

**Epitope-specific antibody responses differentiate COVID-19 outcome
and variants of concern**

Courtney Voss^{1,9}, Sally Esmail^{1,9}, Xuguang Liu¹, Michael J. Knauer², Suzanne Ackloo³, Tomonori Kaneko¹, Lori Lowes², Peter Stogios⁴, Almagul Seitova³, Ashley Hutchinson³, Farhad Yusifov³, Tatiana Skarina⁴, Elena Evdokimova⁴, Peter Loppnau⁴, Pegah Ghiabi⁴, Taraneh Haijan⁴, Shanshan Zhong¹, Husam Abdoh², Benjamin D. Hedley², Vipin Bhayana², Claudio M. Martin^{5,6}, Marat Slessarev^{5,6}, Benjamin Chin-Yee⁷, Douglas D. Fraser^{5,6,8}, Ian Chin-Yee² and Shawn S-C. Li^{1*}

¹Department of Biochemistry and ²Department of Pathology and Laboratory Medicine, Western University, London, Ontario N6A 5C1; ³Structural Genomics Consortium and ⁴Department of Chemical Engineering and Applied Chemistry, University of Toronto, 101 College St., Toronto, Ontario M5G 1L7; ⁵Department of Medicine, Western University, London, Ontario N6A 5C1; ⁶London Health Sciences Centre, 800 Commissioners Rd E, London, Ontario N6A 5W9; ⁷Division of Hematology and ⁸Department of Pediatrics, Western University, London, Ontario N6A 5C1 Canada.

⁹These authors contributed equally to this work.

*Corresponding author: Email: sli@uwo.ca

Supplemental Table 1. Recombinant proteins and printing concentrations on the proteome array

Protein	Residues	Printed Concentration
IgG	Equitech Bio Inc SLH66-0001	200nM
Spike Receptor Binding Domain	ThermoFisher Scientific RP-87678 319-541	14μM
Spike Ectodomain	1-1208	0.5μM
NSP3-unique	Replicase polyprotein 1ab 413-676	10μM
NSP3-ADRP	Replicase polyprotein 1ab 1024-1192	10μM
NSP3-PLPro	Replicase polyprotein 1ab 1561-1878	10μM
NSP3-nucleic acid binding domain	Replicase polyprotein 1ab 1867-2034	10μM
NSP4-C terminal Domain	Replicase polyprotein 1ab 3173-3569	10μM
NSP5	Replicase polyprotein 1ab 3264-3569	10μM
NSP7	Replicase polyprotein 1ab 3860-3942	10μM
NSP8	Replicase polyprotein 1ab 3943-4140	10μM
NSP9	Replicase polyprotein 1ab 4141-4253	10μM
NSP10	Replicase polyprotein 1ab 4254-4392	10μM
NSP16	Replicase polyprotein 1ab 6799-7096	10μM
Nucleocapsid (full length)	RayBiotech 230-01104 1-419	5μM
Nucleocapsid Dimerization Domain	247-364	5μM
Nucleocapsid RNA Binding Domain	47-173	5μM

Supplemental Table 2. Patient cohort used in this study

Sample ID	ICU	Outcome	Severity
COV+5	N	Alive	Moderate
COV+7	N	Alive	Moderate
COV+8	N	Alive	Moderate
COV+10	N	Alive	Moderate
COV+11*	N	Alive	Moderate
COV+14	N	Alive	Moderate
COV+17*	N	Alive	Moderate
COV+18*	N	Fatal	Moderate
COV+19*	N	Alive	Moderate
COV+26	N	Alive	Moderate
COV+27	N	Alive	Moderate
COV+28	N	Alive	Moderate
COV+29	N	Alive	Moderate
COV+30*	N	Alive	Moderate
COV+33	N	Alive	Moderate
COV+34	N	Alive	Moderate
COV+51	N	Alive	Moderate
COV+52	N	Alive	Moderate
COV+53*	N	Alive	Moderate
COV+55	N	Alive	Moderate
COV+56	N	Alive	Moderate
COV+57*	N	Alive	Moderate
COV+58	N	Alive	Moderate
COV+59	N	Alive	Moderate
COV+62	N	Alive	Moderate
COV+63	N	Alive	Moderate
COV+64	N	Alive	Moderate
COV+65*	N	Alive	Moderate
COV+67	N	Alive	Moderate
COV+70	N	Alive	Moderate
COV+72	N	Alive	Moderate
COV+73	N	Alive	Moderate
COV+74	N	Alive	Moderate
COV+75	N	Alive	Moderate
COV+76	N	Alive	Moderate
COV+81*	N	Alive	Moderate
COV+82*	N	Alive	Moderate
COV+83	N	Alive	Moderate
COV+84*	N	Alive	Moderate
COV+85*	N	Alive	Moderate
COV+86*	N	Alive	Moderate

