., D.B., S.B., P.S.L. and P.H.
940 conducted the VNT assays with pig sera. A.L., G.W. and C.B. conducted the ELISAs
941 with pig sera and swab fluids. J.N., A.S.A, A.B., S. C., T.M., J.H. and R.A. produced
942 recombinant RBD, RBD-biotin and spike protein. R.W., J.A.H., E.T. B.C., T.J.T., and

943 S.P.G. conceived and designed experiments. T.K.T. prepared the manuscript. All
944 authors read, reviewed and approved the manuscript.

945

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963

964

965 **COMPETING INTERESTS**

966 M.H. is an inventor on a patent regarding spontaneous amide bond formation
967 (EP2534484) and a SpyBiotech founder, shareholder and consultant. M.H. and A.H.K.
968 are inventors on a patent application regarding SpyTag003:SpyCatcher003 (UK
969 Intellectual Property Office 1706430.4).

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977 **REFERENCES**

- 978 1. Zhu, N. et al. A Novel Coronavirus from Patients with Pneumonia in China, 2019. *N
979 Engl J Med* **382**, 727-733 (2020).
- 980 2. World Health Organization (2020).
- 981 3. World Health Organization (2020).
- 982 4. Coronaviridae Study Group of the International Committee on Taxonomy of, V. The
983 species Severe acute respiratory syndrome-related coronavirus: classifying 2019-nCoV
984 and naming it SARS-CoV-2. *Nat Microbiol* **5**, 536-544 (2020).
- 985 5. Wu, F. et al. Author Correction: A new coronavirus associated with human respiratory
986 disease in China. *Nature* **580**, E7 (2020).
- 987 6. Walls, A.C. et al. Structure, Function, and Antigenicity of the SARS-CoV-2 Spike
988 Glycoprotein. *Cell* **181**, 281-292 e286 (2020).
- 989 7. Shang, J. et al. Structural basis of receptor recognition by SARS-CoV-2. *Nature* **581**,
990 221-224 (2020).
- 991 8. Yan, R. et al. Structural basis for the recognition of SARS-CoV-2 by full-length human
992 ACE2. *Science* **367**, 1444-1448 (2020).
- 993 9. Zhou, P. et al. A pneumonia outbreak associated with a new coronavirus of probable
994 bat origin. *Nature* **579**, 270-273 (2020).
- 995 10. Gomes, A.C., Mohsen, M. & Bachmann, M.F. Harnessing Nanoparticles for
996 Immunomodulation and Vaccines. *Vaccines (Basel)* **5** (2017).
- 997 11. Jiang, S., Lu, L., Liu, Q., Xu, W. & Du, L. Receptor-binding domains of spike proteins
998 of emerging or re-emerging viruses as targets for development of antiviral vaccines.
Emerg Microbes Infect **1**, e13 (2012).
- 1000 12. Zhou, Y., Jiang, S. & Du, L. Prospects for a MERS-CoV spike vaccine. *Expert Rev
1001 Vaccines* **17**, 677-686 (2018).
- 1002 13. Quinlan BD, M.H., Zhang L, Guo Y, He W, Ojha A, Parcells MS, Luo G, Li W, Zhong
1003 G, Choe H, Farzan M The SARS-CoV-2 receptor-binding domain elicits a potent
1004 neutralizing response without antibody-dependent enhancement. *bioRxiv* (2020).
- 1005 14. Yang, J. et al. A vaccine targeting the RBD of the S protein of SARS-CoV-2 induces
1006 protective immunity. *Nature* (2020).
- 1007 15. Ju, B. et al. Human neutralizing antibodies elicited by SARS-CoV-2 infection. *Nature*
1008 **584**, 115-119 (2020).
- 1009 16. Rogers, T.F. et al. Isolation of potent SARS-CoV-2 neutralizing antibodies and
1010 protection from disease in a small animal model. *Science* **369**, 956-963 (2020).
- 1011 17. Tokatlian, T. et al. Enhancing Humoral Responses Against HIV Envelope Trimers via
1012 Nanoparticle Delivery with Stabilized Synthetic Liposomes. *Sci Rep* **8**, 16527 (2018).
- 1013 18. Rahikainen, R. et al. Overcoming Symmetry Mismatch in Vaccine Nanoassembly with
1014 SpyCatcher003 Amidation. *Angew Chem* (2020).
- 1015 19. Brune, K.D. et al. Plug-and-Display: decoration of Virus-Like Particles via isopeptide
1016 bonds for modular immunization. *Sci Rep* **6**, 19234 (2016).
- 1017 20. Bruun, T.U.J., Andersson, A.C., Draper, S.J. & Howarth, M. Engineering a Rugged
1018 Nanoscaffold To Enhance Plug-and-Display Vaccination. *ACS Nano* **12**, 8855-8866
1019 (2018).
- 1020 21. Escolano, A. et al. Immunization expands B cells specific to HIV-1 V3 glycan in mice
1021 and macaques. *Nature* **570**, 468-473 (2019).
- 1022 22. Thrane, S. et al. Bacterial superglue enables easy development of efficient virus-like
1023 particle based vaccines. *J Nanobiotechnology* **14**, 30 (2016).
- 1024 23. Khairil Anuar, I.N.A. et al. Spy&Go purification of SpyTag-proteins using pseudo-
1025 SpyCatcher to access an oligomerization toolbox. *Nat Commun* **10**, 1734 (2019).

