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#### Supplemental information

#### SARS-CoV-2 ferritin nanoparticle vaccines elicit

#### broad SARS coronavirus immunogenicity

M. Gordon Joyce, Wei-Hung Chen, Rajeshwer S. Sankhala, Agnes Hajduczki, Paul V. Thomas, Misook Choe, Elizabeth J. Martinez, William C. Chang, Caroline E. Peterson, Elaine B. Morrison, Clayton Smith, Rita E. Chen, Aslaa Ahmed, Lindsay Wieczorek, Alexander Anderson, James Brett Case, Yifan Li, Therese Oertel, Lorean Rosado, Akshaya Ganesh, Connor Whalen, Joshua M. Carmen, Letzibeth Mendez-Rivera, Christopher P. Karch, Neelakshi Gohain, Zuzana Villar, David McCurdy, Zoltan Beck, Jiae Kim, Shikha Shrivastava, Ousman Jobe, Vincent Dussupt, Sebastian Molnar, Ursula Tran, Chandrika B. Kannadka, Sandrine Soman, Caitlin Kuklis, Michelle Zemil, Htet Khanh, Weimin Wu, Matthew A. Cole, Debra K. Duso, Larry W. Kummer, Tricia J. Lang, Shania E. Muncil, Jeffrey R. Currier, Shelly J. Krebs, Victoria R. Polonis, Saravanan Rajan, Patrick M. McTamney, Mark T. Esser, William W. Reiley, Morgane Rolland, Natalia de Val, Michael S. Diamond, Gregory D. Gromowski, Gary R. Matyas, Mangala Rao, Nelson L. Michael, and Kayvon Modjarrad



pCoV132: Y453R + L518R

## Figure S1. Structure-based design of SARS-CoV-2 S-domain ferritin nanoparticle immunogens and design pipeline. Related to Figure 1.

Four ferritin nanoparticle immunogen designs were developed focused on (1) Spike ferritin nanoparticles (blue), (2) RBD ferritin nanoparticles (green), (3) RBD-NTD ferritin nanoparticles (black), and (4) S1 ferritin nanoparticles (orange). The design iterations and concepts are indicated, along with select mutations and design name. Lead vaccine candidates from each category are highlighted.



# Figure S2. Negative-stain electron microscopy 2D micrographs of SARS-CoV-2 ferritin nanoparticle vaccine candidates. Related to Figure 2 and 4.

Negative-stain electron microscopy 2D micrographs. The white scale bars represent 100 nm.

- (A) Spike ferritin nanoparticles pCoV1B-05 and pCoV1B-08.
- (B) RBD ferritin nanoparticles pCoV03, pCoV50, pCOV58, pCoV59, pCoV127, pCoV129, pCoV130, pCoV131
- (C) RBD-NTD ferritin nanoparticles pCoV122, pCoV125, pCoV147
- (D) S1 ferritin nanoparticle pCoV110 and pCoV112.



Figure S3. Biophysical and antigenic characterization of S-domain ferritin nanoparticle immunogens. Related to Figure 2 and 3. (A) Size-exclusion chromatography on a Superdex S200 10/300 column of representative SARS-CoV-2 S-based ferritin nanoparticles from

- (B) Expression levels (mg/L supernatant) of representative SARS-CoV-2 Spike-based ferritin nanoparticles.
- (C) Dynamic light scattering analysis of representative SARS-CoV-2 Spike-based ferritin nanoparticles.
- (D) Spike ferritin nanoparticles (E) RBD ferritin, (F) RBD-NTD ferritin and (G) S1 ferritin nanoparticles were assessed for binding to a set of neutralizing antibodies (concentration = 30 μg/ml) by biolayer interferometry.



## Figure S4 SARS-CoV-2 nanoparticle vaccine candidates elicit robust binding and pseudovirus neutralizing antibody responses in mice. Related to Figure 5 and 7.

