- 1 **Title:** Virus-like particle displaying SARS-CoV-2 receptor binding domain elicits neutralizing
- 2 antibodies and is protective in a challenge model
- 3 Author names and affiliations: Julia L. McKechnie<sup>a</sup>\*, Brooke Fiala<sup>a</sup>\*, Clancey Wolf<sup>a</sup>, Daniel
- 4 Ellis<sup>a</sup>, Douglas Holtzman<sup>a</sup>, Andrew Feldhaus<sup>a</sup>
- 5 \*These authors have contributed equally to this work
- 6 <sup>a</sup>Icosavax, Inc.
- 7 1930 Boren Ave, Suite 1000
- 8 Seattle, WA 98101, USA
- 9 Abstract

While the effort to vaccinate people against severe acute respiratory syndrome coronavirus 2 10 (SARS-CoV-2) has largely been successful, particularly in the developed world, the rise of new 11 12 variants as well as waning immunity illustrate the need for a new generation of vaccines that 13 provide broader and/or more durable protection against infection and severe disease. Here we 14 describe the generation and characterization of IVX-411, a computationally designed, two-15 component virus-like particle (VLP) displaying the ancestral SARS-CoV-2 receptor binding domain (RBD) on its surface. Immunization of mice with IVX-411 generates neutralizing 16 antibodies against the ancestral strain as well as three variants of concern. Neutralizing antibody 17 18 titers elicited by IVX-411 are durable and significantly higher than those elicited by immunization with soluble RBD and spike antigens. Furthermore, immunization with IVX-411 19 is shown to be protective in a Syrian Golden hamster challenge model using two different strains 20 21 of SARS-CoV-2. Overall, these studies demonstrate that IVX-411 is highly immunogenic and 22 capable of eliciting broad, protective immunity.

Keywords: SARS-CoV-2 vaccine, virus-like particles, protein nanoparticle, receptor binding
 domain, neutralizing antibody response, long-lived plasma cells

Abbreviations: COVID-19, coronavirus disease 2019; SARS-CoV-2, severe acute respiratory
syndrome coronavirus 2; VOCs, variants of concern; S, spike protein; RBD, receptor binding
domain; ACE2, angiotensin-converting enzyme 2; VLPs, virus-like particles; LLPCs, long-lived
plasma cells; IM, intramuscular; SEC, size exclusion chromatography; DLS, dynamic light
scattering; BLI, biolayer interferometry; nsTEM, negative stain transmission electron
microscopy; PNA, pseudo-particle neutralization assay

## 31 Background

32 The coronavirus disease 2019 (COVID-19) pandemic has led to over 637 million 33 confirmed cases and 6.6 million deaths worldwide as of November 29, 2022 [1]. Our 34 understanding of this disease as well as the causative virus, SARS-CoV-2, has grown 35 dramatically since December, 2019 when it emerged in Wuhan, China [2,3]. Acute SARS-CoV-2 36 infection is characterized by symptoms such as fever, cough, and sore throat. The virus can also 37 lead to a potentially debilitating condition called 'long-haul COVID' [4]. Additionally, the emergence of variants of concern (VOCs) has illustrated the ability of SARS-CoV-2 to evolve, 38 39 leading to increased transmissibility and immune evasion. Given continued viral evolution and 40 transmission, further advances in SARS-CoV-2 vaccination approaches are necessary to stem the public health impact. 41

The SARS-CoV-2 spike (S) protein has been the primary target for vaccine development due to the ability of anti-S protein antibodies to neutralize the virus and protect against severe disease [5–9]. S protein trimers decorate the surface of the virion and are comprised of two subunits: S1 and S2 [10,11]. S1 sits at the apex of the S protein and contains the receptor binding 46 domain (RBD). The RBD binds to angiotensin-converting enzyme 2 (ACE2), a protein expressed 47 in human airway epithelia as well as lung parenchyma [12]. After S1 binds ACE2, the S2 subunit facilitates viral fusion with the host cell. During this process, the S protein trimer undergoes a 48 49 structural transition between its prefusion conformation and its post-fusion conformation, bringing together the fusion peptide and the host cell membrane, mediating virion entry. 50 51 Antibodies which block S1 binding to ACE2 are often potently neutralizing. Consequently, it is 52 unsurprising that the RBD is the primary target of neutralizing antibodies in human serum 53 [13,14].

54 Rapid development of vaccines against SARS-CoV-2 was facilitated by two vaccine 55 platform technologies: messenger RNA (mRNA) and viral vectors. While the deployment of 56 these vaccines was instrumental in slowing viral transmission and saving lives, the associated 57 reactogenicity and durability of these vaccines leave room for improvement. Furthermore, the 58 two viral vector vaccines, Ad26.COV2.S (Janssen) and ChAdOX1-S (AstraZeneca), have been 59 linked to a rare blood clotting disorder, thrombosis with thrombocytopenia syndrome [15]. 60 Concern over this potentially life-threatening side effect has led to limited use of these vaccines in developed countries. Waning antibody responses to vaccination, particularly with the mRNA-61 62 based vaccines, have also been described [16]. While two doses of the mRNA-based vaccines, 63 BNT162b2 (Pfizer) and mRNA1273 (Moderna), were roughly 90% efficacious in preventing 64 COVID-19 up to two months post immunization, waning antibody responses have resulted in 65 recommendations to receive multiple additional doses [17–20]. Critically, these first generation 66 vaccines are also less efficacious against the omicron variant, which has led to an increase in 67 breakthrough infections [21–23]. Considering these limitations, a new generation of SARS-CoV-68 2 vaccines that generate broad, durable immunity with reduced reactogenicity are needed.

69 Virus-like particles (VLPs) are a compelling technology for developing new SARS-CoV-70 2 vaccines. Whereas soluble recombinant viral proteins tend to be poorly immunogenic, 71 particularly in the absence of an adjuvant, VLPs can induce both humoral and cellular immune 72 responses even without adjuvants [24,25]. One advantage of VLPs is the multivalent presentation 73 of antigen, which promotes B cell receptor clustering and activation, facilitating the production 74 of high affinity antibodies [26]. Importantly, VLP-based vaccines displaying native viral 75 antigens are already commercially available for the prevention of hepatitis B (HBV) and human 76 papillomavirus (HPV) infection. These vaccines have excellent safety and durability profiles 77 [27,28]. Vaccination with HBV vaccines generates high antibody titers that are protective for up 78 to 30 years [29]. Similarly, a single dose of the bivalent HPV vaccine elicits antibody titers that 79 are maintained for at least seven years [30].

