

Supplementary Information

Safety and antiviral activity of combination HIV-1 broadly neutralizing antibodies in viremic individuals

Supplementary Figures

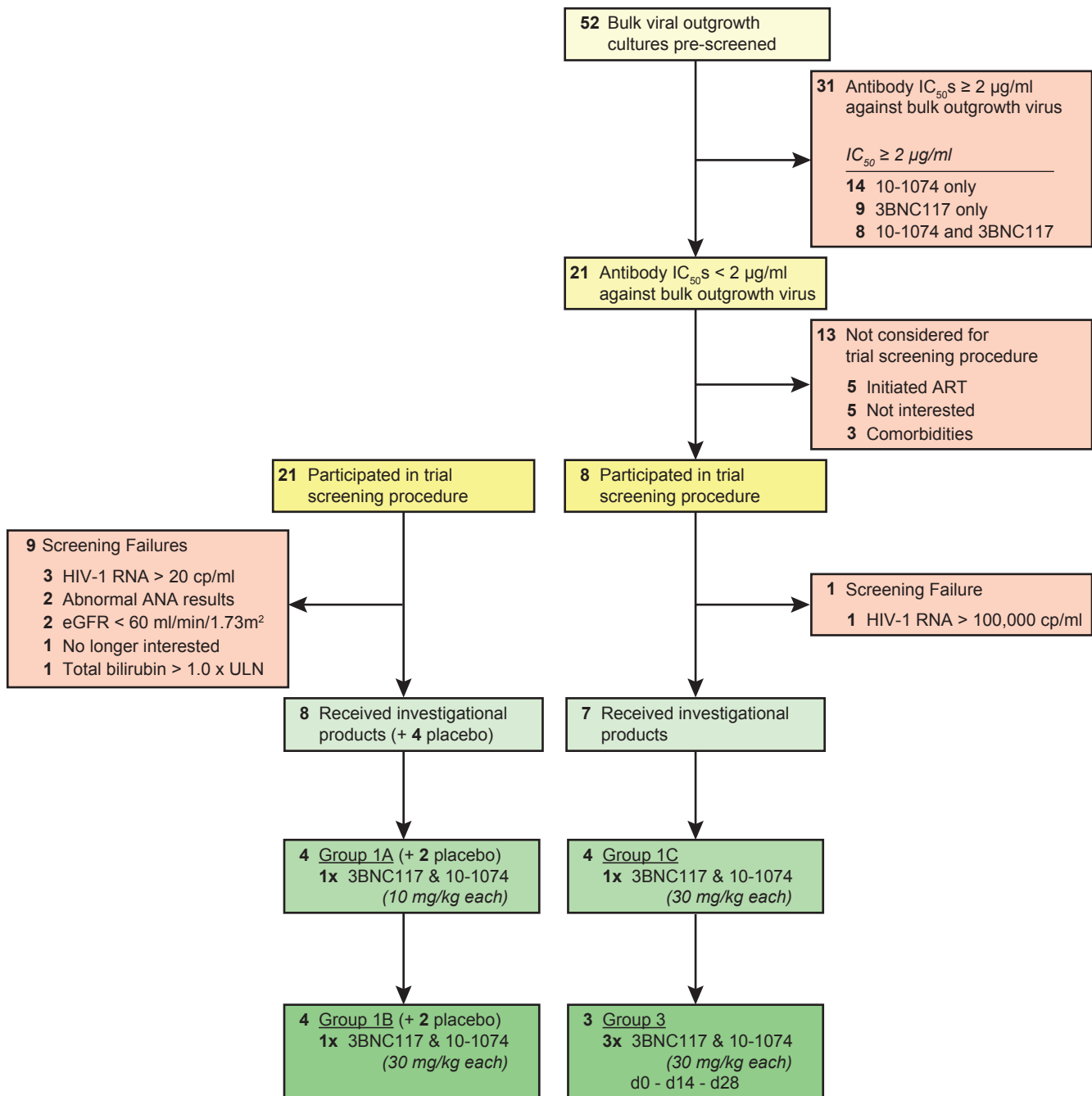
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Individuals on ART