1026 24. Keeble, A.H. et al. Approaching infinite affinity through engineering of peptide-protein
1027 interaction. *Proc Natl Acad Sci U S A* (2019).

1028 25. Hsia, Y. et al. Corrigendum: Design of a hyperstable 60-subunit protein icosahedron.
1029 *Nature* **540**, 150 (2016).

1030 26. Huang, K.Y.A. et al. Plasmablast-derived antibody response to acute SARS-CoV-2
1031 infection in humans. *bioRxiv* (2020).

1032 27. ter Meulen, J. et al. Human monoclonal antibody combination against SARS
1033 coronavirus: synergy and coverage of escape mutants. *PLoS Med* **3**, e237 (2006).

1034 28. Wrapp, D. et al. Structural Basis for Potent Neutralization of Betacoronaviruses by
1035 Single-Domain Camelid Antibodies. *Cell* **181**, 1436-1441 (2020).

1036 29. Wang, Q. et al. Structural and Functional Basis of SARS-CoV-2 Entry by Using Human
1037 ACE2. *Cell* **181**, 894-904 e899 (2020).

1038 30. Preiss, S., Garcon, N., Cunningham, A.L., Strugnell, R. & Friedland, L.R. Vaccine
1039 provision: Delivering sustained & widespread use. *Vaccine* **34**, 6665-6671 (2016).

1040 31. Podda, A. & Del Giudice, G. MF59-adjuvanted vaccines: increased immunogenicity
1041 with an optimal safety profile. *Expert Rev Vaccines* **2**, 197-203 (2003).

1042 32. Meurens, F., Summerfield, A., Nauwynck, H., Saif, L. & Gerdts, V. The pig: a model
1043 for human infectious diseases. *Trends Microbiol* **20**, 50-57 (2012).

1044 33. Overgaard, N.H. et al. Establishing the pig as a large animal model for vaccine
1045 development against human cancer. *Front Genet* **6**, 286 (2015).

1046 34. Graham, S.P. et al. Evaluation of the immunogenicity of prime-boost vaccination with
1047 the replication-deficient viral vectored COVID-19 vaccine candidate ChAdOx1 nCoV-
1048 19. *NPJ Vaccines* **5**, 69 (2020).

1049 35. Altman, M.O., Angeletti, D. & Yewdell, J.W. Antibody Immunodominance: The Key
1050 to Understanding Influenza Virus Antigenic Drift. *Viral Immunol* **31**, 142-149 (2018).

1051 36. Huang, K.Y. et al. Focused antibody response to influenza linked to antigenic drift. *J
1052 Clin Invest* **125**, 2631-2645 (2015).

1053 37. Baum, A. et al. Antibody cocktail to SARS-CoV-2 spike protein prevents rapid
1054 mutational escape seen with individual antibodies. *Science* **369**, 1014-1018 (2020).

1055 38. Pinto, D. et al. Cross-neutralization of SARS-CoV-2 by a human monoclonal SARS-
1056 CoV antibody. *Nature* **583**, 290-295 (2020).

1057 39. Zhou, D. et al. Structural basis for the neutralization of SARS-CoV-2 by an antibody
1058 from a convalescent patient. *Nat Struct Mol Biol* (2020).

1059 40. Ju, B. et al. Potent human neutralizing antibodies elicited by SARS-CoV-2 infection.
1060 *bioRxiv* (2020).

1061 41. Brouwer, P.J.M. et al. Potent neutralizing antibodies from COVID-19 patients define
1062 multiple targets of vulnerability. *Science* **369**, 643-650 (2020).

1063 42. Cao, Y. et al. Potent Neutralizing Antibodies against SARS-CoV-2 Identified by High-
1064 Throughput Single-Cell Sequencing of Convalescent Patients' B Cells. *Cell* **182**, 73-84
1065 e16 (2020).

1066 43. Hansen, J. et al. Studies in humanized mice and convalescent humans yield a SARS-
1067 CoV-2 antibody cocktail. *Science* **369**, 1010-1014 (2020).