80 Recent advances in computational protein design have allowed for the generation of 81 novel, self-assembling VLPs that can display diverse antigens [24,25,31–36]. Here we produced 82 and further characterized a two-component, computationally designed VLP displaying 60 copies 83 of the ancestral SARS-CoV-2 RBD protein [24], referred to here as IVX-411. We show that 84 immunization of naïve animals with IVX-411 elicits high neutralizing antibody titers against the 85 ancestral strain as well as three VOCs. To evaluate the durability of this immune response, as 86 well as the benefit of the VLP platform over soluble protein, we immunized mice with IVX-411 87 or soluble S protein. We found that immunization with IVX-411 generated higher neutralizing titers and increased antigen-specific long-lived plasma cells (LLPCs) compared to immunization 88 89 with soluble protein. Finally, Syrian Golden hamsters immunized with IVX-411 and challenged 90 with SARS-CoV-2 had lower viral loads and reduced disease severity than unimmunized 91 hamsters.

## 92 Methods

Component production and characterization: RBD-CompA gene based on previously described 93 amino acid sequence [24] was synthesized and cloned by Genscript in the pcDNA3.4+ vector. 94 95 DNA was transiently transfected into HEK293F cells, which were incubated at 36 °C with 150 rpm shaking for 4 days before harvest by centrifugation, and 0.2  $\mu$ m filtration. Ni<sup>2+</sup> resin (Indigo, 96 97 Cube Biotech, #75110) was added to 4 µL/mL of cellular supernatant following addition of 1 M 98 Tris pH 8.0 to 50 mM and 5 M NaCl to 300 mM, and incubated with gentle rocking for 2 hours at room temperature (RT) or 16 hours at 4 °C. The loaded Ni<sup>2+</sup> resin was applied to gravity 99 columns. The columns were washed with 5 column volumes (CV) of wash buffer (20 mM Tris 100 101 pH 8.0, 300 mM NaCl, 30 mM imidazole, 0.75% CHAPS). Proteins were eluted with elution 102 buffer (20 mM Tris pH 8.0, 300 mM NaCl, 500 mM imidazole, 0.75% CHAPS) and dialyzed 103 into 20 mM Tris pH 8.0, 250 mM NaCl, 5% glycerol, 0.75% CHAPS buffer 3X. The purified 104 RBD-CompA was analyzed by SDS-PAGE and UV-Vis (Supplemental Figure 1), and 105 endotoxin levels were determined (Endosafe nexgen-PTS, Charles River; passing value = <10106 EU/mg). For manufacture of the CompB pentamer, a transformed E. coli Master Cell Bank was 107 expanded into a stirred-tank bioreactor for fed-batch production. CompB was purified using a 108 two-column chromatography process and final formulation conducted by tangential flow 109 filtration. Following purification and formulation, CompB was 0.2 mm filtered and stored at <-110 65°C.

111 <u>VLP production:</u> RBD01-CompA and CompB were quantified by UV-Vis prior to mixing in a 1.2X over equimolar ratio. RBD-CompA was added to a tube to a final concentration of 10  $\mu$ M, 113 then 20 mM Tris pH 8.0, 250 mM NaCl, 5% glycerol, 0.75% CHAPS buffer was added to bring 114 the reaction volume up to 1 mL. CompB was added to a final concentration of 8  $\mu$ M. The reaction was mixed and incubated for 1 hour. The resulting VLP was purified by SEC (Superose
6 Increase, Cytiva, #29091596), eluting around 11 mL, using 20 mM Tris pH 8.0, 250 mM NaCl,
5% glycerol, 0.75% CHAPS buffer as the mobile phase. Peak fractions centered around 11 mL
were pooled and filter-sterilized (0.2 µm) prior to analysis. VLP concentrations were quantified
by UV-Vis.

<u>UV-Vis spectroscopy:</u> All protein samples were analyzed by UV-Vis on an Agilent Cary 60.
Wavelength scans from 400 to 200 nm were collected, with baseline correction using a matching
buffer blank. Absorbance at 280 nm was used to quantify the protein concentrations with the
molar extinction coefficients and molecular weights as in the following formula:

$$c\left(\frac{mg}{mL}\right) = \frac{A280}{Ec} \cdot MW$$

<u>Dynamic Light Scattering:</u> DLS measurements were taken on a nanoDSF instrument (UNcle,
 UNchained Laboratories), using autoattenuation of the laser and collecting 10 acquisitions of 5
 seconds each at 20 °C.

127 Size Exclusion Chromatography: A Superose 6 Increase 10/300 GL column (Cytiva, #29091596) 128 was used to purify trimeric RBD-CompA component, IVX-411 in vitro assembly reaction (VLP 129 purification), or IVX-411 purified VLP (analytical), on an AKTA FPLC system (Cytiva, AKTA 130 Go). Columns were equilibrated using 1.2 CV of SEC purification buffer (20 mM Tris pH 8.0, 131 250 mM NaCl, 100 mM L-Arginine, 5% glycerol, 0.75% CHAPS), then 0.5-2 mL of sample was 132 injected onto the column and eluted using 1.2 CV of SEC purification buffer. 133 Biolayer interferometry: Purified RBD-CompA trimeric component, IVX-411 VLP, ACE2-Fc 134 dimerized receptor, and monoclonal antibodies (mAbs) (CR3022, COVA2-39, and CV07-270)

were diluted to 10  $\mu$ g/mL in BLI assay buffer (PBS pH 7.4, 0.5% BSA, 0.05% Tween 20). 200

136  $\mu$ L of each dilution and BLI assay buffer were added to black 96-well microplates. Protein A

biosensors (Sartorius, #18-5010) were hydrated in BLI assay buffer for 10-20 minutes and
loaded onto an Octet Red96 BLI instrument (Pall, FortéBio). Biosensors were dipped into BLI
assay buffer to obtain a baseline (60 s), loaded with dimerized ACE2-Fc or mAbs (120 s), dipped
into BLI assay buffer, transferred to RBD-CompA and IVX-411 wells (150 s), and dipped back
into BLI assay buffer (150 s).

142 Negative Stain Transmission Electron Microscopy: IVX-411 sample was diluted to 75 µg/mL in 143 SEC buffer. Sample was adhered to a thick-carbon/Formvar copper 400 mesh grid (Electron 144 Microscopy Sciences, #CF400-Cu-TH) by pipetting 6 µL of sample directly onto the carbon side 145 of the grid and incubating for 1 minute. The grid was dipped into a 50 µL droplet of sterile 146 filtered DI water followed by blotting with grade 1 filter paper (Whatman, #Z240079). The grid 147 was stained by dipping into a 6 µL drop of 0.75% uranyl formate stain, incubated for 1 minute, 148 and blotted off. The staining step was repeated, and the grid dried for 1 minute prior to storage. 149 A Talos L120C TEM microscope, Leginon software, and Gatan camera were used to image the 150 sample.

151 <u>VLP Prime-Boost study:</u> The in-life portion of this study was conducted at Abcore Inc. Female, 152 BALB/c mice were immunized IM on days 0 and 21 with 0.2 mg of IVX-411 or IVX-411 + 153 MF59. Immunizations, once prepared at room temperature, were used immediately or within two 154 hours of preparation. Serum samples from each animal were collected on Day 0 (prior to 155 immunization), Day 21 (prior to boost), and on Day 35 (terminal bleed).