Data relating to each category of immunogen are colored as follows: SpFN (blue), RFN (green), pCoV146 (black) and pCoV111 (orange). N = 10/group. (A) Biolayer Interferometry binding analysis of C57BL/6 and BALB/c sera from mice immunized with SpFN + Alhydrogel® (B) RFN + Alhydrogel® and (C) pCoV146 + Alhydrogel® to SARS-CoV-2 RBD. Mean values are indicated by a horizontal line, n=10.

(D) Pseudovirus neutralization (ID<sub>50</sub> values) of C57BL/6 and BALB/c sera from mice immunized with SpFN + Alhydrogel® (E) RFN + Alhydrogel® and (F) pCoV146 + Alhydrogel®. Geometric mean values are indicated by a horizontal line, n=10.

(G) Biolayer interferometry analysis of BALB/c mouse sera binding to SARS-CoV-2 RBD at study weeks 2, 5 and 8. Mice were immunized with the four lead candidate vaccines SpFN (blue), RFN (green), pCoV146 (black) and pCOV111 (orange). Binding mean values are indicated by a horizontal line, n=10, sera responses at a given study week were compared for statistical differences using a Kruskal-Wallis test followed by a Dunn's post-test.

(H) ELISA analysis of BALB/c mice immune responses as indicated in (G). Binding geometric mean values of the endpoint titers are indicated by a horizontal line, n=10, sera responses at a given study week were compared for statistical differences using a Kruskal-Wallis test followed by a Dunn's post-test.
(I) ACE2 blocking assay ID50 inhibition titers of study week 10 mouse sera from C57BL/6 mice (left) and BALB/c mice (right) immunized with the fourl ead

(I) ACE2 blocking assay ID50 inhibition titers of study week 10 mouse sera from C57BL/6 mice (left) and BALB/c mice (right) immunized with the fourlead immunogens from each design category (colored as in Figure 5).

(J) Pseudovirus neutralization ID50 titers of BALB/c mice immunized as indicated in (G). Geometric mean values are indicated by a horizontal line, n=10, sera neutralization titers at a given study week for the four immunogens were compared for statistical differences using a Kruskal-Wallis test followed by a Dunn's post-test.

(K) Pseudovirus neutralization ID<sub>80</sub> titers of C57BL/6 (left) and BALB/c mice (right) immunized as indicated in (G). Geometric mean values are indicated by a horizontal line, n=10, sera neutralization titers at a given study week for the four immunogens were compared for statistical differences using a Kruskal-Wallis test followed by a Dunn's post-test.

P values <0.0001 (\*\*\*\*), <0.001 (\*\*\*), <0.01 (\*\*) or <0.05 (\*).



## Figure S5 SARS-CoV-2 SpFN vaccine candidate elicits robust binding and neutralizing antibody responses at reduced doses in mice. Related to Figure 5 and 7.

(A) Biolayer interferometry analysis of C57BL/6 and (B) BALB/c mouse sera binding response to SARS-CoV-2 RBD following immunization with reducing doses of SpFN.

(C, E) ELISA analysis of C57BL/6 and (D, F) BALB/c mouse sera binding response to SARS-CoV-2 RBD or S-2P following immunization with reducing doses of SpFN.

(G) SARS-CoV-2 pseudovirus ID80 neutralization titers of mice immunized with 0.08 µg SpFN + ALFQ.

(H) Authentic SARS-CoV-2 virus ID80 neutralization titers of mice immunized with 10  $\mu$ g (blue) or 0.08  $\mu$ g (light blue) SpFN + ALFQ. Geometric mean titers for each group and time point are indicated by a horizontal line, n =10. Neutralization titers for the two dose groups at each study time point were compared for statistically significant differences using a Mann-Whitney unpaired two-tailed non-parametric test. The two BALB/c time points that showed differences are indicated by bars. P values <0.001 (\*\*\*), <0.01 (\*\*).

(I) Mouse sera from study week 10 was analyzed for hACE2 blocking capacity to SARS-CoV-2 RBD (left) or SARS-CoV-1 RBD using a biolayer interferometry assay format.