1068 44. Kreer, C. et al. Longitudinal Isolation of Potent Near-Germline SARS-CoV-2-
1069 Neutralizing Antibodies from COVID-19 Patients. *Cell* **182**, 843-854 e812 (2020).

1070 45. Liu, L. et al. Potent neutralizing antibodies against multiple epitopes on SARS-CoV-2
1071 spike. *Nature* **584**, 450-456 (2020).

1072 46. Robbiani, D.F. et al. Convergent antibody responses to SARS-CoV-2 in convalescent
1073 individuals. *Nature* **584**, 437-442 (2020).

1074 47. Zhang, N. et al. Identification of an ideal adjuvant for receptor-binding domain-based
1075 subunit vaccines against Middle East respiratory syndrome coronavirus. *Cell Mol*
1076 *Immunol* **13**, 180-190 (2016).

1077 48. Islam, M.R. et al. Genome-wide analysis of SARS-CoV-2 virus strains circulating
1078 worldwide implicates heterogeneity. *Sci Rep* **10**, 14004 (2020).

1079 49. Scudellari, M. (Nature Publishing Group, Nature; 2020).

1080 50. Walls, A.C. et al. Elicitation of potent neutralizing antibody responses by designed
1081 protein nanoparticle vaccines for SARS-CoV-2. *bioRxiv* (2020).

1082 51. Gao, Q. et al. Development of an inactivated vaccine candidate for SARS-CoV-2.
1083 *Science* **369**, 77-81 (2020).

1084 52. van Doremalen, N. et al. ChAdOx1 nCoV-19 vaccination prevents SARS-CoV-2
1085 pneumonia in rhesus macaques. *bioRxiv* (2020).

1086 53. Keech, C. et al. First-in-Human Trial of a SARS CoV 2 Recombinant Spike Protein
1087 Nanoparticle Vaccine. *medRxiv* (2020).

1088 54. Prevost, J. et al. Cross-sectional evaluation of humoral responses against SARS-CoV-
1089 2 Spike. *bioRxiv* (2020).

1090 55. Seow, J. et al. Longitudinal evaluation and decline of antibody responses in SARS-
1091 CoV-2 infection. *medRxiv* (2020).

1092 56. Kanekiyo, M. et al. Mosaic nanoparticle display of diverse influenza virus
1093 hemagglutinins elicits broad B cell responses. *Nat Immunol* **20**, 362-372 (2019).

1094 57. Zakeri, B. et al. Peptide tag forming a rapid covalent bond to a protein, through
1095 engineering a bacterial adhesin. *Proc Natl Acad Sci U S A* **109**, E690-697 (2012).

1096 58. Huo, J. et al. Neutralization of SARS-CoV-2 by Destruction of the Prefusion Spike.
1097 *Cell Host Microbe* (2020).

1098 59. Matrosovich, M., Matrosovich, T., Carr, J., Roberts, N.A. & Klenk, H.D.
1099 Overexpression of the alpha-2,6-sialyltransferase in MDCK cells increases influenza
1100 virus sensitivity to neuraminidase inhibitors. *J Virol* **77**, 8418-8425 (2003).

1101 60. Peng, Y. et al. Broad and strong memory CD4 (+) and CD8 (+) T cells induced by
1102 SARS-CoV-2 in UK convalescent COVID-19 patients. *bioRxiv* (2020).

1103 61. Lin, Y. et al. Optimisation of a micro-neutralisation assay and its application in
1104 antigenic characterisation of influenza viruses. *Influenza Other Respir Viruses* **9**, 331-
1105 340 (2015).

1106 62. Caly, L. et al. Isolation and rapid sharing of the 2019 novel coronavirus (SARS-CoV-
1107 2) from the first patient diagnosed with COVID-19 in Australia. *Med J Aust* **212**, 459-
1108 462 (2020).

1109 63. Amanat, F. et al. A serological assay to detect SARS-CoV-2 seroconversion in humans.
1110 *medRxiv* (2020).

1111 64. Fairhead, M. & Howarth, M. Site-specific biotinylation of purified proteins using BirA.
1112 *Methods Mol Biol* **1266**, 171-184 (2015).

1113 65. Pedrera, M. et al. Bovine Herpesvirus-4-Vectored Delivery of Nipah Virus
1114 Glycoproteins Enhances T Cell Immunogenicity in Pigs. *Vaccines (Basel)* **8** (2020).