<u>VLP versus soluble protein duration study</u>: The in-life portion of this study was conducted at
Aragen Bioscience Inc. BALB/c mice were immunized on days 0 and 21 with 0.2 mg of IVX411, RBD-CompA, or S-2P formulated with or without MF59. Serum samples were collected on

days 0, 20, 35, 63, 91, 119, and 154. Animals were sacrificed on Day 154. Bone marrow wascollected for ELISPOT analysis of LLPCs.

ELISpot: The Mouse IgG ELISpot<sup>BASIC</sup> kit, Protocol II (Mabtech, #3825-2A) was used by 161 162 Aragen Bioscience Inc to perform LLPC quantification. SARS-CoV-2 antigen RBD01-dn5B was 163 diluted to 80 mg/mL and used to coat wells of PVDF plates (Mabtech, #3654-WP-10) according 164 to the assay kit protocol. Anti-mouse IgG was diluted to 20 mg/ mL and was used to coat 165 additional wells on the same plate. Plates were washed and blocked with assay media for 30 minutes at RT. Cells isolated from femoral bone marrow were resuspended at  $1 \times 10^6$  cells/mL. 166 100 mL was added to the coated wells. Plates were incubated at 33°C for 14-16 hours. Spots 167 counts/well were determined using a Zellnet Consulting ELISpot reader. 168

169 Syrian Golden hamster efficacy study: The in-life portion of this study was conducted at 170 Lovelace Biomedical. 36 male, SGHs were purchased from Charles River. 16 hamsters were 171 immunized IM with 0.2 mg IVX-411 formulated with MF59 on days 0 and 21. Remaining 172 hamsters were immunized with PBS. Serum samples were collected on days 0, 21, and 35. Body 173 weights were measured starting on Day 39. On Day 42 8 IVX-411 immunized hamsters and 8 PBS immunized hamsters were challenged via intranasal instillation with 4.64 x  $10^5$ 174 175 TCID<sub>50</sub>/animal of WA1/2020; 8 IVX-411 immunized hamsters and 8 PBS immunized hamsters were challenged with  $1.53 \times 10^6$  TCID<sub>50</sub>/animal of B.1.617.2. Animals were sacrificed on Day 176 177 47 and lung weights were measured. Lungs were fixed in NBF, trimmed, paraffin embedded, 178 sectioned at 4 mm, and stained with hematoxylin and eosin for microscopic examination. Histopathologic findings were graded subjectively on a scale of 1 to 5. The Provantis<sup>TM</sup> (Instem 179 180 LSS Ltd., Staffordshire, England) computer software was used for necropsy and histopathology 181 data acquisition, reporting, and analysis.

182	RT-qPCR: RT-qPCR analysis on lung and nasal swab samples collected from the Syrian Golden
183	hamster study were performed at Lovelace Biomedical. Samples were processed in Trizol using
184	a TissueLyser and centrifuged at 4000 x g for 5 minutes. RNA extraction was performed on
185	supernatants using the QIAGEN RNeasy kit according to the manufacturer's instructions.
186	Samples were run in triplicate and genome copies per mL or gram equivalents were calculated
187	from a standard curve generated from RNA standards of known copy concentration. The N and E
188	gene primers and probe sequences were as follows:
189	N gene:
190	SARS-CoV-2 Forward: 5' TTACAAACATTGGCCGCAAA 3'
191	SARS-CoV-2 Reverse: 5' GCGCGACATTCCGAAGAA 3'
192	SARS-CoV-2 Probe: 6FAM-ACAATTTGCCCCCAGCGCTTCAG-BHQ-1
193	E gene:
194	SARS-CoV-2 Forward: 5' ACAGGTACGTTAATAGTTAATAGCGT 3'
195	SARS-CoV-2 Reverse: 5' ATATTGCAGCAGTACGCACACA 3'

196 SARS-CoV-2 Probe: 6FAM-ACACTAGCCATCCTTACTGCGCTTCG-BHQ-1

197 SARS-CoV-2 Pseudo-particle Neutralization Assay (PNA): PNA assays were performed at 198 Nexelis. ACE-2 expressing Vero E6 cells (ATCC CRL-1586) were seeded in a 96-well 199 microtiter plate at 20,000 cells/well. Serum samples and controls were heat-inactivated at 56°C 200 for 30 minutes, diluted in duplicate in cell medium, and serial two-fold dilutions were performed. Each SARS-CoV-2 pseudovirus was diluted to reach a desired concentration, added to the 201 202 diluted serum samples, and incubated at 37°C in 5% CO<sub>2</sub> for 1 hour. This mixture was added to the cells at 80% confluency. Plates were incubated for 18-22 hours at 37°C in 5% CO<sub>2</sub> before 203 204 supernatants were removed. 50 mL of ONE-Glo EX Luciferase Assay Substrate (Promega,

#E8110) diluted 1:2 in cell media was added to each well and incubated at RT for 3 minutes with agitation. Luminescence across all wavelengths was measured for 0.5 seconds using a SpectraMax iD3 microplate reader and SoftMax Pro v7.0.1 (Molecular Devices). A titration curve using a 4-parameter logistic regression was made for each dilution. The reciprocal dilution of the sample for which the luminescence was equal to a pre-defined cut-point of 50 was reported as the titer. The cut-point was determined using linear regression using 50% flanking points.

212 **Results** 

# 213 Production and characterization of two-component I53-50 VLPs displaying ancestral SARS-

214 <u>CoV-2 RBD</u>

215 A two-component, computationally designed protein VLP, referred to as I53-50 [35], was 216 utilized to display the ancestral (Wuhan-Hu-1) SARS-CoV-2 RBD to improve the 217 immunogenicity of the monomeric antigen as previously described [24]. I53-50 is a 120-subunit 218 VLP comprised of 20 homotrimeric (CompA) and 12 homopentameric (CompB) subunits, 219 capable of *in vitro* assembly following purification and subsequent mixing of the individual 220 components [35,37]. The his-tagged SARS-CoV-2 RBD antigen was displayed on I53-50 by 221 genetically fusing the C-terminus of the antigen to the N-terminus of the CompA subunit using a 222 16-residue glycine-serine linker (RBD-CompA) [24]. RBD-CompA and CompB were 223 individually expressed in HEK293F and E. coli cells, respectively, and purified prior to mixing 224 in equimolar ratios to induce spontaneous self-assembly of VLPs in vitro (Figure 1A). 225 Following purification by size exclusion chromatography (SEC) to remove residual components, 226 the resulting RBD-I53-50 VLPs (IVX-411) were characterized for identity, aggregation state, 227 antigenicity and VLP structural integrity. UV-Vis spectroscopy wavelength scan analysis

228 showed a peak at 280 nm and low-level scattering typical of non-aggregated, well-formed VLPs 229 (ratio of absorbance at 320 nm to 280 nm =  $\sim 0.1$ ) (Figure 1B). Dynamic light scattering (DLS) 230 measurements suggested non-aggregated, monodispersed VLPs with a polydispersity index of 231 11.8% (Figure 1C). Analytical SEC of purified IVX-411 resulted in a resolved peak centered 232 around 11 mL, consistent with the calculated molecular weight of the VLP (Figure 1D). IVX-233 411 eluted earlier than the constituent components (Figure 1D). The antigen appeared intact 234 based on binding to ACE2-Fc, as well as three additional anti-RBD monoclonal antibodies, as 235 measured by biolayer interferometry (BLI) (Figure 1E). Finally, negative stain transmission 236 electron microscopy (nsTEM) confirmed that the VLP sample consisted of monodispersed, intact 237 VLPs of the expected diameter (Figure 1F).

