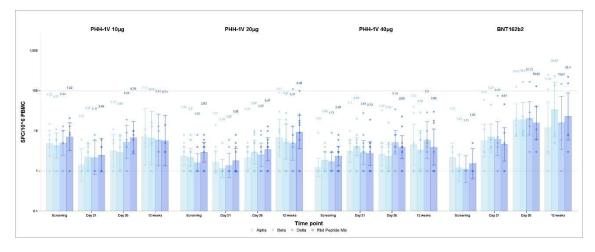
#### SUPPLEMENTARY SECTION

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# Supplementary figure 1

Total SARS-CoV-2 specific T-cell responses measured by IFN- $\gamma$  ELISPOT against variants of concern (VOC) alpha (B.1.1.7), beta (B.1.351), delta (B.1.617.2) and RBD peptide mix variants, at screening, 21 and 35 days after first vaccination and 12 weeks after second vaccination are shown for all participants distributed by vaccine group PHH-1V 10 $\mu$ g n= 5, PHH-1V 20 $\mu$ g n=10, Cohort 3 PHH-1V 40 $\mu$ g n=10, control n=5. T-cell responses are expressed as mean spot forming cells / 10<sup>6</sup> PBMC. Bars indicate 95% confidence intervals and data points represent individual results.



# **Supplementary Table 1**

# Humoral and cellular immunogenicity data censored obtained at 24W and 48W visits.

Censored data on total binding antibodies titres measured by ELISA at 24W and 48W time-points are shown by vaccine group: 24W PHH-1V  $10\mu g$  n=2, PHH-1V  $20\mu g$  n=6, Cohort 3 PHH-1V  $40\mu g$  n=4, control n=4; 48W PHH-1V  $10\mu g$  n=1, PHH-1V  $20\mu g$  n=3, Cohort 3 PHH-1V  $40\mu g$  n=3, control n=3. Total binding antibodies titers are expressed in GMT shown as adjusted treatment mean (95% CI). GMT geometric mean titer; CI confidence interval.

Neutralization antibodies titres measured by pseudovirus-based neutralization assay (PBNA) against variants of concern assessed alpha (B.1.1.7), beta (B.1.351), delta (B.1.617.2) and gamma (P.1) at 24W and 48W time-points are shown by vaccine group: 24W PHH-1V 10 $\mu$ g n=2, PHH-1V 20 $\mu$ g n=6, Cohort 3 PHH-1V 40 $\mu$ g n=4, control n=4; 48W PHH-1V 10 $\mu$ g n=1, PHH-1V 20 $\mu$ g n=3, Cohort 3 PHH-1V 40 $\mu$ g n=3, control n=3. Neutralization titers measured as IC50 are expressed in GMT shown as adjusted treatment mean (95% CI).

T-cell responses measured by IFN- $\gamma$  ELISPOT against variants of concern assessed alpha (B.1.17), beta (B.1.351), delta (B.1.617.2) and RBD peptide mix at 24W time-point are shown by vaccine group: PHH-1V 10 $\mu$ g n=2, PHH-1V 20 $\mu$ g n=6, Cohort 3 PHH-1V 40 $\mu$ g n=4, control n=4. T cell response is expressed as mean spot forming cells / 10<sup>6</sup> PBMC (95% CI).

	PHH-1V 10µg	PHH-1V 20μg	PHH-1V 40µg	BNT162b2
Binding Antibodies				
24 Week	(n=2) 981.01 (101.55 - 9476.81) (n=1) 927.55	(n=6) 392.21 (84.51 - 1820.2) (n=3) 1717.39	(n=3) 4370.95 (804.39 - 23751.11) (n=3) 2440.96	(n=4) 1837.21 (232.97 - 14488.49) (n=3) 1955.56
40 Week	(87.15 - 9872.42)	(401.39 – 7348)	(407.61 - 14617.57)	(322.83 - 11845.87)
Neutralizing Antibodies				
ALPHA B.1.1.7				
24 Week	(n=2) 167.41 (13.43 - 2086.51)	(n=6) 104.52 (20.71 - 527.52)	(n=3) 2979.36 (419.78 – 21145.74)	(n=4) 375.29 (46.18 – 3049.67)
48 Week	(n=1) 586.37 (73.7 - 4665.25)	(n=3) 669.75 (196.92 - 2277.88)	(n=3) 1385.86 (268.87 - 7143.14)	(n=3) 429.62 (95.26 - 1937.49)
BETA B.1.351 (n=2) (n=6) (n=3) (n=4)				
24 Week	250.84 (17.79 -3537.15) (n=1)	139.97 (25.14 - 779.29) (n=3)	2902.13 (376.43 - 22374.29) (n=3)	559.57 (59.92 - 5225.64) (n=3)
48 Week	250.91 (34.68 - 1815.5)	856.35 (261.77 - 2801.41)	À991.44 (1027.14 - 24256.2)	801.07 (181.61 – 3533.44)
DELTA B.1.617.1	(*)	(==:,	(**=****	()
24 Week	(n=2) 294.19 (27.53 - 3144.12)	(n=6) 64.74 (15.61 - 268.45) (n=3)	(n=3) 10143.23 (1518.42 - 67758.2) (n=3)	(n=4) 413.47 (68.9 - 2481.06) (n=3)
48 Week	(n=1) 215.25 (47.74 - 970.62)	(11–3) 110.8 (45.29 - 271.05)	978.46 (305.79 - 3130.85)	356.31 (118.7 - 1069.57)
GAMMA P.1				
24 Week	(n=2) 348.09 (26.6 - 4555.03)	(n=6) 125.3 (26.08 - 602)	(n=3) 4562.92 (589.43 - 35322.89)	(n=4) 480.87 (65.63 - 3523.35)
48 Week	(n=1) 644.05 (72.46 - 5724.85)	(n=3) 362.42 (100.78 - 1303.26)	(n=3) 3058.97 (524.22 - 17849.91)	(n=3) 560.44 (114.77 - 2736.8)
T-cell responses				
ALPHA B.1.1.7	(n=2)	(n=6)	(n=3)	(n=4)
24 Week	5.32 (0.24 - 117.38)	2.11 (0.25 - 17.49)	77.41 (7.16 - 837.38)	79.49 (4.11 - 1536.36)
BETA B.1.351	(n=2)	(n=6)	(n-2)	(n=4)
24 Week	(n=2) 0.63 (0.09 - 4.27)	(n=6) 7.17 (2.07 - 24.81)	(n=3) 27.55 (5.72 - 132.8)	(n=4) 100.87 (16.51 - 616.32)
DELTA B.1.617.1	(n=2)	(n=6)	(n=3)	(n=4)
24 Week	15.48 (4.01 - 59.73)	5.54 (2.39 - 12.8)	17.05 (5.24 - 55.52)	41.29 (12.38 - 137.74)
RBD PEPTIDE MIX				
04.14/	(n=2)	(n=6)	(n=3)	(n=4)
24 Week	8.85 (1.17 - 67.23)	5.62 (1.67 - 18.93)	24.91 (3.54 - 175.04)	34.44 (7.14 - 166.1)