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1116

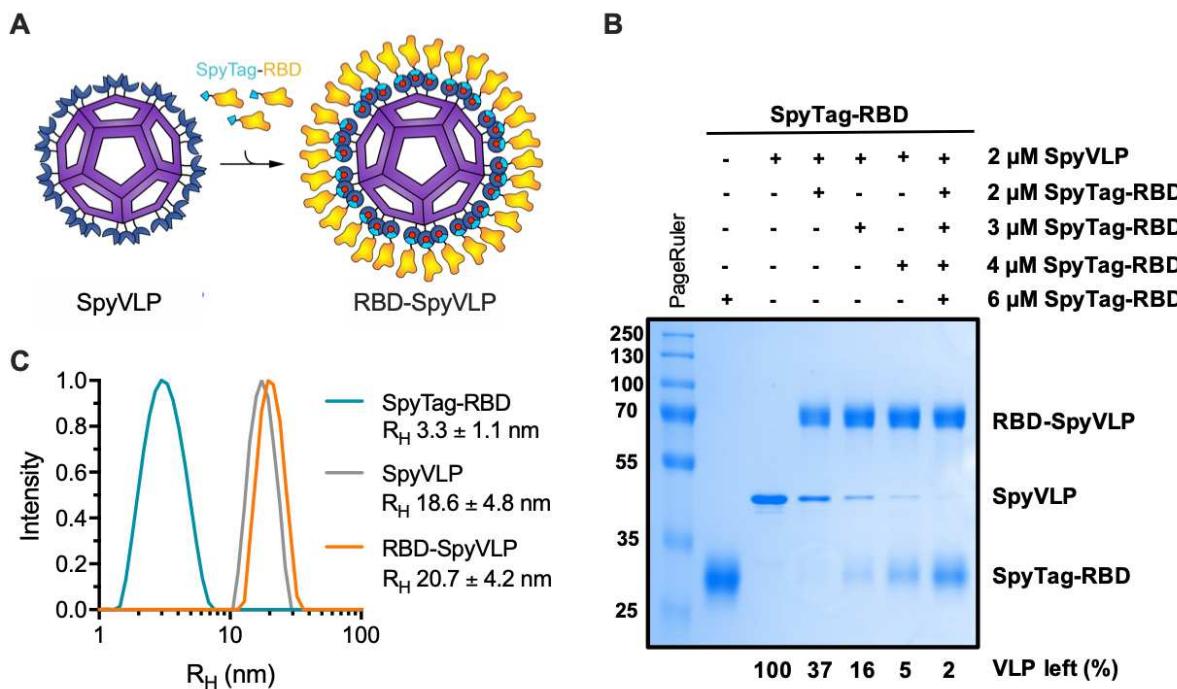
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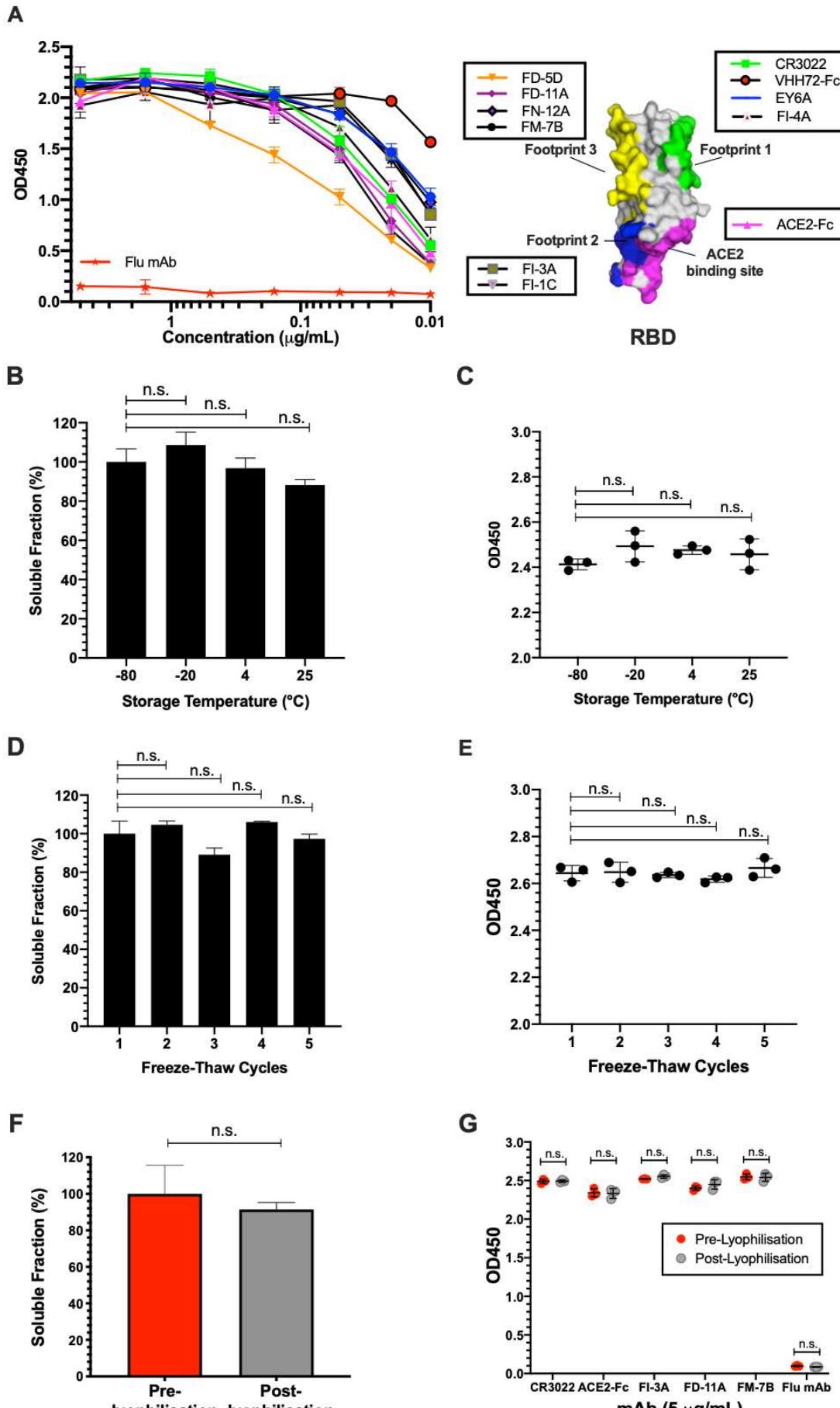
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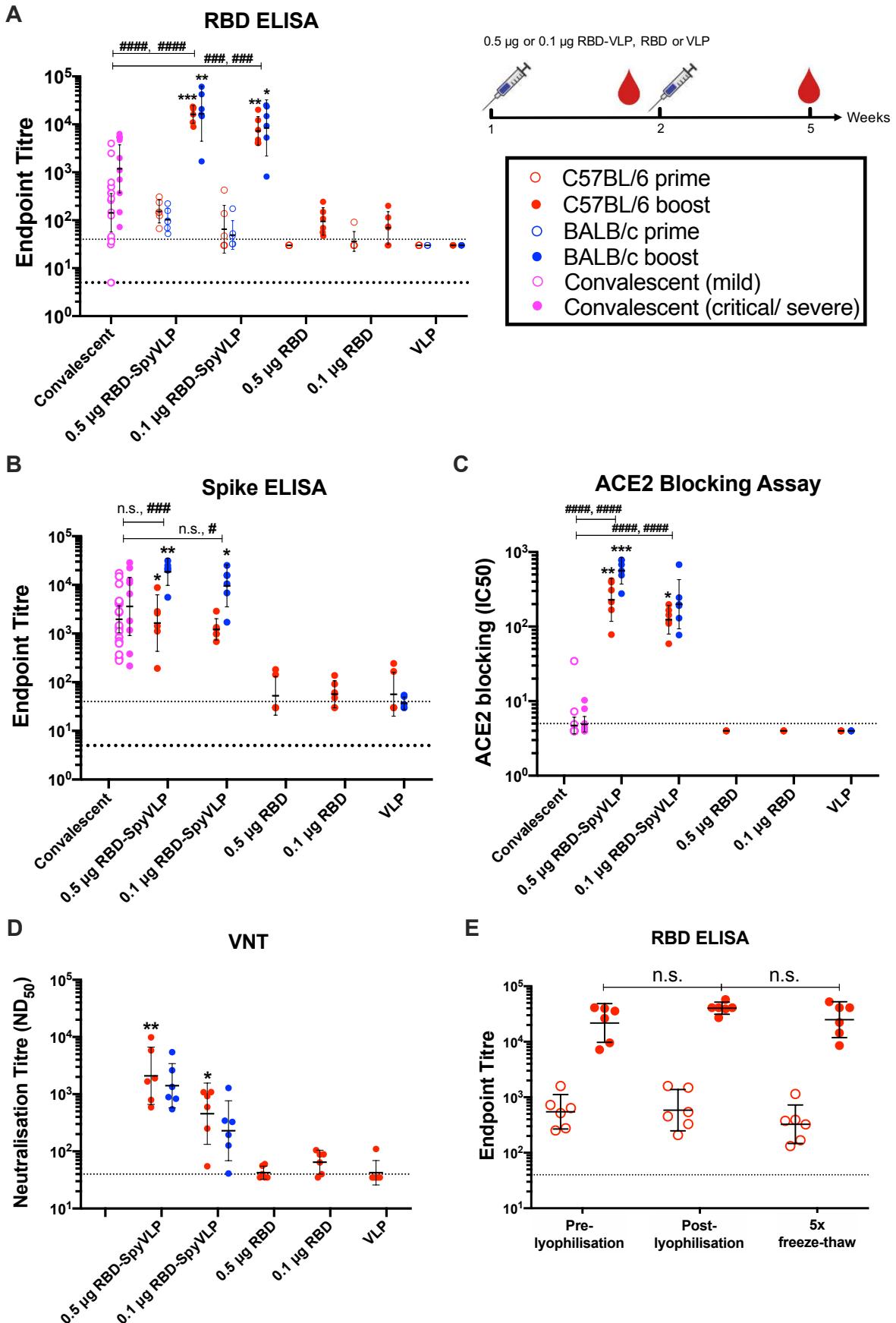
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1124 **Figure 1. SpyTag-RBD can be efficiently conjugated to SpyCatcher003-mi3 VLP.**
1125 (A) Schematic diagram of the RBD-SpyVLP vaccine candidate, consisting of
1126 SpyCatcher003-VLP conjugated with SpyTag-RBD. The isopeptide bonds formed
1127 spontaneously between SpyTag and SpyCatcher are indicated with red dots. (B)
1128 Conjugation of SpyCatcher003-mi3 with SpyTag-RBD at various ratios. Reactions
1129 were performed at 4 °C overnight and analysed using SDS-PAGE with Coomassie
1130 staining and densitometry, with the percentage of unreacted VLP shown. (C) Dynamic
1131 light scattering (DLS) characterisation of SpyTag-RBD, SpyVLP, and conjugated
1132 RBD-SpyVLP (n=3, values shown as mean±SD). R_H = hydrodynamic radius.