### 238 Animals primed and boosted with IVX-411 develop robust neutralizing titers against three VOCs

239 We evaluated the ability of IVX-411 to induce neutralizing antibody titers against the 240 ancestral SARS-CoV-2 strain as well as three variants of concern (beta, gamma, and delta). 241 Naïve BALB/c mice were immunized intramuscularly (IM) on Day 0 and Day 21 with 0.2 mg of 242 IVX-411 formulated with CSL Segirus' proprietary oil-in-water adjuvant, MF59 (Figure 2A). 243 Mice were bled on Day 0 (pre-immunization), Day 21 (pre-boost), and on Day 35 (14 days post-244 boost). Neutralizing titers against the ancestral strain and the beta variant were measured at all 245 time points using a cell-based pseudo-particle neutralization assay (PNA). Neutralizing titers 246 against gamma and delta variants were measured using only the Day 35 samples.

Three weeks after the initial priming dose, mice immunized with IVX-411 had an average Day 21 neutralizing antibody titer (NT50) of 2.1 x  $10^2$  against the ancestral strain (**Figure 2B**) and an average neutralizing antibody titer of 1.4 x  $10^2$  against the beta variant (**Figure 2C**). After receiving a booster immunization, the Day 35 neutralizing titers against the

251 ancestral strain increased 45-fold. Similarly, a 78-fold increase in neutralizing titers against the 252 beta variant was also observed. Mice primed and boosted with IVX-411 had an average neutralizing titer against the gamma variant of  $2.58 \times 10^4$  (Figure 2D). Neutralizing titers against 253 254 the delta variant were lower than those against gamma, beta, and the ancestral strain, with an average neutralizing titer of  $3.71 \times 10^3$  (Figure 2E). Day 35 neutralizing antibody titers against 255 256 all four strains were higher than those of control human convalescent sera. Together, these 257 results demonstrate that immunization with IVX-411 results in a broad, potent antibody 258 response.

# 259 <u>Immunization with IVX-411 induces a more potent humoral immune response than</u> 260 <u>immunization with soluble protein</u>

261 Having confirmed the immunogenicity of IVX-411, we next sought to evaluate its ability 262 to elicit durable, humoral immunity compared to trimeric spike-based and RBD-based soluble 263 antigens. Naïve BALB/c mice were immunized on Day 0 and boosted on Day 21 with either 0.2 264 mg of IVX-411, an equivalent antigen dose of RBD-CompA (0.15  $\mu$ g), or an equivalent RBD 265 antigen dose of S-2P (0.4  $\mu$ g), a stabilized prefusion version of the S protein ectodomain (Figure 266 **3A**). Each antigen was formulated with either MF59 or an aqueous buffer. Serum samples were 267 collected on days 0, 20, 35, 63, 91, 119, and 154. On Day 154 the animals were sacrificed, at 268 which point LLPCs were isolated from the bone marrow and assessed by ELISpot.

Serum analysis by a cell-based PNA revealed that immunization with IVX-411 generated higher neutralizing antibody titers than immunization with soluble proteins. On Day 20, after a single immunization, only animals immunized with IVX-411 had neutralizing antibody titers above the lower limit of quantitation (**Figure 3B**). Even at this early timepoint, MF59 enhanced the neutralizing antibody response in naïve animals with the IVX-411 MF59 group having 4.4274 fold higher Day 20 neutralizing antibody titers compared to animals immunized with IVX-411 275 alone. This trend was consistent throughout the duration of the study. Compared to the aqueous 276 formulations, immunization with antigen plus MF59 resulted in 16, 3.1, and 116-fold increases 277 in Day 35 neutralizing antibody titers for IVX-411, RBD-CompA, and S-2P, respectively. 278 Importantly, the Day 35 neutralizing antibody titers in the IVX-411 MF59 group were 198-fold 279 higher than those in the RBD-CompA MF59 group and 5.3-fold higher than those in the S-2P 280 MF59 group. These results suggest that immunization with IVX-411 results in higher 281 neutralizing antibody titers than immunization with soluble protein, particularly compared with 282 soluble trimerized RBD, which is poorly immunogenic on its own. There were no statistically 283 significant differences in the neutralizing antibody titers within the different treatment groups 284 from Day 35 to Day 154, except for the RBD-CompA MF59 group. Unlike the other groups, neutralizing antibody titers following a second administration of RBD-CompA MF59 did not 285 286 reach their peak until Day 154. The 7.9-fold increase in titers on Day 154 compared to Day 35 287 rose to the level of statistical significance. While the overall consistency of neutralizing antibody 288 titers post-boost reveals the durability of the humoral immune response elicited by immunization 289 in general, these results show that immunization with IVX-411 leads to higher neutralizing titers.

To further evaluate the ability of IVX-411 to generate a long-term humoral immune response, an ELISpot assay was performed on isolated LLPCs. Similar to the neutralizing antibody titer results, these results demonstrated adjuvantation was key to generating a sizable RBD-specific LLPC compartment in naïve animals. There were no statistically significant differences in the number of RBD-specific LLPCs between the aqueous conditions. (Figure 3C). Compared to the aqueous formulations, immunization with antigen plus MF59 resulted in 4.3, 1.8, and 4.4-fold increases in LLPC counts for IVX-411, RBD-CompA, and S-2P, respectively. Immunization with IVX-411 MF59 resulted in a statistically significant, 3.9-fold increase in the number of LLPCs compared to immunization with RBD-CompA MF59. The 1.2-fold increase in LLPC counts observed in the IVX-411 MF59 immunized group compared to the S-2P MF59 immunized group was modest and did not reach the level of statistical significance. These results demonstrate that immunization with IVX-411 in the presence of an oil-in-water emulsion induces durable neutralizing antibody titers, which along with LLPC counts, are superior to those induced by immunization with soluble antigens.

