Supporting Information

Potent Anti-SARS-CoV-2 Efficacy of COVID-19 Hyperimmune Globulin from Vaccine-Immunized Plasma

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Figure S1. Assessment of the antiviral activity of IVIG against SARS-CoV-2 pseudotyped viruses *in vitro*. (a-c) IVIG showed no neutralization effect against pseudotyped SARS-CoV-2 Wuhan-1 (a), Beta (b), and Delta (c) strains. Pseudotyped viruses were pre-incubated with serial dilutions of COVID-HIG at different concentrations for 1 h at 37° C. Next, Huh-7 cells were incubated with the pseudotyped viruses for 24 h. Luciferase was detected to assess infection. The y-axis represents percent inhibition. Data are shown as mean \pm SD of three independent experiments (n = 3). IVIG, human immune globulin intravenous.



Figure S2. Plaque reduction neutralization assay of COVID-HIG. COVID-HIG-001 (a), COVID-HIG-002 (b), and COVID-HIG-003 (c) significantly inhibited infection by SARS-CoV-2 WIV04, Beta, and Delta strains in Vero E6 cells. Maltose was used as a negative control. Viruses were incubated with COVID-HIG at different concentrations at 37°C for 1 h. Thereafter, Vero E6 cells were infected with SARS-CoV-2 WIV04, Beta and Delta strains. The infected cells were stained with hematoxylin/eosin at 48 h (Beta and Delta strains) or 72 h (WIV04 strain) post-infection. WIV04, nCoV-2019BetaCoV/Wuhan/WIV04/2019.

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Figure S3. Gross observation of the lungs of mice treated with COVID-HID 6 days after SARS-COV-2 infection. Mice in the adjuvant control group had a large range of diffuse crimson lesions in the lungs, whereas the area of crimson lesions in the lungs of mice in the COVID-HIG prevention and treatment groups was significantly reduced. Maltose adjuvant-treated mice were used as the control group. Scale bar = 0.5 cm.



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Figure S4. Virological analyses of treated and untreated mouse lungs using immunofluorescence staining of SARS-CoV-2 nucleocapsid proteins. Changes in virus protein expression (red fluorescence) in the mouse lungs. Tissues were dissected and collected for fixation, embedding, slicing, and immunofluorescence detection. Maltose-treated mice were used as the adjuvant control group. The figure on the right (scale bar = 100 nm) is an enlarged view of the white dotted box in the figure on the left (scale bar = 1 μ m).

| Lot | Product | Total protein (mg/mL) | IgG (mg/mL) |
|---------------|-----------|-----------------------|-------------|
| | PBP | 64.7 | 10.0 |
| COVID-HIG-001 | COVID-HIG | 53.2 | 52.1 |
| COVID-HIG-002 | PBP | 63.0 | 9.8 |
| | COVID-HIG | 53.0 | 52.0 |
| COVID-HIG-003 | PBP | 64.9 | 8.8 |
| | COVID-HIG | 54.2 | 53.5 |

Table S1. Total protein and IgG concentrations of PBP and COVID-HIG

IgG, immunoglobulin G. PBP, pooled BBIBP-CorV plasma.

| COVID-HIG Lot | Distribution of the four IgG subclasses (%) * | | | | | |
|---------------|---|----------------------|---------------------|---------------------|--|--|
| | IgG1 | IgG2 | IgG3 | IgG4 | | |
| COVID-HIG-001 | 67.38% | 28.09% | 2.61% | 1.93% | | |
| COVID-HIG-002 | 64.91% | 29.64% | 3.32% | 2.13% | | |
| COVID-HIG-003 | 65.27% | 29.05% | 3.51% | 2.17% | | |
| Mean ± SD | $65.85\% \pm 1.33\%$ | $28.93\% \pm 0.78\%$ | $3.14\% \pm 0.48\%$ | $2.08\% \pm 0.13\%$ | | |

Table S2. IgG subclass distribution of COVID-HIG

*Immunoturbidimetry was used for detection.

 Table S3. Summary values of binding affinity, blocking potency, and neutralization

 potency of COVID-HIG-003

| Strain | K _D (nM) | | IC ₅₀ (nM) of ELISA | IC ₅₀ (nM) of PBNA | PRNT ₅₀ (nM) | Ratio (K _D /PRNT ₅₀) |
|-----------------------|---------------------|-------|-----------------------------------|-----------------------------------|-------------------------|--|
| | S protein | 4.60 | 2.27×10^{3} | 0.55×10 ³ (Wuhan-1) | 0.19×10^{3} | 2.4×10^{-2} |
| WIV04 (Or Wuhan-1) | NTD | 14.80 | | | | 7.8×10^{-2} |
| | NP | 14.30 | | | | N/A |
| | RBD | 99.50 | | | | 5.2×10^{-1} |
| Beta | RBD | 5.33 | 4.66×10^{3} | 3.95×10^{3} | 1.44×10^{3} | 3.7×10^{-3} |
| Delta | RBD | 21.90 | 1.13×10^{3} | 1.52×10^{3} | 1.61×10^{3} | 1.4×10^{-2} |

WIV04, nCoV-2019BetaCoV/Wuhan/WIV04/2019 strain. S, spike. NTD, N-terminal domain of S protein. NP, nucleocapsid protein. RBD, receptor-binding domain of S protein. K_D , equilibrium dissociation constant. Competitive ELISA assays were performed to calculate the 50% inhibitory concentration values for the blocking ability of COVID-HIG. PBNA, pseudotyped virus-based neutralization assay. PRNT₅₀, the 50% plaque reduction neutralization test concentration of COVID-HIG against strains. N/A, not applicable.