COV+87	N	Alive	Moderate
COV+88	N	Alive	Moderate
COV+89*	N	Alive	Moderate
COV+90*	N	Alive	Moderate
COV+91	N	Alive	Moderate
COV+1	Y	Alive	Severe
COV+2	Y	Fatal	Severe
COV+3	Y	Alive	Severe
COV+4	Y	Alive	Severe
COV+6	Y	Alive	Severe
COV+9	Y	Fatal	Severe
COV+12	N	Fatal	Severe
COV+13*	N	Fatal	Severe
COV+15*	Y	Fatal	Severe
COV+16*	N	Fatal	Severe
COV+20	Y	Fatal	Severe
COV+22	Y	Alive	Severe
COV+24	Y	Alive	Severe
COV+25*	Y	Alive	Severe
COV+31	Y	Alive	Severe
COV+32	Y	Alive	Severe
COV+35	Y	Fatal	Severe
COV+36	Y	Fatal	Severe
COV+37	Y	Fatal	Severe
COV+38	Y	Fatal	Severe
COV+39	Y	Alive	Severe
COV+40	Y	Fatal	Severe
COV+41	Y	Alive	Severe
COV+42	Y	Alive	Severe
COV+43	Y	Alive	Severe
COV+44	Y	Alive	Severe
COV+45	Y	Alive	Severe
COV+46	Y	Alive	Severe
COV+47	Y	Fatal	Severe
COV+48	Y	Alive	Severe
COV+49	Y	Fatal	Severe
COV+50	Y	Fatal	Severe
COV+54	Y	Alive	Severe
COV+60	Y	Alive	Severe
COV+61	Y	Alive	Severe
COV+66	Y	Alive	Severe
COV+68*	Y	Alive	Severe
COV+69*	N	Fatal	Severe