Α

В

С

Figure S6. SARS-CoV-2 S-domain ferritin nanoparticle vaccine candidates elicit durable binding and neutralizing antibody responses. Related to Figure 5 and 7.

(A) ELISA binding of mouse sera to S-2P and RBD antigens (Square: S-2P; Circle: RBD). Sera samples up to week 20 of the mouse studies were assessed for durability of binding responses.

(B) Pseudovirus neutralization ID50 values of SARS-CoV-2 by mouse sera following SpFN (blue) or RFN (green) immunization at multiple timepoints. Samples were assessed from both C57BL/6 or BALB/c mice.

(C) Biolayer Interferometry binding of study week 10 immunized C57BL/6 mouse serum to SARS-CoV-2 RBD, and SARS-CoV-2 RBD variants. Immunogens are indicated at the top left of each graph. Mean values are indicated by a horizontal line, n=10, Significance was assessed using a Kruskal-Wallis test followed by a Dunn's post-test.

(D) Pseudovirus ID50 neutralization of SARS-CoV-2 VoC by immune sera from study week 20 for SpFN-immunized, or study week 17 for RFN-immunized mice



В



Figure S7 Histopathological analysis of SARS-CoV-2 infection in K18-ACE2 mice. Related to Figure 7 (A, B) Hematoxylin and eosin staining of lung sections from K18-hACE2 mice following intranasal infection with 1.25 x 10<sup>4</sup> PFU SARS-CoV-2. Images show two magnifications. Images are representative of n = 10 per group.

Α

#### Table S1. Spike-domain ferritin immunogens, Related to Figure 1.

Spike-Ferritin (all	based on S-2P variant with Dfurin and PP)		
Construct ID	Description		
		Shortened ectodomain - no coiled coil (closest to	
pCoV1B-01	S2P(1-1137)-del-4-Ferritin	flu HA pass off)	NL
		Shortened ectodomain - no coiled coil (closest to	
pCoV1B-02	S2P(1-1137)-del-6-Ferritin	flu HA pass off)	NL
pCoV1B-03	S2P(1-1208)-del-Ferritin	Full ectodomain	NL
pCoV1B-04	S2P(1-1208)-GCN4-Ferritin	Full ectodomain with GCN4	NL
		Shortened ectodomain with ending with a couple	
pCoV1B-05	S2P(1-1154)-del-Ferritin	turns of coiled coil	NL
pCoV1B-06	S2P(1-1158)op1-del-Ferritin	Optimized HR ending (end on glycan N1158)	NL
		Optimized HR ending (IIe) (end on glycan	
pCoV1B-07	S2P(1-1158)op2-del-Ferritin	N1158)	NL
		Optimized HR ending (N1158 glycan removed,	
pCoV1B-08	S2P(1-1158)op1x2-del-Ferritin	but exists on the repeated HR)	NL
		Optimized HR ending (IIe) (N1158 glycan	
pCoV1B-09	S2P(1-1158)op2x2-del-Ferritin	removed, but exists on the repeated HR)	NL
		Optimized HR ending with GCN4 fused in	
pCoV1B-10	S2P(1-1158)op1-fGCN4-del-Ferritin	register (no glycan N1158)	NL
		Shortened ectodomain - no coiled coil (closest to	PI
pCoV1B-01-PL	PL-S2P(12-1137)-del-4-Ferritin	flu HA pass off)	• =
		Shortened ectodomain - no coiled coil (closest to	PL
pCoV1B-02-PL	PL-S2P(12-1137)-del-6-Ferritin	flu HA pass off)	
pCoV1B-03-PL	PL-S2P(12-1208)-del-Ferritin	Full ectodomain	PL
pCoV1B-04-PL	PL-S2P(12-1208)-GCN4-Ferritin	Full ectodomain with GCN4	PL
		Shortened ectodomain with ending with a couple	PL
	PL-S2P(12-1154)-del-Ferritin	turns of colled coll	
pcov-IB-06-PL	PL S2P(12,1158) and del Earritin	Optimized UP anding (and an glycon N1159)	PL
(aka Sprin)	PL-52P(12-1156)0p1-dei-Femilin	Optimized HR ending (le) (and on glycan NT 56)	
	DL S2D(12 1159)on2 dol Earritin	N1159)	PL
pcovid-07-PL	PL-32P(12-1156)0p2-dei-Permin	NTTOO) Optimized UP opding (N1159 glycop removed	
nCo\/1B_08_PI	PL-S2P(12-1158)op1x2-del-Earritin	but exists on the repeated HP)	PL
PC0112-00-FL		Ontimized HR ending (IIe) (N1158 alvcan	
nCo\/1B-09-PI	PL-S2P(12-1158)on2x2-del-Ferritin	removed but exists on the repeated HR)	PL
P00110001E		Ontimized HR ending with GCN4 fused in	
pCoV1B-10-PI	PL-S2P(12-1158)op1-fGCN4-del-Ferritin	register (no glycan N1158)	PL

