

1 **SUPPLEMENTAL APPENDIX**

2 **Supplement to:**

3 **Therapeutic potential of convalescent plasma and SARS-CoV-2 hyperimmune**
4 **immunoglobulins against BQ.1, BQ.1.1 and XBB variants**

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10 **TABLE OF CONTENTS**

11 **CONTENTS**

12 **1. Materials and Methods**

13 Samples and study design

14 Ethics Statement

15 Neutralization assay

16 Quantification and statistical analysis

17 **2. Table S1.** SARS-CoV-2 variants mutations introduced in the spike plasmid for production of
18 SARS-CoV-2 pseudovirions for analysis in PsVNA

19 **3. Table S2.** Summary of CP and IVIG lots and neutralization titers

20 **4. Acknowledgments**

21 **5. Supplementary References**

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23 **MATERIALS AND METHODS**

24 **Samples and study design**

25 IVIG products approved in the United States are polyclonal antibody preparations made
26 from 10,000 or more U.S. plasma donors and may include cold alcohol fractionation (Cohn-
27 Oncley), anion-exchange and size-exclusion chromatographies. The final product is sterile-
28 filtered IgG (>95%) and formulated at 100 mg/mL. Twenty intravenous immunoglobulin batches
29 were produced from plasma collected prior to August 2019 (2019-IVIG) and eight IVIG lots made
30 from plasma donations in 2020 (2020-IVIG) and manufactured between October 2020 and
31 January 2021 (each lot derived from >10,000 donors), were obtained from six manufacturers.

32 Nineteen hCoV-2IG batches prepared from COVID-19 CP (~200-1000 donors per lot)
33 were obtained/purchased from four commercial companies for blinded antibody analysis. The
34 plasma units used in the manufacturing of the hCoV-2IG batches were collected in 2020 prior to
35 emergence of the Delta and Omicron VOCs or prior to availability of COVID-19 vaccines.

36 Eight random CP lots were obtained from recovered COVID-19 patients between May-
37 September 2020 (at least 30-days post-recovery) prior to COVID-19 vaccinations. At the time of
38 collection SARS-CoV-2 D614G was the predominant strain in the US. Nine CP lots were
39 collected in February 2022 from recovered individuals following Omicron breakthrough infections
40 (probably BA.1), who received at least two doses of COVID-mRNA vaccination.

41

42 **Ethics Statement**

43 This study was approved by the Food and Drug Administration's Research Involving
44 Human Subjects Committee (RIHSC #2020-04-02). This study complied with all relevant ethical

45 regulations for work with human participants, and informed consent was obtained. Samples were
46 collected from adult subjects who provided informed consent to participate in the study. All
47 assays performed fell within the permissible usages in the original informed consent.

48

49 **Neutralization assay**

50 Samples were evaluated in a qualified SARS-CoV-2 pseudovirion neutralization assay
51 (PsVNA) using SARS-CoV-2 WA1/2020 strain and circulating Omicron subvariants: BA.4/BA.5,
52 BA.2.75, BA.2.75.2, BQ.1, BQ.1.1 and recombinant XBB. The mutations in spike protein of these
53 Omicron subvariants are shown in Supplementary Table S1. SARS-CoV-2 neutralizing activity
54 measured by PsVNA correlates with PRNT (plaque reduction neutralization test with authentic
55 SARS-CoV-2 virus) in previous studies (1-3). However, some antibodies targeting the N-terminal
56 domain of SARS-Cov-2 spike may not show neutralization in the pseudovirus neutralization
57 assay.

58 Neutralization assays were performed as previously described (2, 4). Briefly, 50 μ L of
59 SARS-CoV-2 S pseudovirions (counting \sim 200,000 relative light units) were pre-incubated with
60 an equal volume of medium containing serial dilutions (starting at 1:10) of all samples at room
61 temperature for 1 h. Then, 50 μ L of virus-antibody mixtures were added to 293T-ACE2-
62 TMPRSS2 cells [10^4 cells/50 μ L; gift from Carol Weiss (1)] in a 96-well plate. The input virus with
63 all SARS-CoV-2 strains was the same (2×10^5 relative light units/50 μ L/well). After a 3 h
64 incubation, fresh medium was added to the wells. Cells were lysed 24 h later, and luciferase
65 activity was measured using One-Glo luciferase assay system (Promega). The assay of each
66 sample was performed in duplicate, and the 50% neutralization titer was calculated using Prism
67 9 (GraphPad Software). The limit of detection for the neutralization assay is 1:20. Two

68 independent biological replicate experiments were performed for each sample and variation in
69 PsVNA50 titers was <10% between replicates.