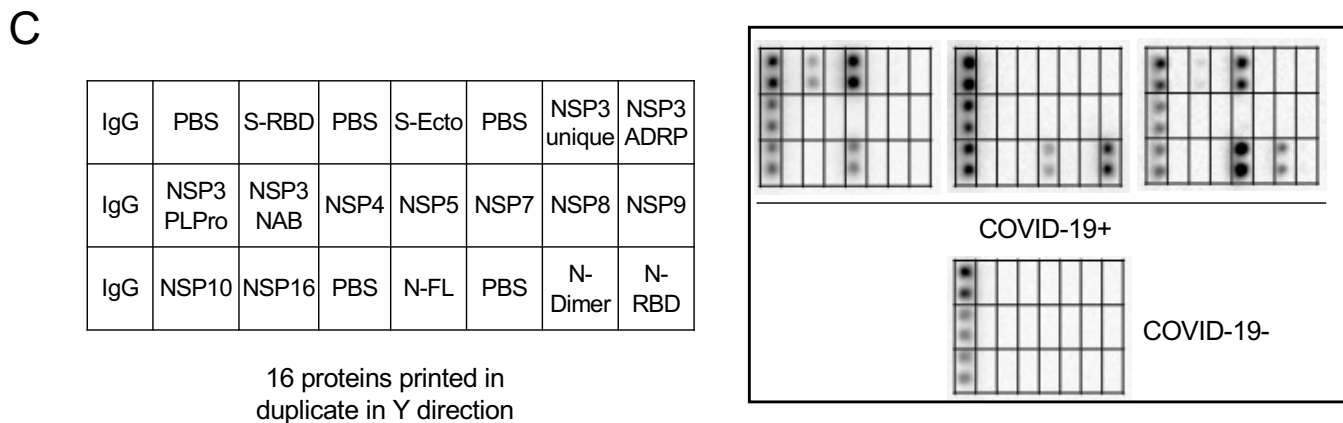
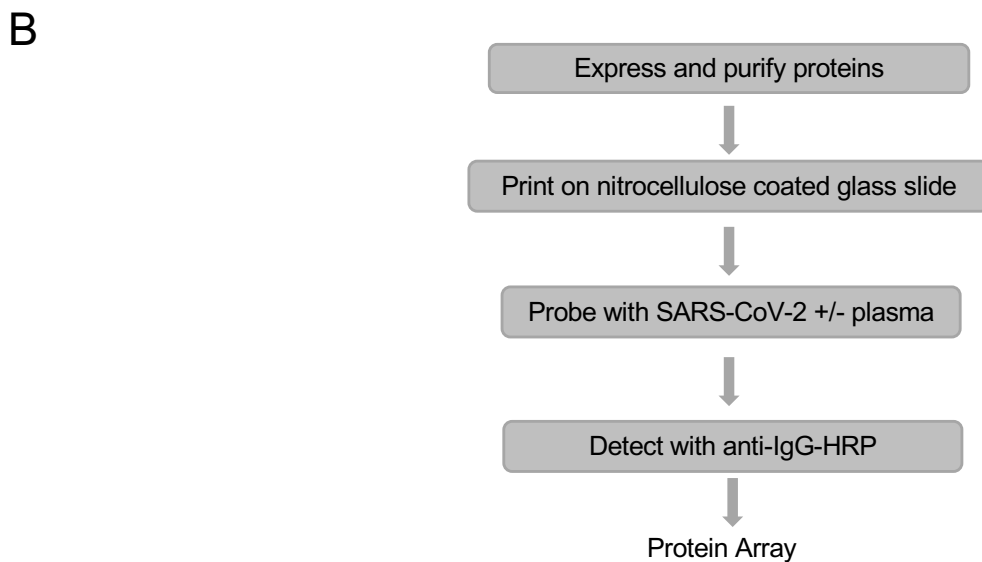
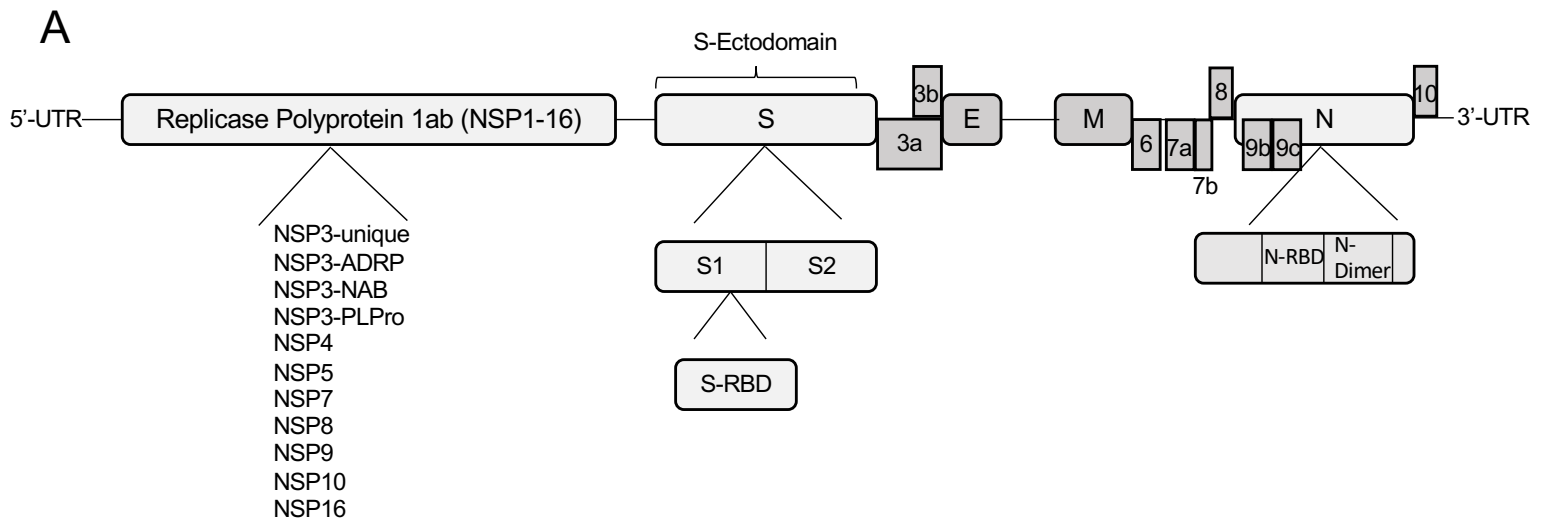
COV+71	Y	Fatal	Severe
COV+77	Y	Fatal	Severe
COV+78	N	Fatal	Severe
COV+79*	Y	Fatal	Severe
COV+80*	N	Fatal	Severe
COV-1	N	Alive	Negative
COV-2	N	Alive	Negative
COV-3	N	Alive	Negative
COV-4	N	Alive	Negative
COV-5	N	Alive	Negative
COV-6	N	Alive	Negative
COV-7	N	Alive	Negative
COV-8	N	Alive	Negative
COV-9	N	Alive	Negative
COV-10	N	Alive	Negative
COV-11	N	Alive	Negative
COV-35	N	Alive	Negative
COV-36	N	Alive	Negative
COV-37	N	Alive	Negative
COV-38	N	Alive	Negative
COV-39	N	Alive	Negative
COV-40	N	Alive	Negative
COV-41	N	Alive	Negative
COV-42	N	Alive	Negative
COV-43	N	Alive	Negative
COV-44	N	Alive	Negative
COV-45	N	Alive	Negative
COV-46	N	Alive	Negative
COV-47	N	Alive	Negative
COV-48	N	Alive	Negative
COV-49	N	Alive	Negative
COV-50	N	Alive	Negative