RBD-Ferritin			
Construct ID	Description	Comment	Leader
		N-terminal His8 with HRV-3C cleavage site,	
pCoV03	His8-3C-RBD(331-527)-Ferritin	GSGGGG linker between RBD and Ferritin	PL
pCoV29	His8-3C-RBD-3-Ferritin	SGG linker	PL
pCoV30	His8-3C-RBD-3-del-Ferritin	SGG linker, D first 10 residues in ferritin, then DIEK changed to DIIK	PL
pCoV31	His8-3C-RBD-6-del-Ferritin	P527G, D first 8 residues in ferritin, then SKDIEK changed to DIIK	PL
pCoV1A-01	His8-3C-RBD-PPII-Ferritin	Extend distance between RBD and ferritin - using polyproline Helix	PL
pCoV1A-02	His8-3C-RBD-alpha1-Ferritin	Extend distance between RBD and ferritin - using alpha Helix from bottom of S protein	PL
pCoV1A-03	His8-3C-RBD-alpha2-Ferritin	Extend distance between RBD and ferritin- using alpha Helix from bottom of S protein	PL
pCoV1A-04	His8-3C-RBD-GCN4-del-Ferritin	Extend distance between RBD and ferritin + stabilize ferritin - using GCN4 trimerization motif	PL
	Hige 20 PPD 1111 1159ap1 dol Egritin	Extend distance between RBD and ferritin + stabilize ferritin - using semi-native trimerization	PL
PC0V IA-05	1130-30-RDD-1141_11300p1-001-F011011	Extend distance between RBD and ferritin +	PI
pCoV1A-06	His8-3C-RBD-1141_1158op1x2-del-Ferritin	motif	

RBD-Ferritin (continued)					
Construct ID	Description	Comment	Leader		
pCoV49	His8-3C-RBD-F456N/K458T-Ferritin	RBD with indicated point mutations	PL		
pCoV50	His8-3C-RBD-L455R/Y449K/F490R-Ferri	tin RBD with indicated point mutations	PL		
pCoV51	His8-3C-RBD-L455R-Ferritin	RBD with indicated point mutation	PL		
pCoV52	His8-3C-RBD-I468R-Ferritin	RBD with indicated point mutation	PL		
pCoV53	His8-3C-RBD-Y453R-Ferritin	RBD with indicated point mutation	PL		
pCoV54	His8-3C-RBD-L452R-Ferritin	RBD with indicated point mutation	PL		
pCoV55	His8-3C-RBD-L492R-Ferritin	RBD with indicated point mutation	PL		
pCoV56	His8-3C-RBD-F490R-Ferritin	RBD with indicated point mutation	PL		
pCoV57	His8-3C-RBD-F490A-Ferritin	RBD with indicated point mutation	PL		
pCoV58	His8-3C-RBD-L517N/L518K/H519S-Ferri	tin RBD with indicated point mutations	PL		
pCoV59	His8-3C-RBD-L518R-Ferritin	RBD with indicated point mutation	PL		
pCoV60	His8-3C-RBD-V367T/L335N-Ferritin	RBD with indicated point mutations	PL		
pCoV61	His8-3C-RBD-T385N/L387T-Ferritin	RBD with indicated point mutations	PL		
pCoV62	His8-3C-RBD-V382R-Ferritin	RBD with indicated point mutation	PL		
pCoV63	His8-3C-RBD-F377R-Ferritin	RBD with indicated point mutation	PL		
	His8-3C-RBD-F490A/L518N/L519K/H520S-				
pCoV127	Ferritin	RBD with indicated point indiations	ΓL		
pCoV128	His8-3C-RBD-F490A/L518R-Ferritin	RBD with indicated point mutations	PL		
	His8-3C-RBD-				
	L455R/Y449K/F490R/L517N/L518K/	RBD with indicated point mutations	PL		
pCoV129	H519S-Ferritin				
	His8-3C-RBD-L455R/Y449K/F490R/	RBD with indicated point mutations	PI		
pCoV130	L518R-Ferritin	NBD with indicated point mutations	ΓL		
pCoV131 (aka	His8-3C-RBD-Y453R/L517N/L518K/	RBD with indicated point mutations	PI		
RFN)	H519S-Ferritin	Rep with indicated point indiations			
pCoV132	His8-3C-RBD-Y453R/L518R-Ferritin	RBD with indicated point mutations	PL		