Table S4. Neutralization potency of PBP and COVID-HIG against the pseudotypedSARS-CoV-2 Wuhan-1 strain

| Lot | Product | IC ₅₀ (mg/mL) |
|---------------|-----------|--------------------------|
| | PBP | 0.2759 |
| COVID-HIG-001 | COVID-HIG | 0.0856 |
| | PBP | 0.2537 |
| COVID-HIG-002 | COVID-HIG | 0.0693 |
| | PBP | 0.4582 |
| COVID-HIG-005 | COVID-HIG | 0.0827 |

Wuhan-1, SARS-CoV-2 Wuhan-Hu-1 strain. PBP, pooled BBIBP-CorV plasma.

| Cassia | | Severity grade [*] | | La diat | | | |
|---------------------|---------------|-----------------------------|----------|----------|-------|-----------------------|---------------|
| ID : | ID # | Alveolar wall | Alveolar | Bronchus | Blood | dual [#] mea | Group mean |
| Maltose | 1 | | | +++ | +++ | 5 | 5.0 |
| Waltose | $\frac{1}{2}$ | +++ | +++ | +++ | +++ | 5 | 5.0 |
| | 3 | +++ | +++ | +++ | +++ | 6 | |
| | 4 | +++ | ++ | +++ | +++ | 5 | |
| | 5 | +++ | +++ | +++ | +++ | 6 | |
| | 6 | +++ | ++ | +++ | +++ | 5 | |
| | 7 | ++ | ++ | ++ | +++ | 4 | |
| | 8 | + | + | ++ | ++ | 3 | |
| COVID-HIG | 1 | + | - | ++ | ++ | 2 | 2.8 |
| 300 mg/kg, -24h | 2 | ++ | ++ | + | - | 2 | |
| | 3 | ++ | ++ | ++ | ++ | 3 | |
| | 4 | ++ | ++ | ++ | +++ | 4 | |
| | 5 | - | - | - | + | 1 | |
| | 6 | + | + | ++ | +++ | 3 | |
| | 7 | ++ | ++ | +++ | ++ | 4 | |
| | 8 | + | ++ | ++ | +++ | 3 | |
| COVID-HIG | 1 | + | + | +++ | +++ | 4 | 2.9 |
| 300 mg/kg, multiple | 2 | + | - | + | ++ | 2 | |
| | 3 | ++ | + | ++ | +++ | 3 | |
| | 4 | + | ++ | ++ | +++ | 4 | |
| | 5 | ++ | +++ | +++ | +++ | 5 | |
| | 6 | + | - | + | + | 1 | |
| | 7 | ++ | ++ | ++ | ++ | 3 | |
| | 8 | - | - | - | + | 1 | |
| COVID-HIG | 1 | + | + | ++ | +++ | 3 | 3.2 |
| 600 mg/kg, +2 h | 2 | + | ++ | +++ | ++ | 4 | |
| | 3 | + | ++ | ++ | +++ | 4 | |
| | 4 | +++ | ++ | ++ | ++ | 4 | |
| | 5 | + | - | + | + | 1 | |
| | 6 | + | + | + | ++ | 2 | |
| | / | + | + | +++ | ++ | 5 | |
| | 8 | ++ | +++ | +++ | +++ | 5 | 2.0 |
| 200 mg/kg + 2 h | 1 | ++ | + | ++ | +++ | 4 | 3.9 |
| 500 mg/kg, +2 m | 2 | +++ | +++ | +++ | +++ | 0 | |
| | 3 4 | + | - | + | + | 1 | |
| | + 5 | ++ ++ | ++ ++ | ++ ++ | +++ | 4 | |
| | 5 | ++ | ++ | ++ | +++ | + 1 | |
| | 0 7 | | ++++ | ++++ | ++ | т Д | |
| | 8 | ++ | + | ++ | +++ | т Д | |
| COVID-HIG | 1 | +++ | ++ | +++ | +++ | 5 | 36 |
| 100 mg/kg + 2 h | 2 | +++ | +++ | ++ | +++ | 5 | 2.0 |
| | 3 | + | + | ++ | +++ | 3 | |
| | 4 | ' ++ | , ++ | ++ | +++ | 4 | |
| | 5 | +++ | ++ | ++ | +++ | 5 | |
| | 6 | ++ | ++ | + | +++ | 4 | |
| | 7 | + | - | - | + | 1 | |
| | 8 | + | + | + | ++ | 2 | |

Table S5. Gross pathological findings

*In a single score, - indicates no pathological changes; + indicates mild lesions; ++ indicates moderate lesions; +++ indicates severe lesions. #The comprehensive score was assigned

according to the standards introduced in the Materials and Methods; 0, no lesion; 1–2, mild lesions; 3–4, moderate lesions; 5–6, severe lesions.