Viremic individuals

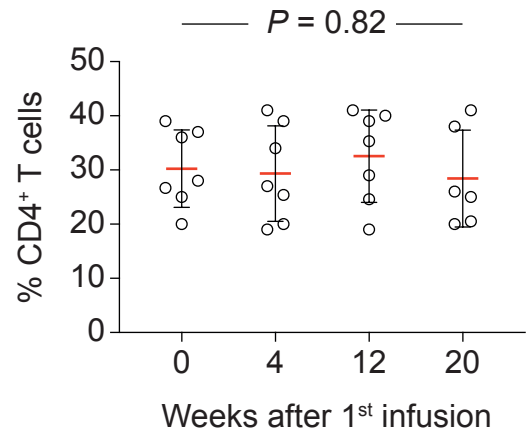
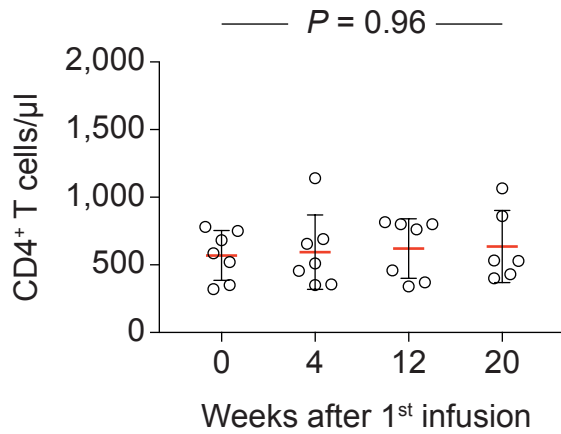


Supplementary Figure 1. Subject enrollment and study design

HIV-1-infected individuals on antiretroviral therapy were screened for enrollment into group 1A (single infusion of 3BNC117 + 10-1074 (10 mg/kg each)) and group 1B (single infusion of 3BNC117 + 10-1074 (30 mg/kg each)). Enrollment into groups 1A and 1B was double-blind and individuals were randomized to receive the antibodies or placebo at a 2:1 ratio (total $n=6$ per group). Placebo recipients are not included in the data analysis. Viremic individuals off antiretroviral therapy were invited for participation if virus obtained by bulk CD4⁺ T cell outgrowth culture was determined to be sensitive against both 3BNC117 and 10-1074 ($IC_{50} < 2 \mu\text{g/ml}$ for each antibody in TZM-bl assay). Enrolled viremic individuals received a single infusion of 30 mg/kg of each antibody (group 1C) or three infusions of 30 mg/kg of each antibody every two weeks (d0, d14, d28).

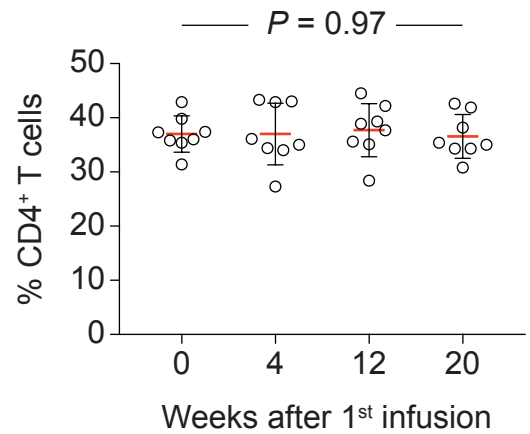
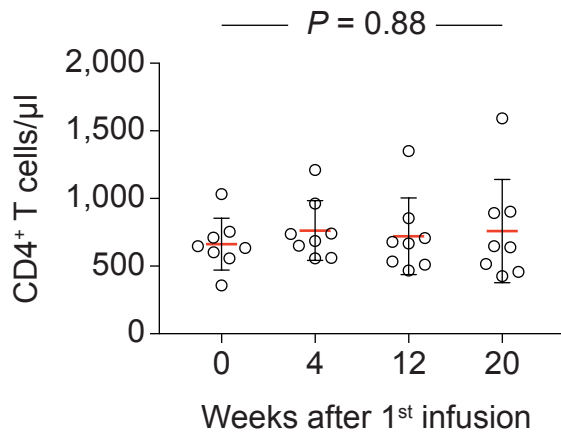
a