#### **RBD-NTD-Ferritin**

Construct ID	Description		
pCoV122	His8-3C-RBD(331-527)-GSGGSG-NTD(12 303)-Ferritin	N-terminal HIs8 with HRV-3C cleavage site, GSGGSG linker between RBD and NTD, GSGGGG linker between NTD and Ferritin	PL
pCoV123	His8-3C-RBD-F490R-NTD-Ferritin	RBD with indicated point mutation	PL
pCoV124	His8-3C-RBD-F490A-NTD-Ferritin	RBD with indicated point mutation	PL
pCoV125	His8-3C-RBD-L517N/L518K/H519S-NTD- Ferritin	RBD with indicated point mutations	PL
pCoV126	His8-3C-RBD-L518R-NTD-Ferritin	RBD with indicated point mutation	PL
pCoV146	His8-3C-RBD-Y453R- L517N/L518K/H519S-NTD-Ferrritin	RBD with indicated point mutations	PL
pCoV147	His8-3C-RBD-F490A-L517N/L518K/ H519S-NTD-Ferrritin	RBD with indicated point mutations	PL

S1-Ferritin			
Construct ID	Description		
pCoV68	S1(12-678)-Ferritin	GSGGSG linker between S1 and Ferritin	PL
pCoV107	S1(12-655)-Ferritin	24 residues removed from the C-terminus	PL
pCoV108	S1(12-655)-L611N/Q613T-Ferritin	24 residues removed from the C-terminus, S1 with indicated point mutations	PL
pCoV109	S1(12-696)-Ferritin	Extended the sequence to include a portion of S2	PL
pCoV110	S1(12-676)-G-S2(689-696)-Ferritin	Extended the sequence to include a portion of S2 with the indicated leader between the two regions	PL
pCoV111	S1(12-676)-GG-S2(689-696)-Ferritin	Extended the sequence to include a portion of S2 with the indicated leader between the two regions	PL
nCo\/112	S1(12-676)-PG-S2(689-696)-Ferritin	Extended the sequence to include a portion of S2 with the indicated leader between the two regions	PL
pC0V112	S1-V312NI/O313V/T31/T-Forritin	S1 with indicated point mutations	PI
pCoV113	S1-1651N/A653S-Ferritin	S1 with indicated point mutations	PI
pCoV115	S1-S316C/V595C-Ferritin	S1 with indicated point mutations	PI
pCoV116	S1-V320C/S591C-Ferritin	S1 with indicated point mutations	PL
pCoV117	S1-L560Q/F562H-Ferritin	S1 with indicated point mutations	PL
pCoV118	S1-F562N/Q564T-Ferritin	S1 with indicated point mutations	PL
pCoV119	S1-F490R-Ferritin	S1 with indicated point mutation	PL
pCoV120	S1-F490A-Ferritin	S1 with indicated point mutation	PL
pCoV02	S1(16-678)-Ferritin	4 residues removed from N-terminus	PL
pCoV67	His8-3C-S1-Ferritin	His8 and HRV-3C cleavage site added to N- terminus	PL