-24 h, COVID-HIG prophylactic groups; +2 h, COVID-HIG treatment groups.

Table S6. RBD-IgG titer of PBP and COVID-HIG

| Lot | Product | RBD-IgG Titer |
|---------------|-----------|----------------------|
| | PBP | 1:944 |
| | COVID-HIG | 1:5,184 |
| | PBP | 1:674 |
| COVID-HIG-002 | COVID-HIG | 1:5,919 |
| | PBP | 1:551 |
| COVID-HIG-003 | COVID-HIG | 1:4,737 |

RBD, receptor-binding domain; PBP, pooled BBIBP-CorV plasma.

| Pseudotyped virus | S gene source | Spike mutation site | Manufacturer |
|----------------------|-------------------|--|---------------------------------------|
| Wuhan-1 | | / | |
| D614G | | D614G | |
| Alpha | | delta69-70HV, delta144Y, N501Y, A570D, D614G, P681H, T716I, S982A, D118H | |
| Beta | | L18F, D80A, D215G, delta242- 244LAL, R246I, K417N, E484K, N501Y, D614G, A701V | |
| Gamma | | L18F, T20N, P26S, D138Y, R190S, K417T, E484K, N501Y, D614G, H655Y, T1027I, V1176F | Gobond Science |
| Kappa | GenBank: MN908947 | G142D, E154K, L452R, E484Q, D614G, P681R, Q1071H, H1101D | and Technology (Beijing) Co., Ltd. |
| Delta | | T19R, G142D, F157del, R158del, L452R, T478K, D614G, P681R, D950N | |
| Omicron | | A67V, H69del, V70del, T95I, G142D, V143del, Y144del, Y145del, N211del, L212I, ins214EPE, G339D, S371L, S373P, S375F, K417N, N440K, G446S, S477N, T478K, E484A, Q493R, G496S, Q498R, N501Y, Y505H, T547K, D614G, H655Y, N679K, P681H, N764K, D796Y, N856K, Q954H, N969K, L981F | |

Table S7. Basic information of pseudotyped SARS-CoV-2 viruses

Wuhan-1, SARS-CoV-2 Wuhan-Hu-1 strain.

Table S8. Gross severity grades recorded for the lung based on the percent lobe affected

by gross changes

| Pathological | Judgment basis | Evaluation | of |
|--------------|--|---------------|----|
| change score | | damage rating | |
| 0 | More than 75% of the lungs had diffuse alveolar injury; the | Severe | |
| | arveolar wan was significantly inickened, an arveolar hyanne | | |
| | hemorane was formed, and horosis was significant, most small | | |
| | of a solution of a solution of the solution of | | |
| | significant: inflammatory cell infiltration was diffuse | | |
| 5 | Here 51% 75% of the lung had diffuse alweolar injury: alweolar | | |
| 5 | wall thickening was obvious an alveolar hvaline membrane was | | |
| | formed and fibrosis was obvious: most small bronchial | | |
| | epithelium proliferated and dropped off considerably: edema | | |
| | and hyperplasia of vascular endothelial cells were obvious: | | |
| | inflammatory cell infiltration was diffuse | | |
| 4 | Here, 36%–50% of the lung had moderate injury; some alveolar | Moderate | |
| | walls were thickened, alveolar fibrin exudation and fibrosis were | | |
| | obvious; some bronchiolar epithelium hyperplasia and necrosis | | |
| | were present; edema and hyperplasia of some vascular | | |
| | endothelial cells; significant infiltration of inflammatory cells. | | |
| 3 | In this case, 21%–35% of the lung had mild to moderate injury; | | |
| | some alveolar walls were thickened; fibrin exudation and | | |
| | fibrosis were rare in alveoli; partial bronchiolar epithelial | | |
| | hyperplasia was observed; significant infiltration of | | |
| | inflammatory cells was present | 2 611 1 | |
| 2 | Here, 5%–20% of the lung had mild pathological injury; most | Mild | |
| | alveolar walls and alveolar cavities had normal structures; a | | |
| | small part of the alveolar walls was thickened; fibrin exudation | | |
| | and horosis were seen in alveon; small amount of bronchiolar | | |
| | monocyte and lymphocyte infiltration was observed | | |
| 1 | Finally $<5\%$ of the lung had slight nathological injury: most | | |
| 1 | alveoli bronchi and blood vessels had normal structures; very | | |
| | few alveolar walls were thickened: fibrin exudation and fibrosis | | |
| | were rare in alveoli; individual bronchiolar epithelial | | |
| | hyperplasia was observed; a small amount of mononuclear and | | |
| | lymphocyte infiltration was occasionally was observed. | | |
| 0 | The alveolar structure was intact; there was no inflammatory | None | |
| | infiltration and no congestion; the bronchial and vascular | | |
| | structures were normal. | | |