

**Supplemental information**

**SARS-CoV-2 prolonged infection during advanced**

**HIV disease evolves extensive immune escape**

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Supplementary materials for SARS-CoV-2 prolonged infection during advanced HIV disease evolves extensive immune escape

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# 1 Table S1: Characteristics of SARS-CoV-2 convalescent or vaccinated study participants related to Figures 1 and 2

Participant number	Infected or BNT162b2 vaccinated	Variant/strain by infection date	Variant/strain by sequencing	Sequence ID	GISAID Accession	Age range (y)	Sex	Symptom onset or vaccination to plasma collection (days)
1	Infected	Ancestral	B.1.1.273	K008646	EPI_ISL_2397308	30-39	F	6, 20, 26, 34, 71, 106, 190
2	Infected	Ancestral	B.1.1	K003667	EPI_ISL_602623	50-59	M	29
3	Infected	Ancestral	B.1.1.273	K003675	EPI_ISL_602631	40-49	F	32
4	Infected	Ancestral	B.1.1.117	K003668	EPI_ISL_602624	70+	F	29
5	Infected	Ancestral	B.1.1	K004289	EPI_ISL_660170	60-69	F	27
6	Infected	Ancestral	B.1.140	K004295	EPI_ISL_660176	60-69	F	28
7	Infected	Ancestral	B.1.1.84	K003673	EPI_ISL_602629	40-49	F	28
8	Infected	Ancestral	B.1	K004291	EPI_ISL_660172	30-39	F	30
9	Infected	Ancestral	C.9	K004302	EPI_ISL_660181	60-69	M	26
10	Infected	Beta	Beta	K008635	N/A*	40-49	F	30
11	Infected	Beta	Beta	K008636	N/A*	40-49	F	41
12	Infected	Beta	Beta	K008628	N/A*	40-49	M	32
13	Infected	Beta	Beta	K008637	EPI_ISL_1229368	50-59	M	42
14	Infected	Beta	N/A <sup>\$</sup>	N/A <sup>\$</sup>	N/A	30-39	F	32
15	Infected	Beta	Beta	K010372	N/A*	30-39	M	33
16	Infected	Beta	Beta	K010356	N/A*	70+	F	48
17	Infected	Beta	Beta	K008633	EPI_ISL_1229367	60-69	F	29
18	Infected	Beta	Beta	K010370	N/A*	30-39	F	31
19	Infected	Delta	Delta	K021407	EPI_ISL_3722338	40-49	F	26
20	Infected	Delta	Delta	K021399	EPI_ISL_3722335	40-49	M	23 <sup>#</sup>
21	Infected	Delta	Delta	K021401	N/A*	50-59	M	31
22	Infected	Delta	Delta	K021225	N/A*	40-49	M	13 <sup>#</sup>
23	Infected	Delta	Delta	K021400	N/A*	40-49	M	44
24	Infected	Delta	Delta	K021226	N/A*	50-59	M	44
25	Infected	Delta	Delta	K020186	EPI_ISL_3939068	40-49	M	31
26	Infected	Delta	Delta	K020187	EPI_ISL_3939088	30-39	M	31
27	Infected	Delta	Delta	K020214	EPI_ISL_3447779	50-59	F	30 <sup>#</sup>
28	Infected	Delta	Delta	K021404	N/A*	30-39	M	32
29	BNT162b2	-	-	-	-	60-69	M	11
30	BNT162b2	-	-	-	-	60-69	M	10
31	BNT162b2	-	-	-	-	50-59	M	18
32	BNT162b2	-	-	-	-	40-49	F	9
33	BNT162b2	-	-	-	-	60-69	M	10
34	BNT162b2	-	-	-	-	20-29	F	8
35	BNT162b2	-	-	-	-	60-69	M	128
36	BNT162b2	-	-	-	-	30-39	F	134
37	BNT162b2	-	-	-	-	20-29	M	131
38	BNT162b2	-	-	-	-	50-59	M	152
39	BNT162b2	-	-	-	-	30-39	F	153
40	BNT162b2	-	-	-	-	30-39	M	158