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1135 **Figure 2. RBD-SpyVLPs are reactive to SARS-CoV-2 binders and thermostable**
1136 **and resilient.** (A) Binding of RBD-SpyVLP to a panel of monoclonal antibodies
1137 isolated from COVID-19 recovered patients that target independent epitopes on the
1138 RBD determined using competitive ELISA (Huang et al., 2020). The boxed antibodies
1139 form groups that compete with each other and with antibodies or nanobodies with
1140 structurally defined footprints: CR3022 (PDB 6W41) (footprint 1), H11-D4-Fc (footprint
1141 2) (PDB 6YZ5), S309 (footprint 3) (PDB 6WPT) and the ACE2 binding site (PDB 6M0J).
1142 Each point represents the mean of duplicate readings and error bars represent ± 1 SD.
1143 The diagram of the RBD, created in PyMOL, shows the ACE2 binding site and the
1144 three binding footprints highlighted. (B) Solubility and (C) immunoreactivity of RBD-
1145 SpyVLP after storage for two weeks at various temperatures, determined using SDS-
1146 PAGE and densitometry or ELISA (10 μ g/mL CR3022 mAb). (D) Solubility and (E)
1147 immunoreactivity of RBD-VLP after freeze-thaw determined using SDS-PAGE and
1148 ELISA (10 μ g/mL CR3022 mAb) after one to five cycles of freeze-thawing. (F) RBD-
1149 SpyVLP soluble fraction, before and after lyophilisation reconstituted in the same
1150 buffer volume and (G) immunoreactivity determined using ELISA with ACE2-Fc and
1151 mAbs that target non-overlapping epitopes on the RBD. Error bars in B, E & F
1152 represent group mean ($n=3$). Error bars in C, E & G represent mean ± 1 SD ($n= 3$).
1153 Statistical difference in B to E was determined using Kruskal-Wallis test followed by
1154 Dunn's multiple comparison test. Statistical difference in F & G was determined using