Viremic



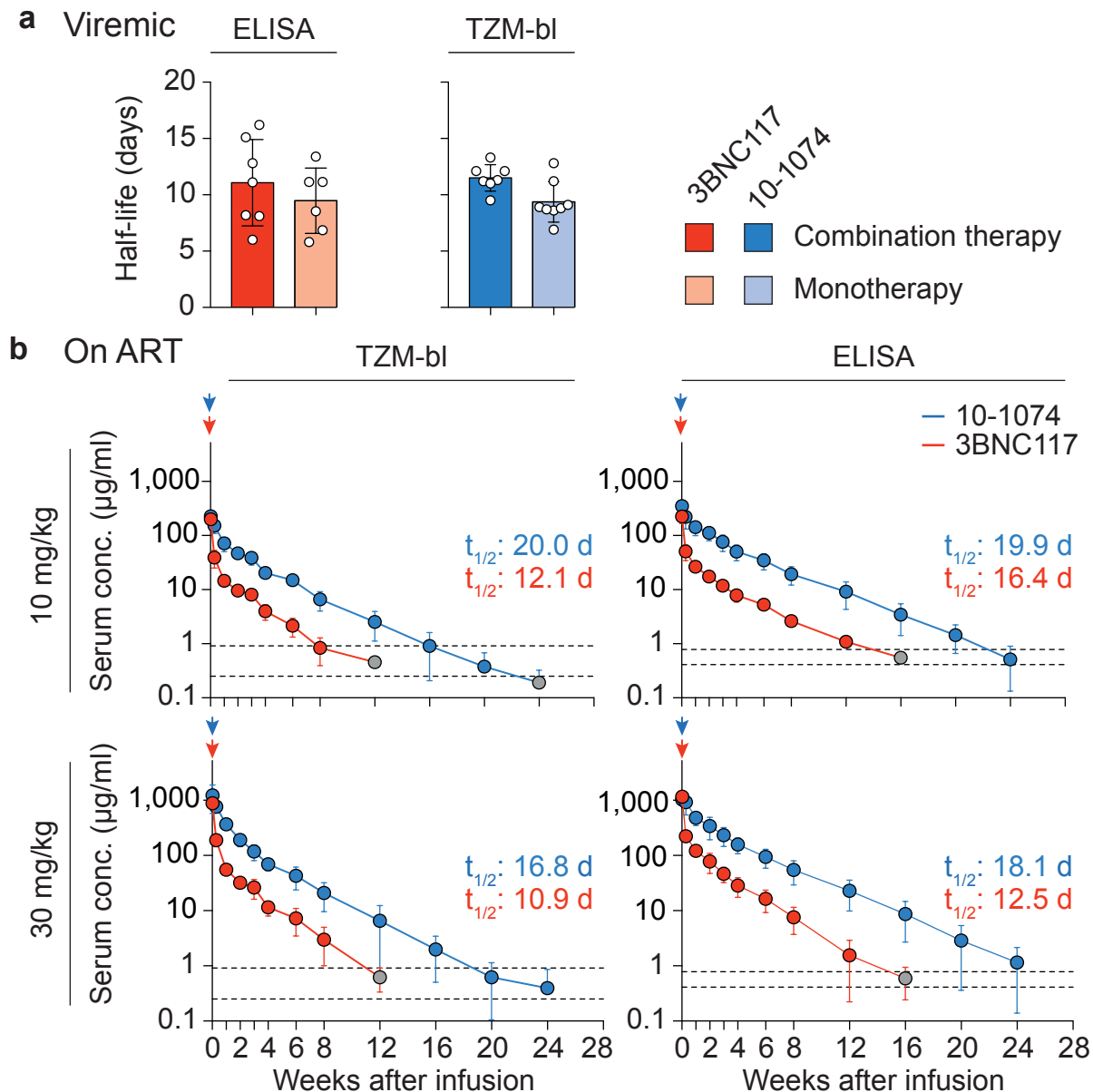
b

On ART



Supplementary Figure 2. CD4⁺ T cell counts

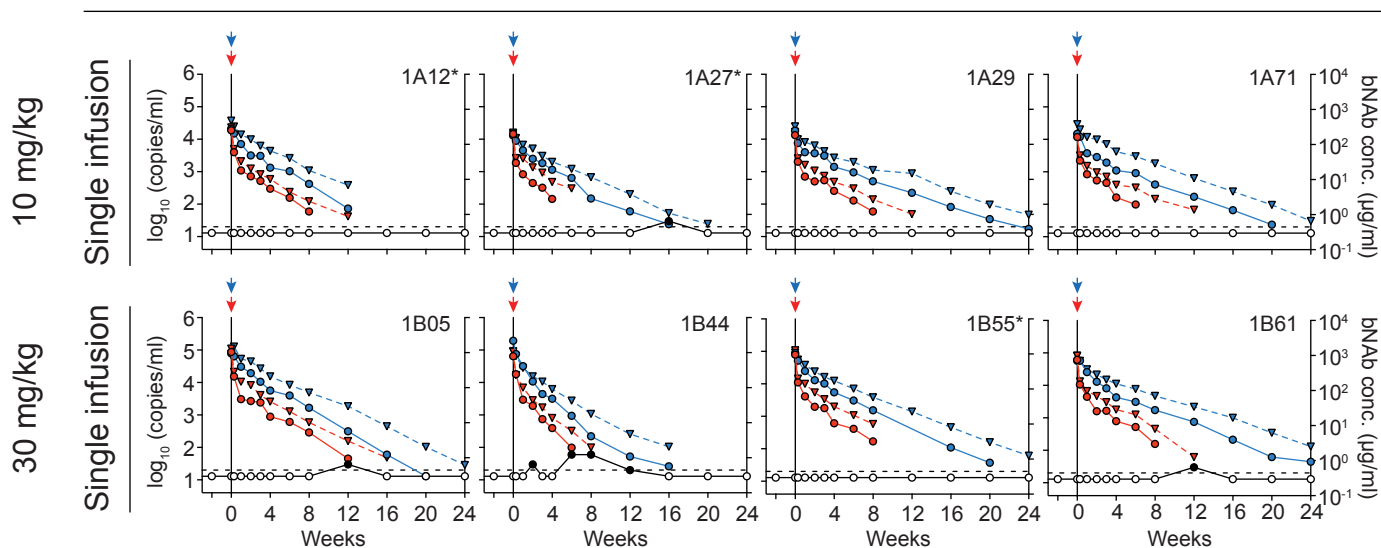
Absolute CD4⁺ T cell counts (left) and relative CD4⁺ T cell frequency among CD3⁺ lymphocytes (right) in (a) viremic individuals ($n=7$) and (b) individuals