*No antibodies detected; sample was removed from analysis. Moderate cases defined as those with intensive care unit (ICU) admission; Severe cases defined as patients admitted to ICU or fatal; cases defined as Negative were determined by quantitative PCR analysis for SARS-CoV-2.

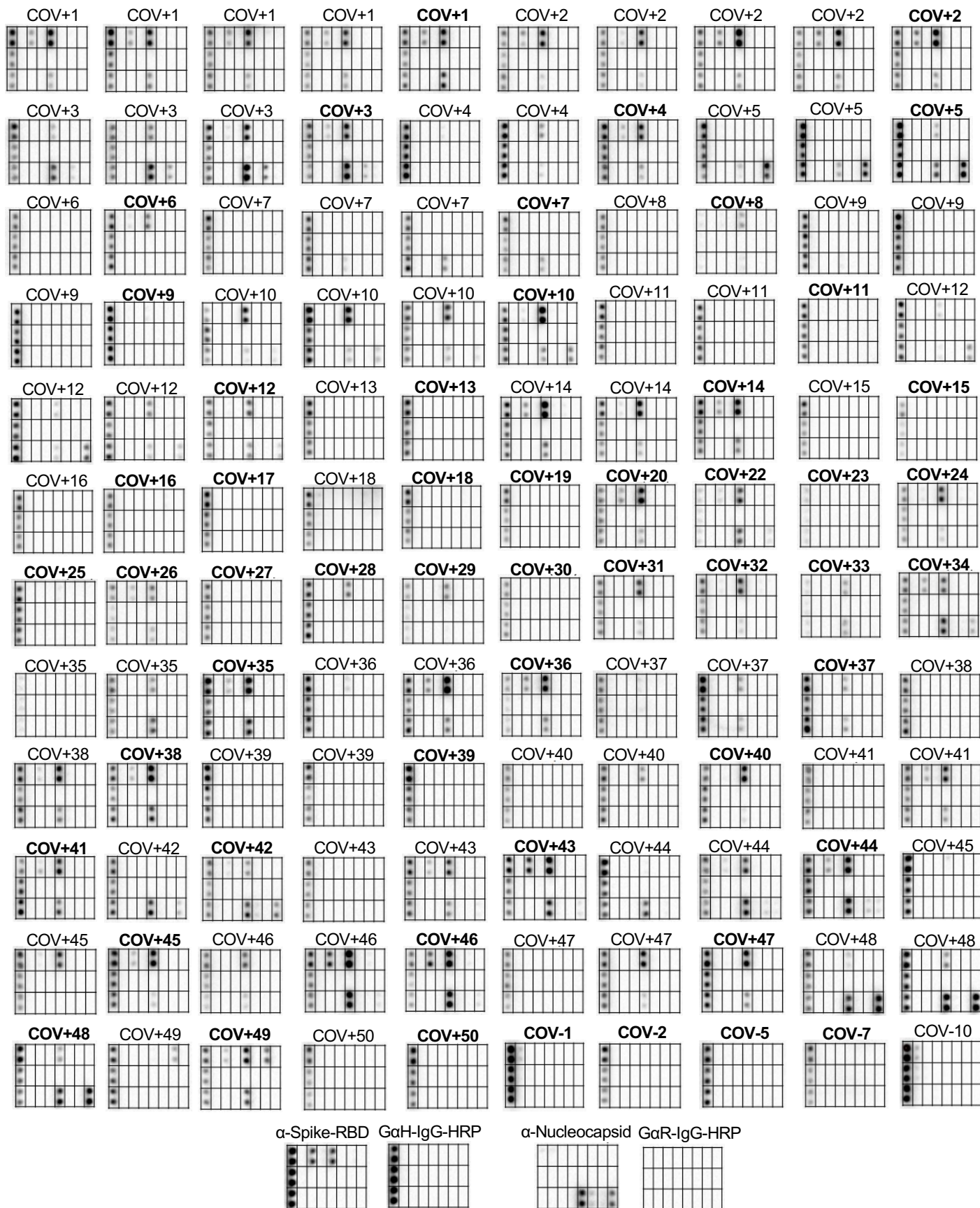
Supplemental Table 3. A list of literature reported epitopes examined in this study