1155 Mann-Whitney U test. n.s. = not significant.
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1160 **Figure 3. RBD-SpyVLPs induce strong antibody responses in mice that are**
1161 **comparable to the responses in recovered patients.** C57BL/6 (red) or BALB/c
1162 (blue) mice (n=6 in each group) were dosed twice IM, two weeks apart with 0.1 µg or
1163 0.5 µg purified RBD, RBD-SpyVLP or VLP alone with AddaVax adjuvant added to all.
1164 Sera were harvested at two weeks after the first dose (open circles) and at three weeks
1165 after the second dose (closed circles). Sera were analysed in (A) ELISA against RBD
1166 (B) ELISA against full-length spike glycoprotein, (C) in an ACE2 competition assay,
1167 and (D) in virus neutralisation assays (VNT) against wild-type SARS-CoV-2 virus. (E)
1168 Antibody response analysed by RBD ELISA for mice dosed twice with 0.5 µg RBD-
1169 SpyVLP (pre-lyophilised, post-lyophilised or freeze-thawed 5 times (5x FT). Data are
1170 presented as the group geometric means \pm 95% confidence intervals. COVID-19
1171 convalescent plasma from humans with mild (open mauve circles) or critical/severe
1172 (closed mauve circles) disease were included for comparison. *p<0.05, ** p<0.01,
1173 ***p<0.001 determined by Kruskal-Wallis test followed by Dunn's multiple comparison
1174 test. #p<0.05, ###p<0.001, #####p<0.0001 determined by Mann-Whitney U test to
1175 compare against convalescent human plasma. Dotted lines represent the lowest
1176 mouse sera dilutions tested. Bold dotted line represents the lowest human sera dilution
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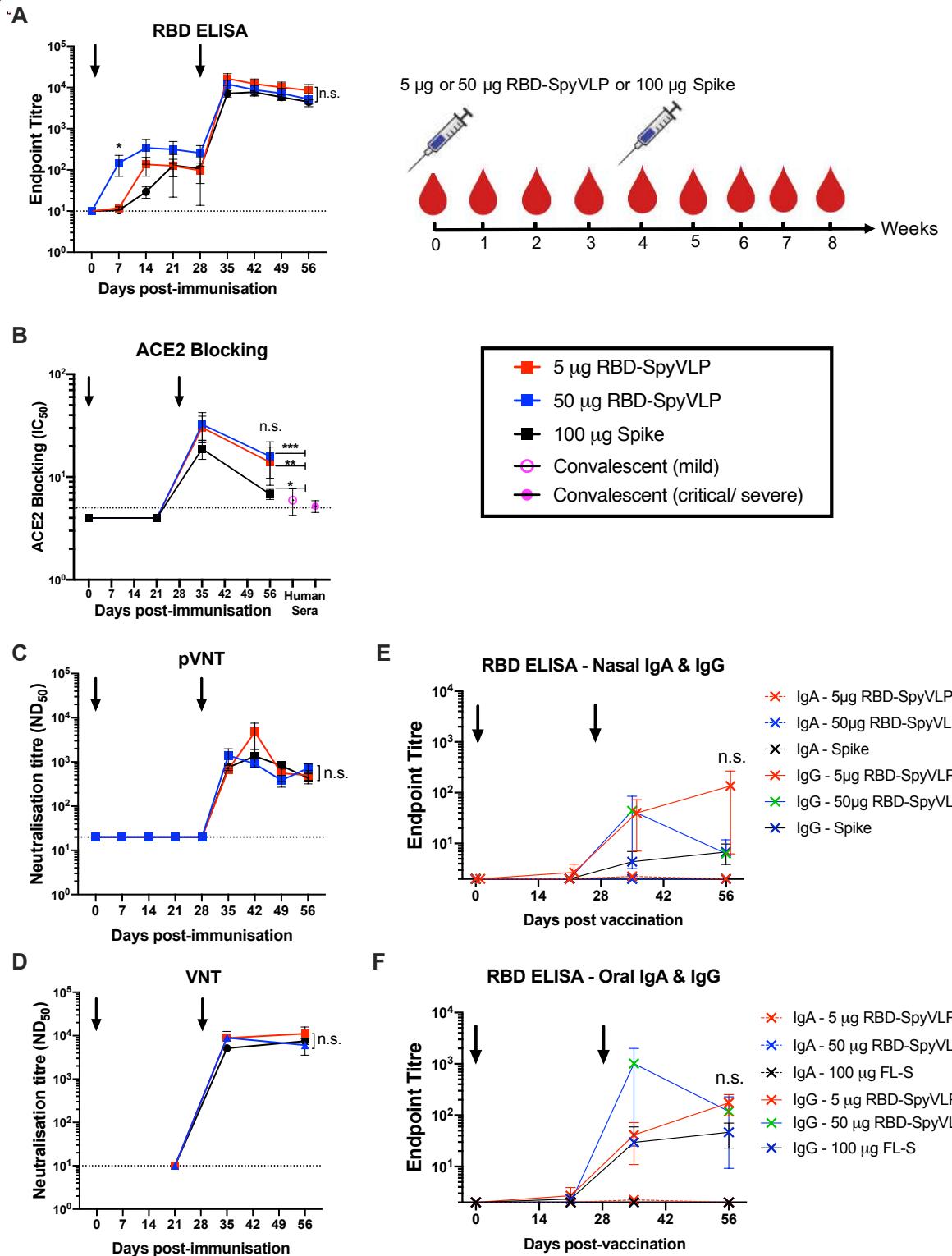
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1193 **Figure 4. RBD-SpyVLPs induce persistent and strong neutralising antibody**