on antiretroviral therapy ($n=8$). Red lines indicate arithmetic mean and error bars indicate standard deviation. P values were calculated using one-way ANOVA comparing all groups.



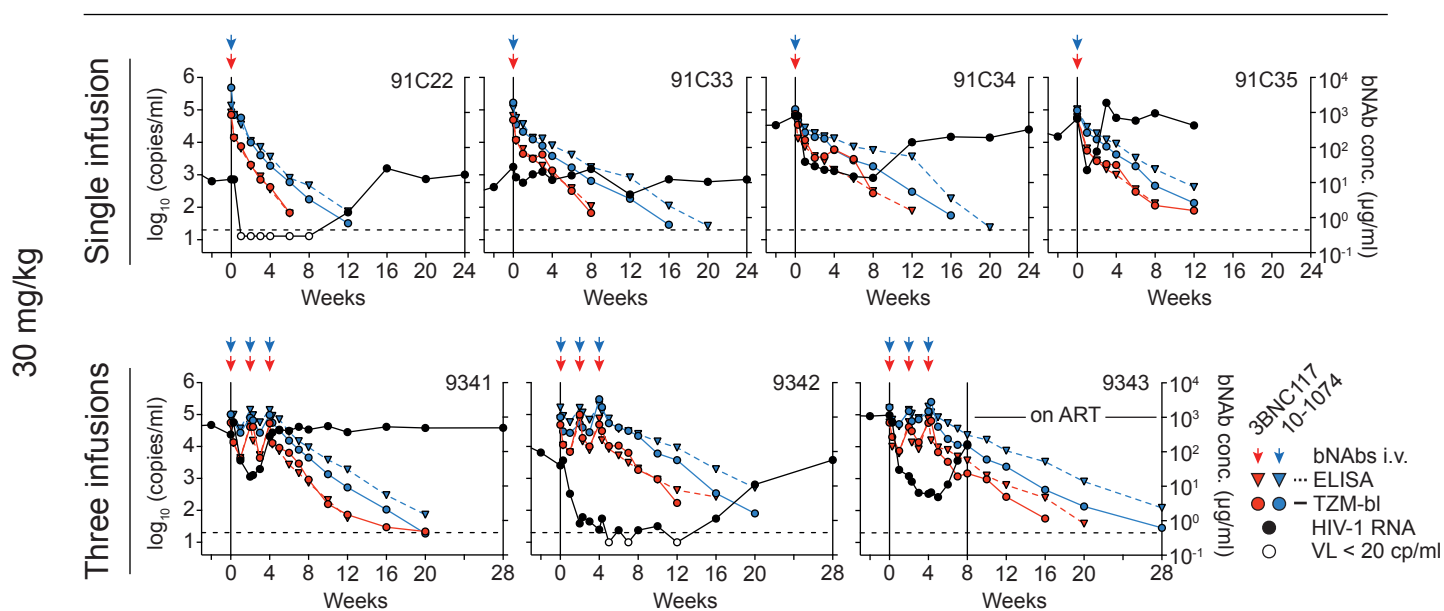
Supplementary Figure 3. BNAbs half-lives in HIV-1-infected individuals