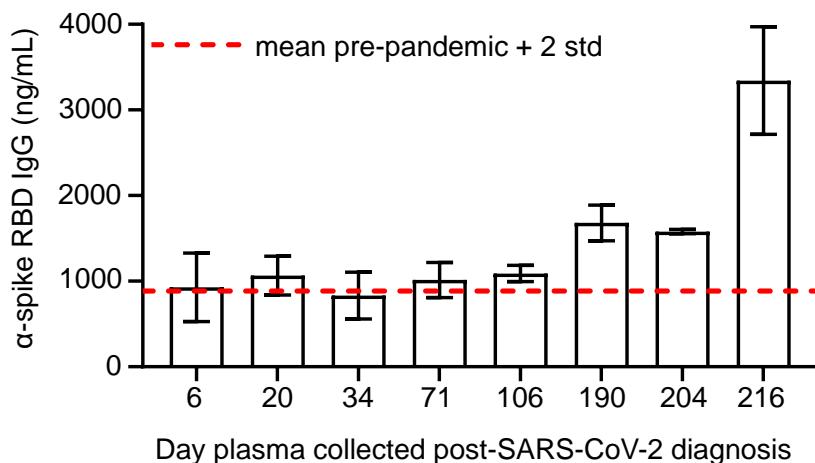
N/A – Not available. \* <90% coverage. Not submitted to GISAID but sufficient sequence for variant call. <sup>\$</sup>Not sequenced. <sup>#</sup>Asymptomatic at diagnosis, date post-diagnostic test used instead of symptom onset date.

**2 Table S2: Read counts of majority and minority genotypes detected in swab and outgrown virus related to Figure 2**

	AA change	Nucleotide change	Codon change	Day 0	Day 6	Day 20	Day 34	Day 71	Day 106	Day 190
				Swab	Swab	Outgrowth	Swab	Outgrowth	Swab	Outgrowth
T95I	21846C>T	21845	ACT>ATT	ACT -143	ACT -605	ACT -1342	ACT -2667	ACT -1540	ACT -935	ACT -96
				ATT -0	ATT -9	ATT -10	ATT -10	ATT -298	ATT -33	ATT -1626
G142V	21987G>T	21986	GGT -10	GGT -1386	GGT -316	GGT -880	GGT -1352	GGT -954	GGT -1238	GGT -480
				GTT -40	GTT -622	GTT -808	GTT -313	GTT -4	GTT -14	GTT -5
R190K	22131G>A	22130	AGG>AAG	AGG -46	AGG -591	AGG -119	AGG -1702	AGG -357	AGG -1685	AGG -324
				AAG -0	AAG -8	AAG -0	AAG -1	AAG -3	AAG -2	AAG -22
*L244del	22293	22811	ACA>del	ACA -12	ACA -1378	ACA -371	ACA -50	ACA -0	ACA -17	ACA -17
				ACA>del	del -1	del -0	del -4	del -158	del -176	del -4
K417I	22812A>G	22811	AAG>ACG	AGC -117	AAG -3237	AAG -141	AAG -3209	AAG -122	AAG -297	AAG -96
				ACG -0	ACG -1	ACG -0	ACG -0	ACG -0	ACG -1	ACG -0
D422Y	22841G>T	22841	GAT>TAT	GAT -107	GAT -1281	GAT -1273	GAT -3434	GAT -1164	GAT -3434	GAT -1079
				TAT -0	TAT -3	TAT -7	TAT -0	TAT -1	TAT -0	TAT -42
I455F	22927G>T	22925	TTC>TTT	TTC -11	TTC -314	TTC -886	TTC -3219	TTC -978	TTC -4331	TTC -888
				TTT -0	TTT -9	TTT -0	TTT -0	TTT -2	TTT -2	TTT -3
F456L	22928T>C	22928	TTT>CTT	TTT -12	TTT -3739	TTT -1112	TTT -3248	TTT -1109	TTT -4361	TTT -1011
				CTT -0	CTT -6	CTT -31	CTT -0	CTT -19	CTT -20	CTT -3
A475V	22986C>T	22985	GCC>GTC	GCC -17	GCC -3409	GCC -1180	GCC -3629	GCC -11	GCC -2874	GCC -43
				GTC -0	GTC -11	GTC -30	GTC -1218	GTC -1369	GTC -2421	GTC -1224
E484K	23012G>A	23012	GAA>AAA	GAA -17	GAA -56	GAA -363	GAA -1632	GAA -27	GAA -2395	GAA -55
				AAA -0	AAA -2063	AAA -917	AAA -2881	AAA -1276	AAA -2139	AAA -1133
F490S	23031T>C	23030	TTT>TCT	TTT -18	TTT -1605	TTT -1196	TTT -4067	TTT -1226	TTT -4007	TTT -1031
				TCT -0	TCT -2	TCT -15	TCT -0	TCT -10	TCT -30	TCT -2912
M501Y	23063A>T	23063	AAT>TAT	AAT -13	AAT -810	AAT -1428	AAT -2034	AAT -1498	AAT -2855	AAT -1354
				TAT -0	TAT -2	TAT -2	TAT -2	TAT -1	TAT -29	TAT -0
D614G	23403A>G	23402	GAT>GGT	GAT -3	GAT -8	GAT -4	GAT -0	GAT -6	GAT -3	GAT -14
				GGT -158	GGT -6668	GGT -811	GGT -5299	GGT -904	GGT -2863	GGT -886
R682W	23606C>T	23606	CGG>TGG	CGG -98	CGG -1831	CGG -189	CGG -27	CGG -618	CGG -59	CGG -607
				TGG -1	TGG -4	TGG -407	TGG -0	TGG -5	TGG -0	CGG -124
D796Y	23948G>T	23948	GAT>TAT	GAT -16	GAT -4293	GAT -1155	GAT -1235	GAT -1409	GAT -568	GAT -1313
				TAT -70	TAT -70	TAT -70	TAT -477	TAT -5	TAT -477	TAT -569
A1078V	24795C>T	24794	GCT>GTT	GCT -219	GCT -772	GCT -772	GCT -4397	GCT -3119	GCT -2519	GCT -1270
				GTT -3	GTT -4833	GTT -358	GTT -1	GTT -18	GTT -12	GCT -1418