	-				
Protein	SpFN_1B-06-	RFN_131	pCoV146	pCoV111	pCoV1B-05
	PL	_			1
Immunogen Fused	Spike (S2P)	RBD	RBD-NTD	<b>S</b> 1	Spike (S2P)
EMDB Code	EMD-25448	EMD-25449	EMD-25450	EMD-25451	N/A
<b>Data Collection</b>					
Microscope	Tecnai	Tecnai	Tecnai	Tecnai	Talos
	T20	T20	T20	T20	L120C
Voltage (kV)	200 kV	200 kV	200 kV	200 kV	120 kV
Camera	Eagle 4K	Eagle 4K	Eagle 4K	Eagle 4K	Ceta
Software	SerialEM	SerialEM	SerialEM	SerialEM	EPU
Pixel Size (Å/pix)	2.195	2.195	2.195	2.195	2.542
Underfocus range	0.7-1.3	0.8-1.3	0.6-1.5	0.8-1.6	0.5-0.9
Image Processing					
Software	RELION	RELION	RELION	RELION	RELION
	3.0.8	3.0.8	3.0.8	3.0.8	3.1.1
# Particle Images	11502	3383	832	2121	2143
Pixel Size (Å/pixel)	4.39	4.39	4.39	4.39	5.084
Box Size (pixels)	160	160	160	160	200
Symmetry (3D)	0	0	0	0	
Initial Lowpass (Å)	100	80	100	100	
(RELION)	100	80	100	100	
Resolution (Å)	25	21	30	30	

Table S2. Negative-stain Electron Microscopy Data Collection and Refinement, Related to Figure 4.

	Turning and the law		D - 11- / -		D - 11- / -	
pCOv no.	Immunogen design	C3/BL/	Balb/c	C3/BL/6	Balb/c	
		6 ALFQ	ALFQ	Alhydrogel	Alhydrogel	
1B-05	S-Trimer-Ferritin	X	Х			
1B-06-PL	S-Trimer-Ferritin	X	Х	X	Х	
<b>RBD-Ferrit</b>	in constructs					
pCOV no.	Immunogen design	C57BL/	Balb/c	C57BL/6	Balb/c	
		6 ALFQ	ALFQ	Alhydrogel	Alhydrogel	
50	RBD-Ferritin		Х			
58	RBD-Ferritin	X	Х	X	Х	
59	RBD-Ferritin		Х			
127	RBD(57+58)-Ferritin	X	Х	X	Х	
129	RBD(50+58)-Ferritin	X	Х	X	Х	
130	RBD(50+59)-Ferritin		Х			
131	RBD(53+58)-Ferritin	X	Х	X	Х	
S1-Ferritin	S1-Ferritin constructs					
pCOV no.		C57BL/	Balb/c	C57BL/6	Balb/c	
		6 ALFQ	ALFQ	Alhydrogel	Alhydrogel	
111	S1-Ferritin	X	Х			
RBD-NTD-Ferritin constructs						
pCOV no.	Immunogen design	C57BL/	Balb/c	C57BL/6	Balb/c	
		6 ALFQ	ALFQ	Alhydrogel	Alhydrogel	
122	RBD-NTD-Ferritin	X	X			
125	RBD(58)-NTD-Ferritin		X		X	
146	RBD(53+58)-NTD-Ferritin	X	X	X	X	
147	RBD(57+58)-NTD-Ferritin	X				

**Table S4. Animal immunogenicity SARS-CoV-2 pseudovirus neutralization ID50 and ID80, Related to Figure 5.** Numbers shown are the ID50/ID80 geometric mean titers for a group, with study week 2, 5, and 8 shown in vertical order.