(a) Terminal bNAb half-lives based on serum concentrations determined by ELISA or TZM-bl assay in viremic study participants receiving a single or three infusions of 30 mg/kg of each 3BNC117 and 10-1074 (combination) ($n = 7$), and viremic individuals receiving a single infusion of 3BNC117 or 10-1074 alone (monotherapy) at a dose of 30 mg/kg ($n = 6$ for 3BNC117 and $n = 8$ for 10-1074) (3BNC117, Caskey and Klein *et al.*, *Nature* **522**, 487-491 (2015); 10-1074, Caskey, Schoofs and Gruell *et al.*, *Nat. Med.* **23**, 185-191 (2017)). Columns indicate arithmetic means and bars indicate standard deviation. Circles indicate individual participants. Note that different pseudovirus strains were used to determine antibody concentrations (3103.v3.C10 and X2088_c9 for mono- and combination therapy, respectively). (b) Serum concentrations of 3BNC117 (red) and 10-1074 (blue) after a single infusion of 10 mg/kg (top, $n = 4$) or 30 mg/kg (bottom, $n = 4$) of each antibody to individuals on continuous ART as determined by TZM-bl (left) or ELISA (right). For the TZM-bl assay, time points with detectable activity against MuLV ($\text{ID}_{80} > 20$) were excluded. In cases of detectable baseline levels by ELISA, these were subtracted, and only measurements above 3-fold baseline levels are included. Circles and lines indicate arithmetic mean and standard deviation, respectively. Dashed lines indicate detection limits for 3BNC117 (top) and 10-1074 (bottom). Grey circles indicate levels below the limit of detection. Arrows indicate antibody infusions.