1194 **responses in pigs.** Pigs (n=3 in each group) were dosed twice (prime and boost) IM,

1195 a month apart with 5 µg or 50 µg RBD-VLP or 100 µg of spike glycoprotein, all

1196 adjuvanted with AddaVax and sera were harvested at indicated time-points and

1197 analysed in (A) ELISA against RBD, (B) in an ACE2 competition assay, (C) in

1198 pseudovirus neutralisation assay (pVNT), and (D) wild-type virus neutralisation assay

1199 (VNT). Antibody levels in (E) nasal and (F) oral swabs were measured at indicated

1200 time points using RBD ELISA. Data are presented as the group mean ± 1 SEM. Black

1201 arrows indicate when the vaccines were administered. Kruskal-Wallis test was used

1202 to compare data on day 56 in B-E, n.s.= not significant. *p<0.05, **p<0.01 and

1203 ***p<0.001 on day 56 data compared to convalescent human plasma determined by

1204 Mann-Whitney U Test. Dotted lines represent the lowest dilutions of sera tested in the

1205 assays.

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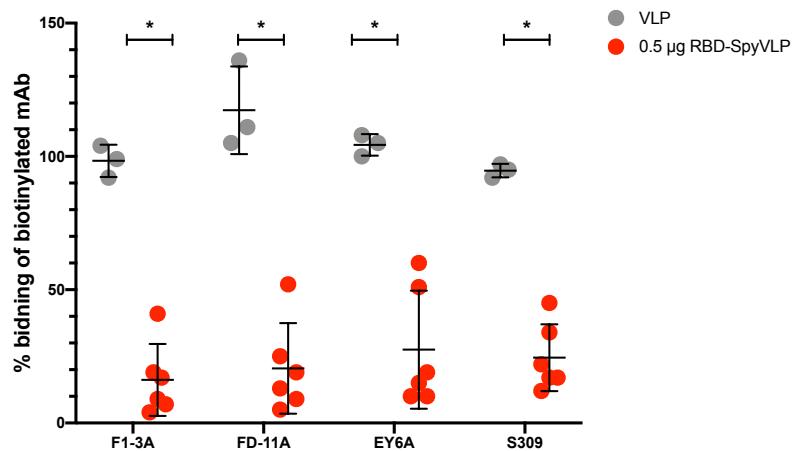
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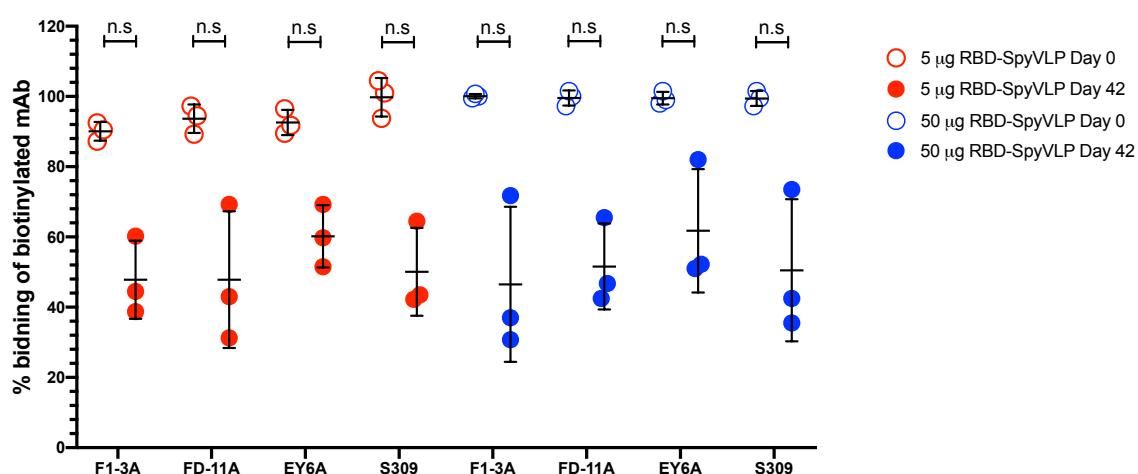
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1215 **Figure 5. RBD-SpyVLP immunisation of mice and pigs elicits polyclonal**
1216 **antibody responses that targets all key epitopes on the RBD.** Competition ELISA
1217 of four human mAbs that target three different key epitopes on the RBD with post-
1218 boost BALB/c sera (0.5 µg RBD-SpyVLP) (A). Competition ELISA of four human mAbs
1219 that target three different key footprints on the RBD with post-boost pig sera (5 µg or
1220 50 µg RBD-SpyVLP) compared to preimmune sera (day 0) (B). Each point in A
1221 represents an average of duplicate readings of a serum sample from one animal three
1222 weeks post boost tested at 1:20 dilution. Each point in B represents an average of
1223 quadruplicate readings of a serum sample from one animal on day 42 tested at 1:20
1224 dilution. Data are presented as group means ± 1 SD. ** p<0.01, determined by Mann-
1225 Whitney U test. n.s. = not significant.

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