Protein	Residues	Sequence	Source	Confirmed on Peptide Array
Spike	21-35	RTQLPPAYTNSFTRG	Δ	Y
Spike	181-195	GKQGNFKNLREFVFK	Δ	N
Spike	351-365	YAWNRRKISNCVADY	Δ	N
Spike	451-465	YLYRLFRKSNLKPFE	Δ	Y
Spike	801-815	NFSQILPDPSKPSKR	Δ	N
Spike	811-825	KPSKRSFIEDLLFNK	Δ	Y
Spike	881-895	TITSGWTFGAGAALQ	Δ	Y
Spike	1041-1055	DFCGKGYHLMSFPQS	Δ	N
Spike	1091-1105	REGVFVSNNGTHWFVT	Δ	N
Spike	612-623	YQDVNCTEVPVA	δ	N
Spike	491-504***	PLQSYGFQPTNGVGT	δ	N
Spike	802-819	FSQILPDPSKPSKRSFIE	δ	Y
Spike	888-902	FGAGAALQIPFAMQM	δ	N
Spike	895-909	QIPFAMQMAYRFNGI	δ	N
Spike	287-305	DAVDCALDPLSETKCTLKS	δ	N
Spike	306-318***	FTVEKGIYQTSNGGGTPCS	δ	N
Spike	588-598***	NGGGTPCSFGGVSVI	δ	N
Spike	524-545	VCGPKKSTNLVKNKCVNFNFN	δ	N
Spike	545-565	GLTGTGVLTESNKKFLPFQQF	δ	Y
Spike	566-587	GRDIADTTDAVRDPQTLEILDI	δ	Y
Spike	601-611	GTNTSNQVAVL	δ	N
Spike	624-640	IHADQLTPTWRVYSTGS	δ	Y
Nucleocapsid	61-75	KEDLKFRGQGVPIIN	Δ	N
Nucleocapsid	91-105	TRRIRGGDGKMKDLS	Δ	N
Nucleocapsid	161-175	LPQGTTLPKGFYAEG	Δ	Y
Nucleocapsid	201-215	SSRGTSPARMAGNGG	Δ	Y
Nucleocapsid	221-235	LLLLDRLNQLESKMS	Δ	Y
Nucleocapsid	251-265	AAEASKKPRQKRTAT	Δ	Y
Nucleocapsid	311-325	ASAFFGMSRIGMEVT	Δ	N
Nucleocapsid	361-375	KTFPPTPEPKDKKKK	Δ	Y
Nucleocapsid	371-385	DKKKKADETQALPQR	Δ	Y
Nucleocapsid	381-395	ALPQRQKKQQTVTLL	Δ	Y
Nucleocapsid	391-405	TVTLLPAADLDDFSK	Δ	N
Nucleocapsid	41-61	RPQGLPNNTASWFTALTQH GK	δ	Y
Nucleocapsid	153**	NNNAATVLQLPQGTTLPKG	δ	Y
Nucleocapsid	355-375	KHIDAYKTFPPTPEPKDKKKK	δ	Y
Nucleocapsid	381-401**	QPLPQRQKKQPTVTLLPAADM	δ	Y
Nucleocapsid	368-391**	PKKDKKKKTDEAQLPQRQKKQP	δ	Y