a On ART

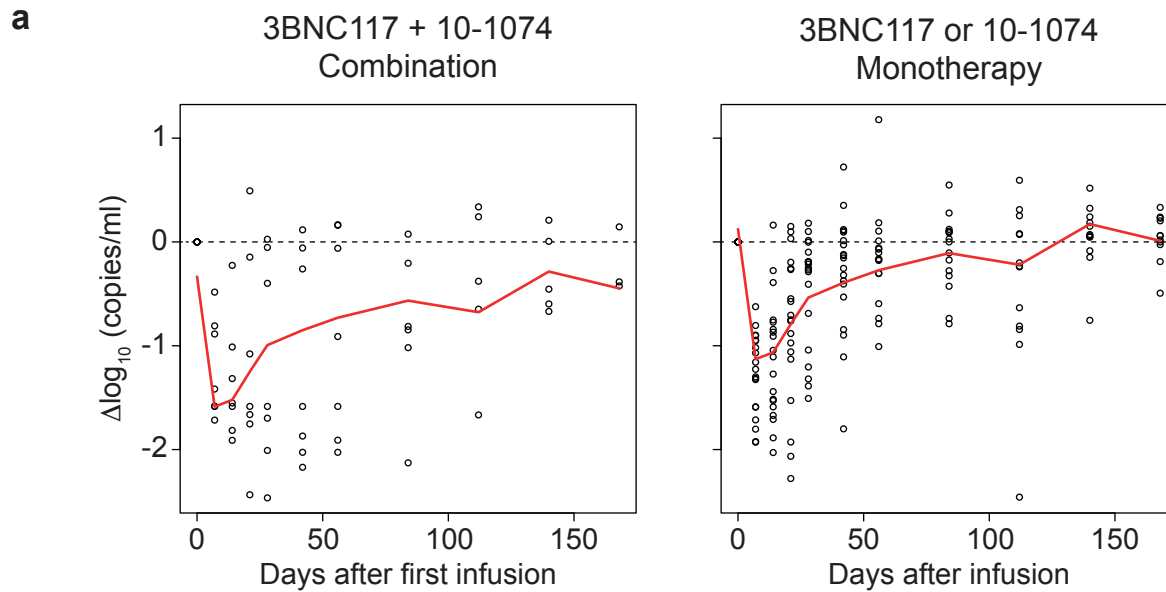


b Viremic



Supplementary Figure 4. HIV-1 RNA and bNAb antibody levels in all study participants

Graphs illustrate viral load (black line; left y-axis) and bNAb serum levels (red, 3BNC117; blue, 10-1074; right y-axis) over time in (a) individuals on antiretroviral therapy (top, single infusion of 10 mg/kg of each antibody; bottom, single infusion of 30 mg/kg of each antibody) and (b) viremic participants (top, single infusion of 30 mg/kg of each antibody; bottom, three infusions of 30 mg/kg of each antibody every two weeks). Only bNAb levels above the respective limit of detection are shown. Antibody levels determined by TZM-bl assay are indicated by colored solid line and circles, and TZM-bl results were excluded if activity against negative control virus (MuLV) was observed. Antibody levels determined by ELISA are indicated by colored dashed line and triangles. Asterisks (*) indicate individuals with detectable baseline levels in ELISA. In these cases, baseline levels were subtracted from subsequent bNAb concentrations determined by ELISA (3BNC117, 1A27 (1.17 µg/ml) and 1B55 (2.48 µg/ml); 10-1074, 1A12 (0.43 µg/ml)) and curves only show concentrations that are 3-fold higher than baseline levels. Dashed black line indicates limit of detection for HIV-1 RNA (20 copies/ml). Arrows indicate antibody infusions.