Spike-Ferritin constructs					
•		C57BL/6	Balb/c	C57BL/6	Balb/c
		ALFQ	ALFQ	Alhydrogel	Alhydrogel
1B-05	S-Trimer-Ferritin (x 2	702/189	115/<80		
	groups)	13.076/5.647	5,546/1.447		
1B-06-PL	S-Trimer-Ferritin	14,976/5,397	1,152/355	297/118	404/<80
		41,237/16,818	16,816/6,662	1074/239	1484/390
	tin constructs	47,323/16,524	25,062/6,540	3901/153	5591/ 1334
KDD-Ferr		C57PL/6	Dalb/a	C57DI /6	Palb/a
pCOV no.	Immunogen design	ALFO	AL FO	C3/BL/0 Alhydrogel	Alhydrogel
50	RBD-Ferritin		X	ringuloger	rinydroger
50	DDD Examitin	577/238	353/123	293/211	232/~80
58	KBD-Fermin	11,224/2,793 31,562/10,09	13,466/3,802 25,340/7,692	1,734/688 5,097/1261	4,836/1,086 9,439/2,569
59	RBD-Ferritin		Х		
127	RBD(57+58)-Ferritin	Х	Х	Х	Х
129	RBD(50+58)-Ferritin	Х	Х	Х	Х
130	RBD(50+59)-Ferritin		Х		
131	RBD(53+58)-Ferritin	358/107	270/95	682/163	119/<40
131	KDD(55+50) Territin	15,950/5,667 38 110/12 824	13,090/3,539 32 969/10 079	1,181/403 2 845/529	182/103 240/99
S1-Ferritir	n constructs	56,110/12,024	52,505/10,075	2,043/329	240/77
pCOV no		C57BL/6	Balb/c	C57BL/6	Balb/c
peevino.		ALFQ	ALFQ	Alhydrogel	Alhydrogel
111	S1-Ferritin	1,770/350 14,893/3,636	450/172 18,112/3,846		
19,157/5,564 17,108/3,886					
KBD-NTD	-Ferritin constructs	05701/6	D . 11. /	05701/6	D 11. /
pCOV no.	Immunogen design	C5/BL/6	Balb/c	C5/BL/6	Balb/C
122	PRD NTD Forritin	X	X	Aiiiyulogel	Allyulogei
122	DDD(58) NTD Eorritin		X		X
125	$\mathbf{RDD}(52, 59) \mathbf{NTD} \mathbf{E}_{\mathbf{r}}$	230/91	2/0/89	~80/~80	662/~80
146	RBD(53+58)-NID-Ferritin	16,678/4,356	31,252/7,190	667/460	2,087/537
1.47	DDD(57 59) NTD Equitin	20,107/6,126 X	24,854/6,744	940/289	2,417/701
147	RBD(57+58)-NTD-Ferriun	А			
Non-Ferrit	in constructs	GEEDI /	D 11 /		D.11./
pCOV no.	Immunogen	C5/BL/6	Balb/c	C5/BL/6 Albydrogel	Balb/c Albydrogel
8	PBD	<80/<80	<80/<80	<80/<80	<80/<80
0	KDD	<80/<80	<80/<80	<80/<80	<80/<80
47	C 2D	1,825/450	1,518/364	<80/<80	<80/<80
47	S-2P	495/100 47,201/16,041 33,853/18012	<80/<80 14,612/4,130 20,954/5660	<80/<80 1175/288 4.619/1.168	<80/<80 3,992/1,213 10.064/3132
Ferritin Construct designs in the literature					
	Immunogen (Reference)		BALB/c		
	S $\Delta$ C-Fer (Powell et al., 2021)	Pseudovirus	Prime + Roost: 34,000		
		neuuanzation:	BALB/c		
	RBD-Ferritin incorporating		SAS aujuvant		
	SpyTag-Spycatcher (Kang et al., 2020)	Live virus neutralization	FRNT <sub>90</sub> : 8,192		