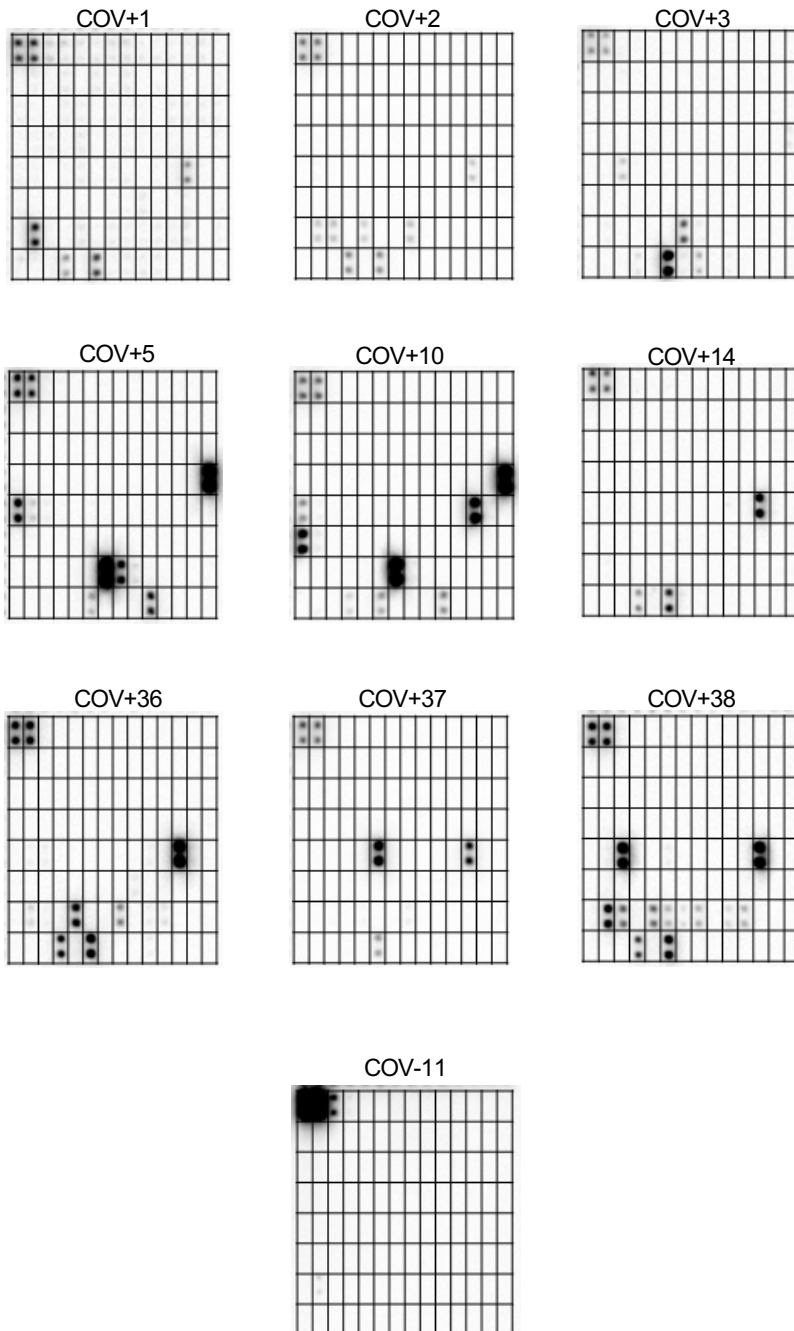
Δ Wang *et al.* (2020) *ACS Cent Sci* **6**: 2238-2249; δ Grifoni *et al.* (2020) *Cell Host Microbe* **27**: 671-680 e2. ** Based off SARS CoV sequence; *** Non-linear peptide from Grifoni *et al.* 2020