# 304 <u>Immunization with IVX-411 reduces disease severity and viral load in Syrian Golden hamster</u> 305 <u>model</u>

306 To evaluate whether IVX-411 could protect against SARS-CoV-2 infection, a viral 307 challenge study using WA1/2020 (ancestral) and B.1.671.2 (delta) strains was performed in 308 Syrian Golden hamsters. Hamsters were randomly assigned to five treatment groups: PBS, 309 Unchallenged; PBS, WA1/2020; PBS, B.1.617.2; IVX-411, WA1/2020; and IVX-411, B.1.671.2 310 (Figure 4A). On Day 0 and Day 21 hamsters in the IVX-411, WA1/2020, and IVX-411, 311 B.1.617.2 groups were immunized with IVX-411 formulated with MF59. Hamsters in the other 312 three treatment groups (PBS, Unchallenged; PBS, WA1/2020; and PBS, B.1.617.2), were 313 injected with PBS. Hamsters were weighed and bled on days 0, 21, and 35. On Day 42 all 314 hamsters, except those in the PBS, Unchallenged group, were inoculated intranasally with either 4.64 x 10<sup>5</sup> TCID<sub>50</sub>/animal of WA1/2020 or 1.53 x 10<sup>6</sup> TCID<sub>50</sub>/animal of B.1.617.2. Body 315 316 weights were recorded daily from Day 39 to Day 47, at which point animals were sacrificed and 317 lung tissue as well as nasal swabs were collected. Hamsters challenged with either WA1/2020 or 318 B.1.617.2 began to lose weight four to five days post challenge (Figure 4B). By six days post 319 challenge, hamsters immunized with IVX-411 had significantly higher body weights compared

to their unimmunized counterparts. Interestingly, this reduction in weight loss was more
pronounced in animals challenged with B.1.617.2 compared to WA1/2020, despite the higher
B.1.617.2 challenge dose. This continued to be the case until the termination of the study on Day
47.

324 As SARS-CoV-2-induced pneumonia develops, lung weight increases relative to body 325 weight due to cellular infiltrates, edema, and a decrease in overall body weight. Consequently, 326 we investigated the ability of immunization with IVX-411 to prevent an increase in lung weight. 327 As expected, hamsters in the PBS, WA1/2020 and PBS, B.1.617.2 groups had increased lung 328 weight compared to hamsters in the PBS, Unchallenged group (Figure 4C). However, this was 329 only statistically significant in animals challenged with B.1.617.2. Similarly, unimmunized 330 animals challenged with either WA1/2020 or B.1.617.2 had a 1.2 or 1.6-fold increase in lung 331 weight as a percent of body weight respectively compared to their IVX-411 immunized 332 counterparts. Immunization with IVX-411 led to a statistically significant reduction in lung 333 weight in the context of challenge with B.1.617.2 and a trend towards lower lung weights in 334 hamsters challenged with WA1/2020.

335 In addition to lung and body weight, the impact of IVX-411 immunization on 336 histopathologic changes associated with SARS-CoV-2 infection were also assessed (Figure 4D). 337 The two challenge strains led to slightly different tissue changes in unimmunized animals. 338 B1.617.2 infection led to more hemorrhage and less centriacinar and bronchiolar epithelial 339 hypertrophy/hyperplasia compared to WA1/2020. The impact of IVX-411 immunization on lung 340 pathology was the most notable in animals challenged with B1.617.2. Animals immunized with 341 IVX-411 and then challenged with B1.617.2 had milder inflammation, reduced pulmonary 342 hemorrhage, and decreased severity of bronchiolar epithelial hyperplasia/hypertrophy compared

to unimmunized animals. In the case of challenge with WA1/2020, immunization resulted in
more subtle decreases in the severity of inflammation, pulmonary hemorrhage, and centriacinar
hyperplasia/hypertrophy. The severity of bronchiolar epithelial hyperplasia/hypertrophy was
markedly improved when animals challenged with WA1/2020 had been previously immunized
with IVX-411.

348 To further assess the level of protection conferred by immunization with IVX-411, lung 349 tissue and nasal swabs were assessed for genomic RNA (N gene) and subgenomic RNA (E gene) 350 by RT-qPCR. Genomic RNA measures both viable and nonviable viral genomes whereas 351 subgenomic RNA is a measure of replicating virus levels. IVX-411 immunization significantly 352 reduced the number of N gene copies per gram of lung tissue 31 and 66-fold and the number of 353 N gene copies per nasal swab 4.4 and 3-fold in hamsters challenged with WA1/2020 and B.1.617.2, respectively (Figure 5A). Consistent with the N gene results, IVX-411 immunization 354 355 reduced the number of E gene copies per gram of lung tissue 60 and 132-fold and the number of 356 E gene copies per nasal swab 5 and 3-fold in hamsters challenged with WA1/2020 and 357 B.1.617.2, respectively (Figure 5B). However, the decreases in N and E gene copies in nasal 358 swab samples collected from immunized animals did not reach the level of statistical 359 significance.

Finally, to determine whether neutralizing antibodies elicited by IVX-411 vaccination were contributing to the observed protection, we evaluated the neutralizing antibody response elicited in the animals immunized with IVX-411 prior to challenge using a cell-based PNA. On Day 35, neutralizing antibody titers against both the ancestral and delta variant were present in six out of eight IVX-411, WA1/2020 hamsters and all eight IVX-411, B.1.617.2 hamsters (**Figure 5C**). Neutralizing titers against the ancestral strain increased approximately 14.5-fold in 366 IVX-411 immunized animals compared to baseline. A roughly 3.8-fold increase in neutralizing 367 titers against the delta variant was also observed. Overall, these results demonstrate that 368 prophylactic immunization with IVX-411 is protective against severe manifestations of SARS-369 CoV-2 infection in Syrian Golden hamsters.

370 **Discussion** 

371 As SARS-CoV-2 becomes endemic, there is great interest in identifying vaccines with 372 improved durability, less reactogenicity, and the potential for combination with other vaccines, 373 such as seasonal influenza. VLPs have been shown to be highly immunogenic and possess many 374 of the characteristics that may be of value for a next-generation SARS CoV-2 vaccine. Here, we 375 show that IVX-411, a computationally designed VLP presenting the RBD of the ancestral SARS-376 CoV-2 strain, is immunogenic in two rodent models. In mice, immunization with IVX-411 elicits 377 neutralizing titers against the ancestral strain as well as three VOCs. These neutralizing antibody 378 titers are durable and higher than those generated in response to immunization with a soluble 379 spike protein. Critically, immunization with IVX-411 is protective in a Syrian Golden hamster 380 challenge model. Together, these results further illustrate the potential of VLPs to be utilized as a 381 modality for future SARS-CoV-2 vaccines.