### Supplementary methods 1

#### Biosafety Approval.

The biologic biosafety committee of the Research Institute Germans Trias i Pujol approved the execution of SARS-CoV-2 experiments at the BSL3 laboratory of the Center for Bioimaging and Comparative Medicine (CSB-20-015-M3).

#### Cell culture and Viral isolation and titration.

Vero E6 cells (ATCC CRL-1586) were cultured in Dulbecco's modified Eagle medium (Invitrogen) supplemented with 10% fetal bovine serum (FBS; Invitrogen), 100 U/ml penicillin, 100 μg/ml streptomycin, (all from Invitrogen).

SARS-CoV-2 was isolated from a nasopharyngeal swab collected in January 2021 in Spain in Vero E6 cells as described in (1). The virus stock was prepared collecting the supernatant from Vero E6 and sequenced as detailed in (1). Genomic sequence was deposited at GISAID repository (http:// gisaid.org) with accession number EPI\_ISL\_1663567. Viral stocks were propagated in Vero E6 cells for two passages and titrated in 10-fold serial dilutions to calculate the TCID50 per mL. Infection was set to achieve a 50% of viral induced cytopathic effect measured with Cell Titer GloTM Luciferase reagent, as described in the next section.

#### Neutralization Assay.

Neutralization assays were performed in duplicate. Briefly, 60 TCID<sub>50</sub> of SARS-CoV-2 were preincubated with serial dilutions of heat-inactivated plasma samples from the indicated individuals for 1 h at 37 °C. Preincubated viruses were added to 60.000 Vero E6 cells per well in 96 well plates. 72 h later, viral-induced cytopathic effect was measured using the Cell Titer Glo Luciferase reagent (Promega) and the Luminoskan Plate Reader from Thermofisher. The relative light units (RLU) were normalized to untreated non-infected cells (without plasma or virus), and the ID<sub>50</sub> (the reciprocal dilution inhibiting 50% of the cytopathic effect) was calculated by plotting and fitting the log of plasma dilution vs. response to a 4-parameter equation in GraphPad Prism 9.3.1, as previously described in (2, 3, 4).

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- 2. Trinité B, Tarrés-Freixas F, Rodon J, Pradenas E, Urrea V, Marfil S, Rodríguez de la Concepción ML, Ávila-Nieto C, Aguilar-Gurrieri C, Barajas A, et al. SARS-CoV-2 infection elicits a rapid neutralizing antibody response that correlates with disease severity. *Sci Rep* (2021) 11:2608. doi:10.1038/s41598-021-81862-9
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# **Supplementary Figure 2**

Example of gating strategy used for the analysis of intracellular cytokine stainings (ICS) from peripheral blood mononuclear cells (PBMCs). First, total cells were selected excluding cellular debris (SSC-A vs FSC-H). Then single cells were separated from doublets (FSC-H vs FSC-A). Next, CD3 positive cells were selected excluding autofluorescent and CD3 negative cells (Empty vs CD3-PerCp). After, CD3 postive living cells were determined (live/dead vs CD3-PerCp) and then CD4 T-cells and CD8 T-cells were selected (CD4-BV421 vs CD8-BV510). Finally, CD4 and CD8 T-cells positive for IL-2, IFNγ and IL-4 were determined (IFNγ-APC vs IL-2-PE) and (IFNγ-APC vs IL-4-PE-Cy7).