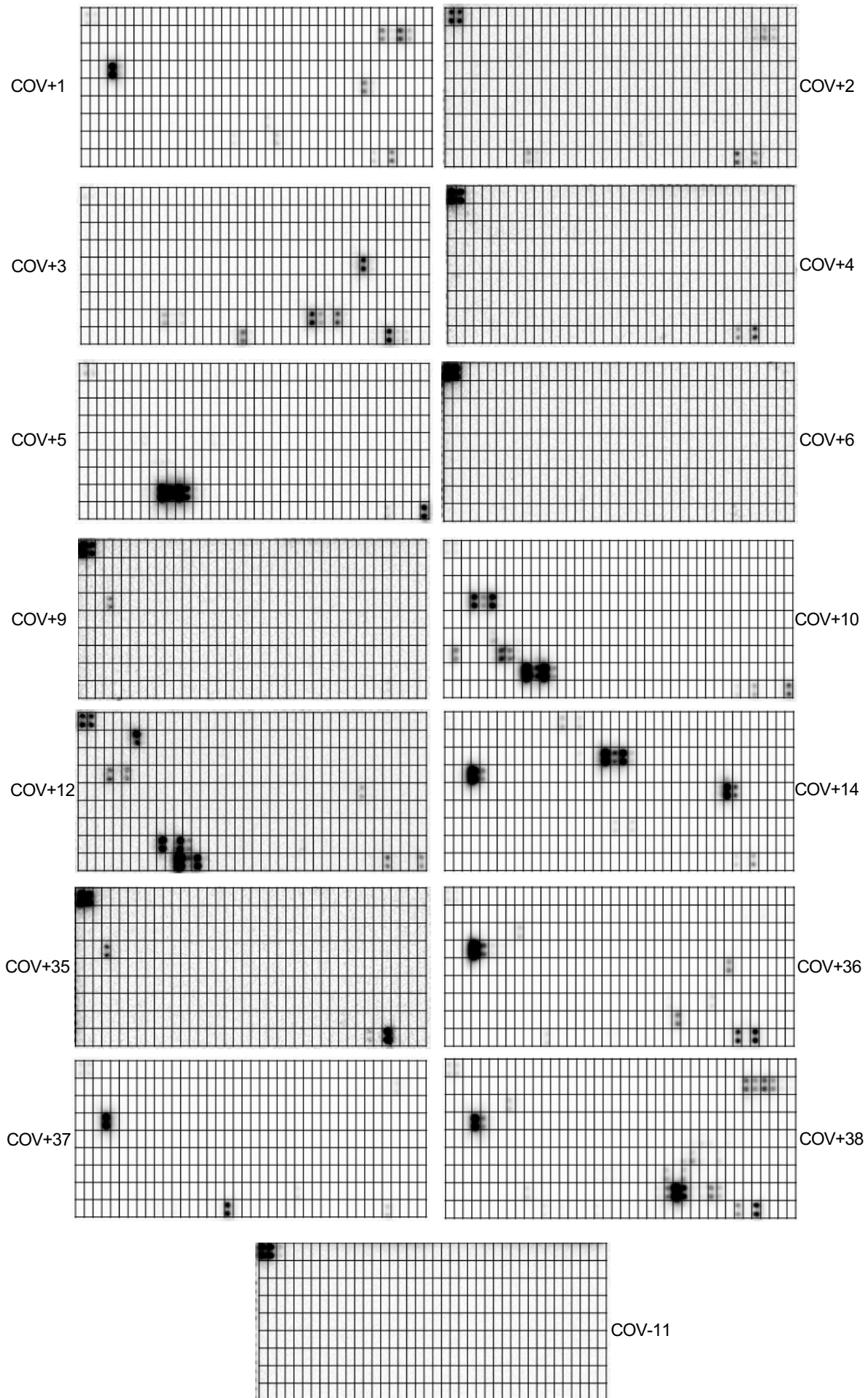
Supplemental Figure 1. Characterizing SARS-CoV-2 antibody responses using a proteome array. (A) A schematic diagram showing the major proteins of SARS-CoV-2. (B) Workflow for the production and screening of a SARS-CoV-2 proteome array. (C) Left, layout of the SARS-CoV-2 proteome array; Right, representative images of the proteome array probed with SARS-CoV-2-positive or -negative (based on NAT) plasma.



Supplemental Figure 2. Antibody responses profiled using the SARS-CoV-2 proteome array. Twenty-four copies of the grid were printed on a single nitrocellulose coated glass slide. Serial patient samples are shown sequentially and bold are the patient samples subsequently analyzed on the master array. All images shown were taken with 5.2 second exposure time.



Supplemental Figure 3. Screening of a peptide array containing literature-reported epitopes. Three copies of the grid were printed on a single nitrocellulose coated glass slide. Images shown were taken with 5.2 second (COV+) or 60 second (COV-) exposure time.



Supplemental Figure 4. Images from epitope mapping using peptide walking array. Two copies of the grid were printed on a single nitrocellulose coated glass slide. Images shown were taken with 5.2 second (COV+) or 60 second (COV-) exposure times.

A

IgG	N-6	N-156	N-221	N-246	N-361	N-381	S-21
IgG	S-176	S-451	S-551	S-671	S-811	S-881	S-1146
IgG	S-1166	S-1216	N-Dimer	N-RBD	N-FL	S-RBD	S-Ecto