70

71 **Quantification and statistical analysis**

72 Descriptive statistics were performed to determine the geometric mean titer values and
73 were calculated using GraphPad. All experimental data to compare differences among groups
74 were analyzed using Ordinary one-way ANOVA with Tukey's pairwise multiple comparison test
75 in GraphPad Prism version 9.3.1. To ensure robustness of the results, absolute measurements
76 were log₂-transformed before performing the analysis. Correlation and regression analyses
77 were performed by computing Pearson correlation (r), assuming that the data were sampled
78 from Gaussian Distribution, in GraphPad Prism version 9.3.1, to correlate the neutralization titers
79 of WA-1 strain with each Omicron subvariant: BA.2.75.2, BA.2.75, BA.4, BQ.1, BQ.1.1 and XBB,
80 for all the SARS-CoV-2 positive CP and IVIG samples. For each graph, we show the value of r
81 (not R square) and a confidence interval of 95%, with two-tailed significance (p) values.

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83 **Data sharing.** All data needed to evaluate the conclusions in the article are present in the
84 manuscript.

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Supplementary Table S1: SARS-CoV-2 variants mutations introduced in the spike plasmid for production of SARS-CoV-2 pseudovirions for analysis in PsVNA.

SARS-CoV-2 variant	Mutations constructed in the spike plasmids
Omicron (BA.2.75)	BA.2 spike mutations (T19I, delL24, delP25, delP26, A27S, G142D, V213G, G339D, S371F, S373P, S375F, T376A, D405N, R408S, K417N, N440K, S477N, T478K, E484A, Q493R, Q498R, N501Y, Y505H, D614G, H655Y, N679K, P681H, N764K, D796Y, Q954H, N969K) + K147E, W152R, F157L, I210V, G257S, D339H, G446S, N460K, Q493reversion
Omicron (BA.2.75.2)	BA.2 spike mutations (T19I, delL24, delP25, delP26, A27S, G142D, V213G, G339D, S371F, S373P, S375F, T376A, D405N, R408S, K417N, N440K, S477N, T478K, E484A, Q493R, Q498R, N501Y, Y505H, D614G, H655Y, N679K, P681H, N764K, D796Y, Q954H, N969K) + K147E, W152R, F157L, I210V, G257S, D339H, R346T, G446S, N460K, F486S, Q493reversion, D1199N
Omicron (BA.4/BA.5)	T19I, delL24, delP25, delP26, A27S, del69 70, G142D, V213G, G339D, S371F, S373P, S375F, T376A, D405N, R408S, K417N, N440K, L452R, S477N, T478K, E484A, F486V, Q498R, N501Y, Y505H, D614G, H655Y, N679K, P681H, N764K, D796Y, Q954H, N969K
Omicron (BQ.1)	BA.4/BA.5 spike mutations (T19I, delL24, delP25, delP26, A27S, del69 70, G142D, V213G, G339D, S371F, S373P, S375F, T376A, D405N, R408S, K417N, N440K, L452R, S477N, T478K, E484A, F486V, Q498R, N501Y, Y505H, D614G, H655Y, N679K, P681H, N764K, D796Y, Q954H, N969K) + K444T and N460K.
Omicron (BQ.1.1)	BQ.1 spike mutations (T19I, delL24, delP25, delP26, A27S, del69 70, G142D, V213G, G339D, S371F, S373P, S375F, T376A, D405N, R408S, K417N, N440K, L452R, S477N, T478K, E484A, F486V, Q498R, N501Y, Y505H, D614G, H655Y, N679K, P681H, N764K, D796Y, Q954H, N969K, K444T and N460K) + R346T.
Omicron (XBB)	BA.2 spike mutations (T19I, delL24, delP25, delP26, A27S, G142D, V213G, G339D, S371F, S373P, S375F, T376A, D405N, R408S, K417N, N440K, S477N, T478K, E484A, Q493R, Q498R, N501Y, Y505H, D614G, H655Y, N679K, P681H, N764K, D796Y, Q954H, N969K) + V83A, Del144, H146Q, Q183E, V213E, G339H, R346T, L368I, V445P, G446S, N460K, F486S, F490S, R493Q

Table S2: Neutralization titers of convalescent plasma, IVIG and hCoV-2IG against SARS-CoV-2 variants*