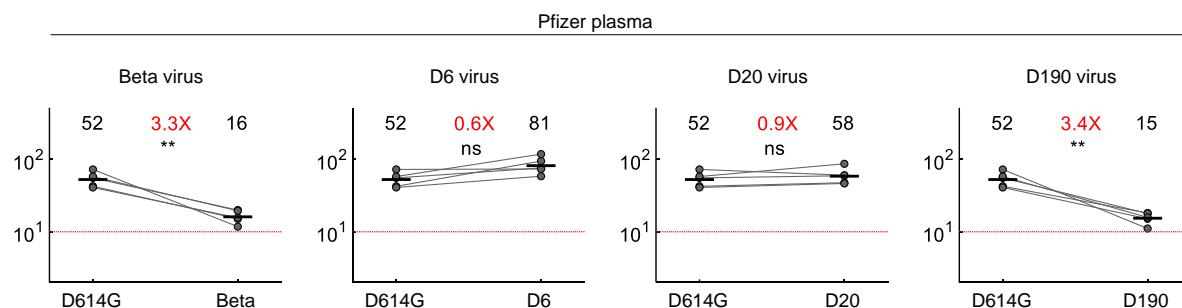
\*Only deletions where the adjacent codon was preserved were counted.

### 3 Figure S1: Spike specific antibody levels with time post-SARS-CoV-2 diagnosis related to Figure 2.



**Fig S 1: Spike specific antibody levels with time post-SARS-CoV-2 diagnosis.** Shown are mean ( $n=4$  replicates per timepoint) and standard deviation of anti-spike RBD antibody concentrations measured in the plasma of the participant with advanced HIV disease by ELISA. Red dashed line denotes the mean + 2 standard deviations of signal from a set of 6 control samples, including plasma of pre-pandemic controls ( $n=4$ ) and pre-pandemic commercial human serum ( $n=2$ ).

### 4 Figure S2: Neutralization of Beta variant and evolved virus by a subset of BNT162b2 plasma donors related to Figure 2



**Fig S 2: Neutralization of Beta variant and evolved virus by a subset of BNT162b2 plasma donors.** Neutralization of Beta, D6, D20, and D190 compared to D614G by Pfizer BNT162b2 plasma ( $n=5$ ). Plasma donors were 136072, 136074, 136075, 136076, 136078. Red horizontal line denotes most concentrated plasma tested. Numbers in black are GMT FRNT50. Numbers in red are fold-change in GMT between virus strain on left and right. \*\*  $<0.01-0.001$ , as determined by the Wilcoxon rank sum test.