These data for IVX-411 are consistent with a large number of preclinical studies on I53-50 immunogens displaying RBD antigens from SARS-CoV-2 and/or other sarbecoviruses [24,31,38], and further demonstrate the precision and immunogenic potency of this platform. In naïve animals, higher neutralizing titers against SARS-CoV-2 were observed after immunization with I53-50 VLPs displaying 60 copies of the RBD compared to soluble S protein trimers and trimerized RBD antigens [24,39]. Interestingly, we observed IVX-411 generated higher neutralizing antibody titers in naïve mice than in Syrian Golden hamsters. However, differences 389 in neutralizing antibody titers between different animal models are consistent with what others 390 have reported [32,40]. Further, we observed improved potency in naïve animals by formulating 391 IVX-411 with MF59, an oil-in-water emulsion. Such results have been externally replicated for 392 I53-50-based SARS-CoV-2 vaccines in naïve human patients [38,41–43]. Our results are 393 immunogenically similar to those observed from other RBD-based, single-component, SARS-394 CoV-2 VLP vaccines, which use nanoparticles such as ferritin [44,45], hepatitis B surface 395 antigen [46], or mi3 (a variant of the I3-01 VLP) [47–50], as well as biochemical conjugation 396 methods such as SpyCatcher and sortase [45,47]. In comparison to these single-component VLP 397 platforms, the two-component nature of I53-50 allows for simplified and modular manufacturing 398 of highly defined immunogen structures through *in vitro* assembly. The precise structure of the 399 final VLP product is enabled both by direct genetic fusion between antigens and CompA, 400 eliminating the need for conjugation, as well as dependable complete and cooperative assembly 401 of the nanoparticle structure due to highly-specific designed interactions between CompA and 402 CompB [35,37]. Finally, the manufacturing and clinical validation of I53-50-based vaccines for 403 SARS-CoV-2 [43] and RSV (Icosavax, Inc. unpublished data) confirms that this platform is 404 scalable, manufacturable, and potently immunogenic while maintaining low reactogenicity, with 405 ample potential to extend to other viral and bacterial indications.

The emergence of the omicron SARS-CoV-2 variant in November, 2021 led to a surge of infections in December, 2021 through March, 2022 [1]. Given the ability of the omicron variant to escape preexisting immunity elicited by both natural infection and immunization, updated booster vaccines targeting the omicron variant as well as the ancestral strain are now recommended for those 12 and older. The I53-50 VLP platform utilized in IVX-411 has the potential to combine multiple SARS-CoV-2 antigens to direct immune responses against several VOCs, including omicron. This platform also has the potency and low reactogenicity needed to generate combination vaccines which could include additional VLP-based vaccines against other viruses. Prior experience with omicron suggests that future updates to SARS-CoV-2 vaccines may be necessary as well as routine booster immunizations. The modular ability for antigen presentation on I53-50 VLPs in response to emerging VOCs, combined with the potential use of high-yield, stabilized RBD designs [39], can enable this platform to meet many needs for future COVID-19 vaccination.

# 419 Declaration of Competing Interest

420 All authors are employees and stockholders of Icosavax, Inc.

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- 429 **References**
- 430 [1] WHO Coronavirus (COVID-19) dashboard n.d. https://covid19.who.int (accessed July 6,
  431 2022).
- 432 [2] Zhou P, Yang X-L, Wang X-G, Hu B, Zhang L, Zhang W, et al. A pneumonia outbreak
  433 associated with a new coronavirus of probable bat origin. Nature 2020;579:270–3.

- 434 [3] Zhu N, Zhang D, Wang W, Li X, Yang B, Song J, et al. A novel Coronavirus from patients
  435 with pneumonia in China, 2019. N Engl J Med 2020;382:727–33.
- 436 [4] Mehandru S, Merad M. Pathological sequelae of long-haul COVID. Nat Immunol
  437 2022;23:194–202.
- 438 [5] Rogers TF, Zhao F, Huang D, Beutler N, Burns A, He W-T, et al. Isolation of potent SARS-
- 439 CoV-2 neutralizing antibodies and protection from disease in a small animal model. Science
  440 2020;369:956–63.
- 441 [6] Zost SJ, Gilchuk P, Case JB, Binshtein E, Chen RE, Nkolola JP, et al. Potently neutralizing
  442 and protective human antibodies against SARS-CoV-2. Nature 2020;584:443–9.
- Liu L, Wang P, Nair MS, Yu J, Rapp M, Wang Q, et al. Potent neutralizing antibodies
  against multiple epitopes on SARS-CoV-2 spike. Nature 2020;584:450–6.
- [8] Brouwer PJM, Caniels TG, van der Straten K, Snitselaar JL, Aldon Y, Bangaru S, et al.
  Potent neutralizing antibodies from COVID-19 patients define multiple targets of
  vulnerability. Science 2020;369:643–50.
- 448 [9] Cao Y, Su B, Guo X, Sun W, Deng Y, Bao L, et al. Potent neutralizing antibodies against
- 449 SARS-CoV-2 identified by high-throughput single-cell sequencing of convalescent patients'
- 450 B cells. Cell 2020;182:73-84.e16.
- [10] Ke Z, Oton J, Qu K, Cortese M, Zila V, McKeane L, et al. Structures and distributions of
  SARS-CoV-2 spike proteins on intact virions. Nature 2020;588:498–502.
- 453 [11] Huang Y, Yang C, Xu X-F, Xu W, Liu S-W. Structural and functional properties of SARS-
- 454 CoV-2 spike protein: potential antivirus drug development for COVID-19. Acta Pharmacol
  455 Sin 2020;41:1141–9.

- 456 [12] Jia HP, Look DC, Shi L, Hickey M, Pewe L, Netland J, et al. ACE2 receptor expression and
- 457 severe acute respiratory syndrome coronavirus infection depend on differentiation of human
  458 airway epithelia. J Virol 2005;79:14614–21.
- 459 [13] Steffen TL, Stone ET, Hassert M, Geerling E, Grimberg BT, Espino AM, et al. The receptor
- binding domain of SARS-CoV-2 spike is the key target of neutralizing antibody in human
- 461 polyclonal sera. BioRxiv 2020. https://doi.org/10.1101/2020.08.21.261727.
- 462 [14] Piccoli L, Park Y-J, Tortorici MA, Czudnochowski N, Walls AC, Beltramello M, et al.
  463 Mapping neutralizing and immunodominant sites on the SARS-CoV-2 spike receptor464 binding domain by structure-guided high-resolution serology. Cell 2020;183:1024465 1042.e21.
- 466 [15] Lai C-C, Ko W-C, Chen C-J, Chen P-Y, Huang Y-C, Lee P-I, et al. COVID-19 vaccines
  467 and thrombosis with thrombocytopenia syndrome. Expert Rev Vaccines 2021;20:1027–35.
- 468 [16] Ferdinands JM, Rao S, Dixon BE, Mitchell PK, DeSilva MB, Irving SA, et al. Waning 2-
- dose and 3-dose effectiveness of mRNA vaccines against COVID-19-associated emergency
- 470 department and urgent care encounters and hospitalizations among adults during periods of
- 471 Delta and Omicron variant predominance VISION Network, 10 states, August 2021-
- 472 January 2022. MMWR Morb Mortal Wkly Rep 2022;71:255–63.
- 473 [17] Moreira ED Jr, Kitchin N, Xu X, Dychter SS, Lockhart S, Gurtman A, et al. Safety and
  474 efficacy of a third dose of BNT162b2 Covid-19 vaccine. N Engl J Med 2022;386:1910–21.
- 475 [18] Menni C, May A, Polidori L, Louca P, Wolf J, Capdevila J, et al. COVID-19 vaccine
- 476 waning and effectiveness and side-effects of boosters: a prospective community study from
- 477 the ZOE COVID Study. Lancet Infect Dis 2022;22:1002–10.