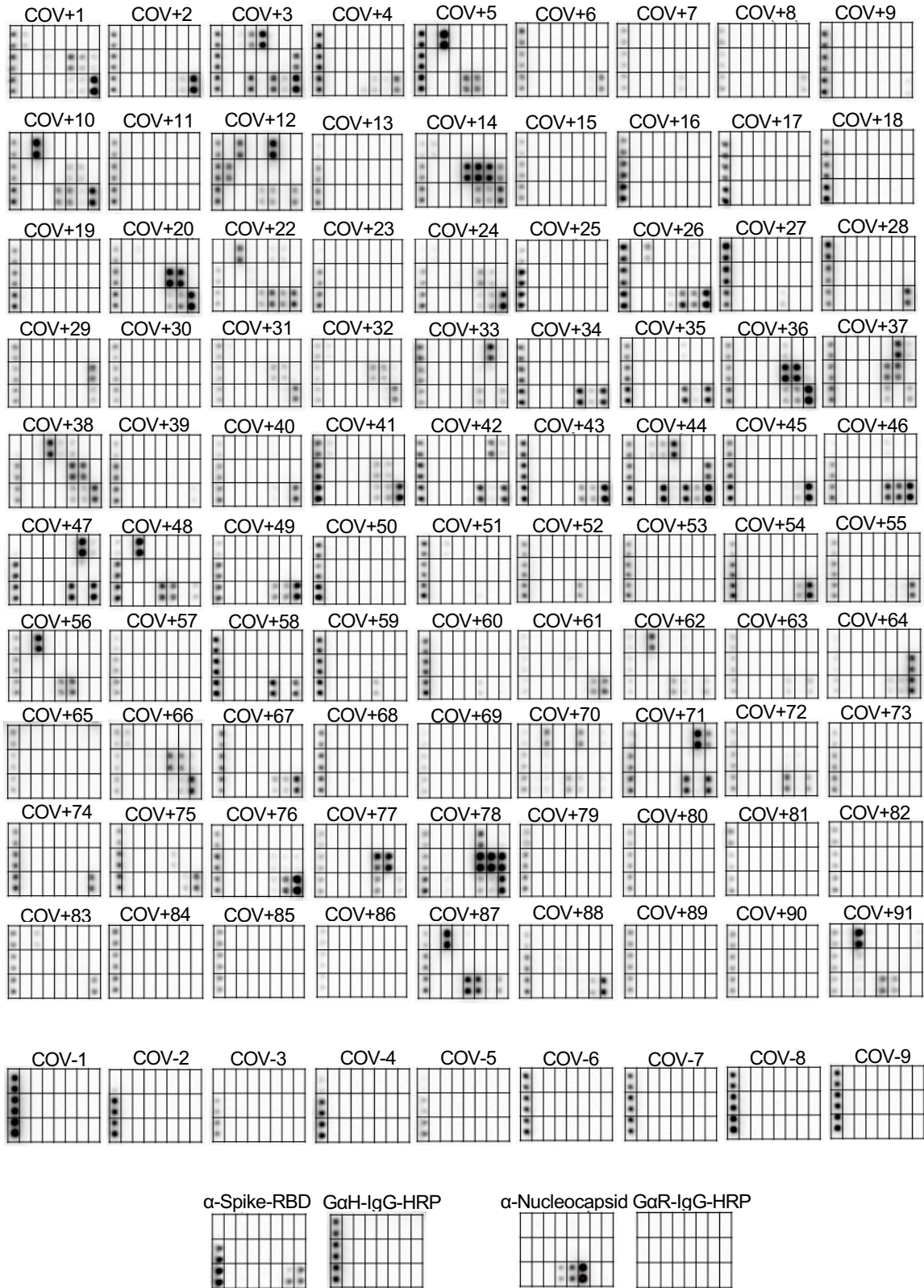
B

Peptide	Sequence
N-6	PQNQRNAPRITFGGP
N-156	AIVLQLPQGTTLPKG
N-221	LLLLDRLNQLESKMS
N-246	VTKKSAEASKKPRQ
N-361	KTFPPTEPKKDKKKK
N-381	ALPQRQKKQQTVTLL
S-21	RTQLPPAYTNSFTRG
S-176	LMDLEGKQGKFNLR
S-451	YLYRLFRRKSNLKPFE
S-551	VLTESNKKFLPFQQF
S-671	CASYQTQNSPRRAR
S-811	KPSKRSEFIEDLLFNK
S-881	TITSGWTFGAGAALQ
S-1146	DSFKEELDKYFKNHT
S-1166	LGDISGINASVVNIQ
S-1216	IWLGFIAGLIAVMV

C

Protein	Printed Concentration
IgG	100nM
Nucleocapsid Dimerization Domain	15 μ M
Nucleocapsid RNA binding domain	1 μ M
Nucleocapsid (Full Length)	2 μ M
Spike-Receptor Binding Domain	14 μ M
Spike-Ectodomain	0.2 μ M

Supplemental Figure 5. SARS-CoV-2 Master Epitope Array. (A) Identity and position of peptides and proteins in the master array. All spots are printed in duplicate in the Y direction. (B) Peptide identities and sequences. (C) Protein concentrations printed on the array.



Supplemental Figure 6. Profiling antibody responses using the master array. Twenty-four copies of the grid were printed on a single nitrocellulose coated glass slide. All images shown were taken with 9.4 second exposure time.