	WA-1	Omicron BA.2.75	Omicron BA.2.75.2	Omicron BA.4/BA.5	Omicron BQ.1	Omicron BQ.1.1	Omicron XBB
IVIG batches produced in 2019 prior to COVID-19							
2019-IVIG-1	10	10	10	10	10	10	10
2019-IVIG-2	10	10	10	10	10	10	10
2019-IVIG-3	10	10	10	10	10	10	10
2019-IVIG-4	10	10	10	10	10	10	10
2019-IVIG-5	10	10	10	10	10	10	10
2019-IVIG-6	10	10	10	10	10	10	10
2019-IVIG-7	10	10	10	10	10	10	10
2019-IVIG-8	10	10	10	10	10	10	10
2019-IVIG-9	10	10	10	10	10	10	10
2019-IVIG-10	10	10	10	10	10	10	10
2019-IVIG-11	10	10	10	10	10	10	10
2019-IVIG-12	10	10	10	10	10	10	10
2019-IVIG-13	10	10	10	10	10	10	10
2019-IVIG-14	10	10	10	10	10	10	10
2019-IVIG-15	10	10	10	10	10	10	10
2019-IVIG-16	10	10	10	10	10	10	10
2019-IVIG-17	10	10	10	10	10	10	10
2019-IVIG-18	10	10	10	10	10	10	10
2019-IVIG-19	10	10	10	10	10	10	10
2019-IVIG-20	10	10	10	10	10	10	10
IVIG batches produced in 2020							
2020-IVIG-1	31	10	10	10	10	10	10
2020-IVIG-2	25	10	10	10	10	10	10
2020-IVIG-3	73	10	10	10	10	10	10
2020-IVIG-4	36	10	10	10	10	10	10
2020-IVIG-5	64	10	10	10	10	10	10
2020-IVIG-6	32	10	10	10	10	10	10
2020-IVIG-7	24	10	10	10	10	10	10
2020-IVIG-8	21	10	10	10	10	10	10
Convalescent plasma batches produced from COVID-19 survivors collected in 2020							
2020-CP-1	538	10	10	10	10	10	10
2020-CP-2	415	10	10	10	10	10	10
2020-CP-3	10	10	10	10	10	10	10
2020-CP-4	302	10	10	10	10	10	10
2020-CP-5	28	10	10	10	10	10	10
2020-CP-6	104	10	10	10	10	10	10
2020-CP-7	122	10	10	10	10	10	10
2020-CP-8	1343	10	79	95	10	10	10
Convalescent plasma batches produced from COVID-19 survivors collected in 2022							
2022-CP-1	1748	327	198	345	136	121	116
2022-CP-2	1067	47	50	53	10	10	10
2022-CP-3	21306	1373	1214	4841	136	61	54
2022-CP-4	567	10	10	10	10	10	10
2022-CP-5	677	10	10	40	10	10	10
2022-CP-6	854	24	58	87	10	10	10
2022-CP-7	973	155	196	163	42	39	37
2022-CP-8	1505	176	169	145	45	42	30
2022-CP-9	603	41	76	41	10	10	10
hCoV-2IG batches produced from COVID-19 convalescent plasma donors							
hCoV-2IG-1	2067	43	52	128	64	53	36
hCoV-2IG-2	1664	49	39	71	10	10	10
hCoV-2IG-3	1602	27	29	81	39	44	39
hCoV-2IG-4	1833	10	22	47	10	10	10
hCoV-2IG-5	2302	126	62	103	10	10	10
hCoV-2IG-6	920	10	10	46	10	10	10
hCoV-2IG-7	3243	430	297	205	48	63	40
hCoV-2IG-8	1730	55	39	99	44	33	28
hCoV-2IG-9	819	20	10	107	34	31	10
hCoV-2IG-10	1428	46	50	99	29	32	10
hCoV-2IG-11	737	23	10	78	26	32	28
hCoV-2IG-12	917	24	28	85	26	37	10
hCoV-2IG-13	1055	64	28	81	10	10	10
hCoV-2IG-14	1896	63	60	104	30	42	25
hCoV-2IG-15	1565	36	31	59	10	10	10
hCoV-2IG-16	2351	84	73	143	32	54	31
hCoV-2IG-17	2486	90	67	151	46	43	25
hCoV-2IG-18	3162	124	82	154	49	57	39
hCoV-2IG-19	1646	29	47	159	53	55	42

* PsVNA titer Cut-off value: 1:10.

87 **Acknowledgments**

88

89 We thank Basil Golding and Keith Peden at FDA for review of the manuscript. We thank Carol
90 Weiss (FDA) for providing plasmid clones expressing SARS-CoV-2 spike variants.

91

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