- 478 [19] Barda N, Dagan N, Cohen C, Hernán MA, Lipsitch M, Kohane IS, et al. Effectiveness of a
- third dose of the BNT162b2 mRNA COVID-19 vaccine for preventing severe outcomes in
  Israel: an observational study. Lancet 2021;398:2093–100.
- 481 [20] Magen O, Waxman JG, Makov-Assif M, Vered R, Dicker D, Hernán MA, et al. Fourth dose
- 482 of BNT162b2 mRNA Covid-19 vaccine in a nationwide setting. N Engl J Med
  483 2022;386:1603–14.
- 484 [21] Gardner BJ, Kilpatrick AM. Estimates of reduced vaccine effectiveness against
  485 hospitalization, infection, transmission and symptomatic disease of a new SARS-CoV-2
  486 variant, Omicron (B.1.1.529), using neutralizing antibody titers. BioRxiv 2021.
  487 https://doi.org/10.1101/2021.12.10.21267594.
- [22] Lopez Bernal J, Gower C, Andrews N, Public Health England Delta Variant Vaccine
  Effectiveness Study Group. Effectiveness of covid-19 vaccines against the B.1.617.2 (delta)
  variant. Reply. N Engl J Med 2021;385:e92.
- 491 [23] Higdon MM, Baidya A, Walter KK, Patel MK, Issa H, Espié E, et al. Duration of
  492 effectiveness of vaccination against COVID-19 caused by the omicron variant. Lancet
- 493 Infect Dis 2022. https://doi.org/10.1016/S1473-3099(22)00409-1.
- 494 [24] Walls AC, Fiala B, Schäfer A, Wrenn S, Pham MN, Murphy M, et al. Elicitation of potent
  495 neutralizing antibody responses by designed protein nanoparticle vaccines for SARS-CoV496 2. Cell 2020;183:1367-1382.e17.
- 497 [25] Marcandalli J, Fiala B, Ols S, Perotti M, de van der Schueren W, Snijder J, et al. Induction
- 498 of potent neutralizing antibody responses by a designed protein nanoparticle vaccine for
- respiratory syncytial virus. Cell 2019;176:1420-1431.e17.

- 500 [26] Bachmann MF, Rohrer UH, Kündig TM, Bürki K, Hengartner H, Zinkernagel RM. The
  501 influence of antigen organization on B cell responsiveness. Science 1993;262:1448–51.
- 502 [27] Wang JW, Roden RBS. Virus-like particles for the prevention of human papillomavirus-
- associated malignancies. Expert Rev Vaccines 2013;12:129–41.
- 504 [28] López-Sagaseta J, Malito E, Rappuoli R, Bottomley MJ. Self-assembling protein
  505 nanoparticles in the design of vaccines. Comput Struct Biotechnol J 2016;14:58–68.
- 506 [29] Bruce MG, Bruden D, Hurlburt D, Zanis C, Thompson G, Rea L, et al. Antibody levels and
- 507 protection after hepatitis B vaccine: Results of a 30-year follow-up study and response to a
- 508 booster dose. J Infect Dis 2016;214:16–22.
- [30] Kreimer AR, Herrero R, Sampson JN, Porras C, Lowy DR, Schiller JT, et al. Evidence for
  single-dose protection by the bivalent HPV vaccine—Review of the Costa Rica HPV
  vaccine trial and future research studies. Vaccine 2018;36:4774–82.
- [31] Walls AC, Miranda MC, Schäfer A, Pham MN, Greaney A, Arunachalam PS, et al.
  Elicitation of broadly protective sarbecovirus immunity by receptor-binding domain
  nanoparticle vaccines. Cell 2021;184:5432-5447.e16.
- 515 [32] Brouwer PJM, Brinkkemper M, Maisonnasse P, Dereuddre-Bosquet N, Grobben M,
  516 Claireaux M, et al. Two-component spike nanoparticle vaccine protects macaques from
  517 SARS-CoV-2 infection. Cell 2021;184:1188-1200.e19.
- 518 [33] Boyoglu-Barnum S, Ellis D, Gillespie RA, Hutchinson GB, Park Y-J, Moin SM, et al.
  519 Quadrivalent influenza nanoparticle vaccines induce broad protection. Nature
  520 2021;592:623–8.
- [34] King NP, Bale JB, Sheffler W, McNamara DE, Gonen S, Gonen T, et al. Accurate design of
  co-assembling multi-component protein nanomaterials. Nature 2014;510:103–8.

523	[35] Bale JB, Gonen S, Liu Y, Sheffler W, Ellis D, Thomas C, et al. Accurate design of
524	megadalton-scale two-component icosahedral protein complexes. Science 2016;353:389-
525	94.

- 526 [36] Ueda G, Antanasijevic A, Fallas JA, Sheffler W, Copps J, Ellis D, et al. Tailored design of
- 527 protein nanoparticle scaffolds for multivalent presentation of viral glycoprotein antigens.528 Elife 2020;9:e57659.
- [37] Wargacki AJ, Wörner TP, van de Waterbeemd M, Ellis D, Heck AJR, King NP. Complete
  and cooperative in vitro assembly of computationally designed self-assembling protein
  nanomaterials. Nat Commun 2021;12:883.
- [38] Arunachalam PS, Walls AC, Golden N, Atyeo C, Fischinger S, Li C, et al. Adjuvanting a
  subunit COVID-19 vaccine to induce protective immunity. Nature 2021;594:253–8.
- [39] Ellis D, Brunette N, Crawford KHD, Walls AC, Pham MN, Chen C, et al. Stabilization of
- the SARS-CoV-2 Spike receptor-binding domain using deep mutational scanning and
  structure-based design. Front Immunol 2021;12:710263.
- 537 [40] Fluckiger A-C, Ontsouka B, Bozic J, Diress A, Ahmed T, Berthoud T, et al. An enveloped
- virus-like particle vaccine expressing a stabilized prefusion form of the SARS-CoV-2 spike
  protein elicits highly potent immunity. Vaccine 2021;39:4988–5001.
- 540 [41] Arunachalam PS, Feng Y, Ashraf U, Hu M, Walls AC, Edara VV, et al. Durable protection
- against the SARS-CoV-2 Omicron variant is induced by an adjuvanted subunit vaccine. Sci
  Transl Med 2022;14:eabq4130.
- 543 [42] Grigoryan L, Lee A, Walls AC, Lai L, Franco B, Arunachalam PS, et al. Adjuvanting a
- 544 subunit SARS-CoV-2 vaccine with clinically relevant adjuvants induces durable protection

545 in mice. NPJ Vaccines 2022;7:55.

- 546 [43] Song JY, Choi WS, Heo JY, Lee JS, Jung DS, Kim S-W, et al. Safety and immunogenicity
- 547 of a SARS-CoV-2 recombinant protein nanoparticle vaccine (GBP510) adjuvanted with
- 548 AS03: A randomised, placebo-controlled, observer-blinded phase 1/2 trial.
- 549 EClinicalMedicine 2022;51:101569.
- 550 [44] Joyce MG, Chen W-H, Sankhala RS, Hajduczki A, Thomas PV, Choe M, et al. SARS-CoV-
- 2 ferritin nanoparticle vaccines elicit broad SARS coronavirus immunogenicity. Cell Rep
  2021;37:110143.
- [45] Saunders KO, Lee E, Parks R, Martinez DR, Li D, Chen H, et al. Neutralizing antibody
  vaccine for pandemic and pre-emergent coronaviruses. Nature 2021;594:553–9.
- 555 [46] Dalvie NC, Tostanoski LH, Rodriguez-Aponte SA, Kaur K, Bajoria S, Kumru OS, et al.
- SARS-CoV-2 receptor binding domain displayed on HBsAg virus-like particles elicits
  protective immunity in macaques. Sci Adv 2022;8:eabl6015.
- [47] Tan TK, Rijal P, Rahikainen R, Keeble AH, Schimanski L, Hussain S, et al. A COVID-19
  vaccine candidate using SpyCatcher multimerization of the SARS-CoV-2 spike protein
  receptor-binding domain induces potent neutralising antibody responses. Nat Commun
  2021;12:542.
- 562 [48] Dalvie NC, Rodriguez-Aponte SA, Hartwell BL, Tostanoski LH, Biedermann AM, Crowell
- LE, et al. Engineered SARS-CoV-2 receptor binding domain improves manufacturability in
  yeast and immunogenicity in mice. Proc Natl Acad Sci U S A 2021;118:e2106845118.
- [49] Cohen AA, Gnanapragasam PNP, Lee YE, Hoffman PR, Ou S, Kakutani LM, et al. Mosaic
  nanoparticles elicit cross-reactive immune responses to zoonotic coronaviruses in mice.
- 567 Science 2021;371:735–41.

568	[50] Hsia Y, Bale JB, Gonen S, Shi D, Sheffler W, Fong KK, et al. Design of a hyperstable 60-
569	subunit protein dodecahedron. [corrected]. Nature 2016;535:136–9.

- 570 **Figure captions**
- 571 Figure 1: Production and characterization of IVX-411 VLPs. (A) Schematic of in vitro
- assembly. (B) UV-Vis spectroscopy. (C) Dynamic Light Scattering. (D) Size Exclusion
- 573 Chromatography. (E) Biolayer Interferometry. (F) Negative stain Transmission Electron
- 574 Microscopy.

575 Figure 2: Immunization with IVX-411 elicits neutralizing antibodies against four SARS-CoV-2 576 strains. (A) Female BALB/c mice were immunized IM on Day 0 and Day 21 with IVX-411 577 formulated with MF59 and bled on Day 0 (pre-immunization), Day 21, and Day 35 (n = 8). 578 Serum neutralizing titers against the ancestral strain (**B**), as well as the beta (**C**), gamma (**D**), and 579 delta (E) variants were measured using a cell-based pseudo-particle neutralization assay. Human 580 convalescent serum (HCS) was used as a positive control. Dashed horizontal line represents the 581 lower limit of quantitation. P-values were calculated in GraphPad Prism 9 using a Wilcoxon 582 matched-pairs signed rank test. \*, P<0.05; \*\*, P<0.01.

583 Figure 3: Immunization with IVX-411 elicits higher neutralizing titers and a larger antigenspecific LLPC compartment than immunization with soluble protein. A) Female BALB/c mice 584 585 were immunized IM with IVX-411, RBD-CompA, or S-2P formulated with or without MF59 on 586 Day 0 and Day 21. Serum was collected on days 0, 20, 35, 63, 91, 119, and 154. Mice were 587 sacrificed and bone marrow was collected on Day 154 for isolation of LLPCs (n = 10). (B) 588 Serum neutralizing titers against the ancestral SARS-CoV-2 strain. (C) ELISpot measuring the 589 frequency of RBD-specific LLPCs in the bone marrow. Dashed horizontal line represents the 590 lower limit of quantitation. P-values were calculated in GraphPad Prism 9 using a Mann591 Whitney test or a Kruskal-Wallis test followed by a Dunn's multiple comparisons test. \*, P< 592 0.05; \*\*, P<0.01; \*\*\*, P<0.001; \*\*\*\*, P<0.0001.

593 Figure 4: Immunization of Syrian Golden hamsters with IVX-411 reduces severity of SARS-

- 594 CoV-2 infection. (A) Male Syrian Golden hamsters were immunized IM with phosphate buffered
- saline (PBS) or IVX-411 formulated with MF59 on days 0 and 21. Hamsters in all groups except
- the PBS, Unchallenged group were inoculated intranasally with one of two SARS-CoV-2
- isolates on Day 42. All hamsters were bled on days 0, 21, and 35. Body weights were recorded
- daily starting on Day 39 until Day 47 when the animals were sacrificed (n = 8). (B) Body
- 599 weights represented as percentage of starting weight prior to challenge. (C) Lung weights

600 represented as a percentage of total body weight post challenge. (D) Histopathological analysis

601 was performed on lungs collected from each animal. Findings were graded on a scale of 1-5 (1 =

Minimal, 2 = Mild, 3 = Moderate, 4 = Marked, 5= Severe). The average severity score for each

group is reported. P-values were calculated in GraphPad Prism 9 using a Mann-Whitney test or a

604 Kruskal-Wallis test followed by a Dunn's multiple comparisons test. \*, P< 0.05; \*\*, P<0.01;

605 \*\*\*, P<0.001; \*\*\*\*, P<0.0001.

606 Figure 5: Prophylactic immunization with IVX-411 elicits neutralizing antibodies and results in 607 decreased viral load upon challenge in Syrian Golden hamsters. Number of N (A) or E (B) gene 608 copies present in lung tissue or nasal swab of Syrian Golden hamsters challenged with 609 WA1/2020 or B.1.617.2 (n = 8). (C) Neutralizing antibody titers against the ancestral strain and 610 delta variant on Day 0 and Day 35 in hamsters immunized with IVX-411. Dashed horizontal line 611 represents the lower limit of quantitation. P-values were calculated in GraphPad Prism 9 using a 612 Mann-Whitney test or a Kruskal-Wallis test followed by a Dunn's multiple comparisons test. \*, 613 P<0.05; \*\*, P<0.01; \*\*\*, P<0.001; \*\*\*\*, P<0.0001.

# 614 Supplemental Figure 1: (A) UV-Vis spectroscopy and (B) SDS-PAGE of purified RBD-

615 CompA trimeric component.











# Figure 5