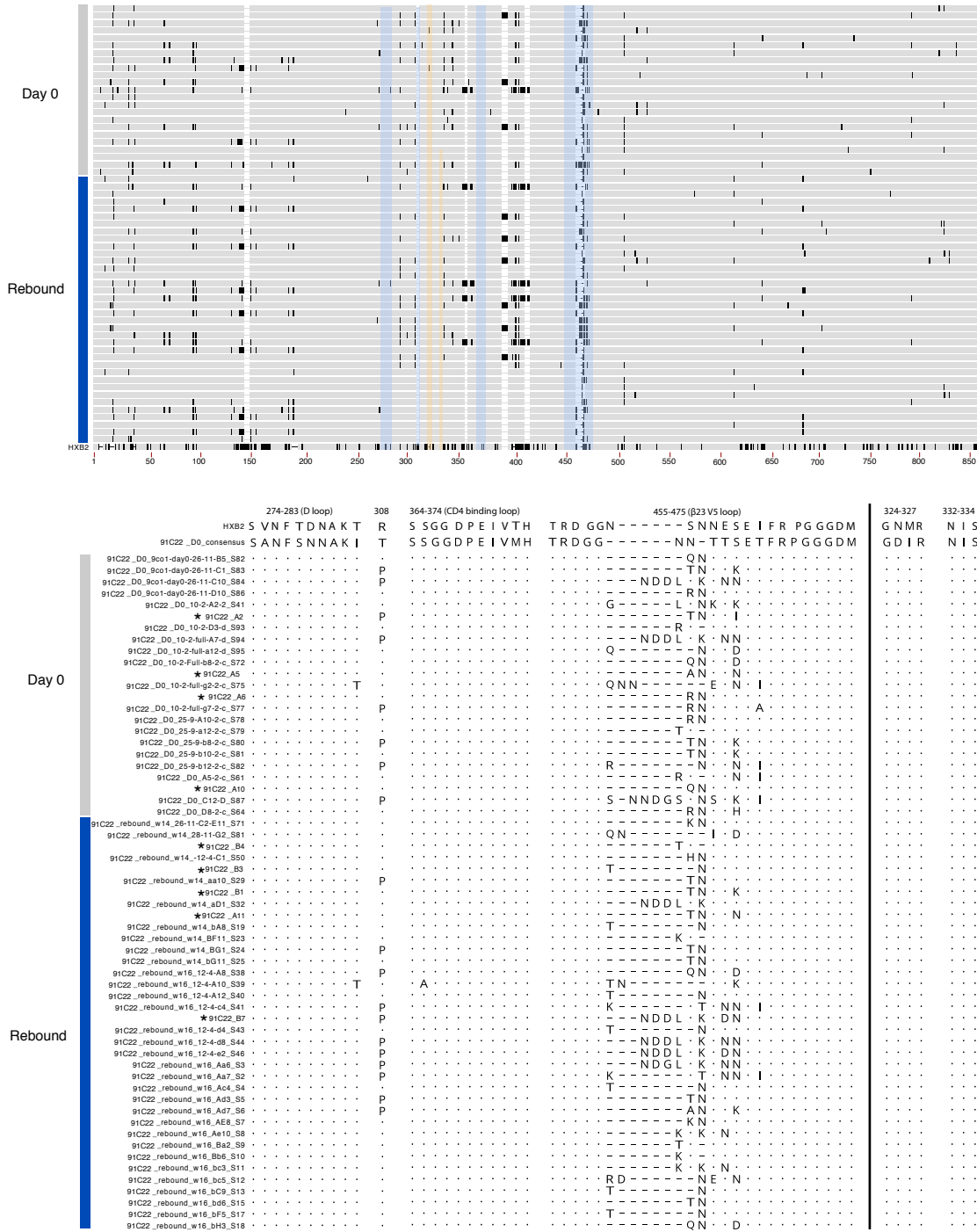
b

Correlation	With Treatment	Without Treatment
none	410.1444777	413.2992212
corARMA(p=0,q=1)	342.3434909	345.3257615
corARMA(p=0,q=2)	334.8421682	337.8340493
corARMA(p=1,q=0)	326.0505608	327.9541902
corARMA(p=1,q=1)	327.2169635	329.4295783
corARMA(p=1,q=2)	329.2051113	331.4291768
corARMA(p=2,q=0)	327.1790532	329.4238417
corARMA(p=2,q=1)	328.5336267	330.5019875
corARMA(p=2,q=2)	330.4939601	332.5006249
corExp	326.0505609	327.9541903
corLin	335.9980408	342.6470672
corRatio	328.314909	331.1412435
corSpher	326.8269772	329.4149949
corGaus	337.8828039	340.8426585
corSymm	296.1530987	308.2251827

Supplementary Figure 5. Viral suppression in combination therapy vs. monotherapy

(a) Marginal means (least-squares means) for the fit linear mixed effects model of individuals receiving 3BNC117 + 10-1074 combination therapy (single infusion or three infusions every two weeks, 30 mg/kg each per infusion) (left) or 3BNC117 or 10-1074 monotherapy (single infusion, 30 mg/kg). Red line indicates marginal means and circles indicate viral loads. Monotherapy data are derived from previously published results in antibody-sensitive viremic individuals (3BNC117, Caskey and Klein *et al.*, *Nature* **522**, 487-491 (2015); 10-1074, Caskey, Schoofs and Gruell *et al.*, *Nat. Med.* **23**, 185-191 (2017)). Only measurements off ART are included (combination, $n=7$ up to d56, $n=6$ on d84, $n=5$ for later time points; monotherapy, $n=19$ up to d42, $n=16$ on d56, $n=15$ on d84, $n=13$ on d102 and d140, $n=12$ on d168). (b) Selection of linear mixed effects model to compare 3BNC117 + 10-1074 combination therapy and 3BNC117 or 10-1074 monotherapy (see (a)). Table shows model performances measured as Akaike information criterion for the different linear mixed effects models with time point as fixed effect. Column 'with treatment' has treatment (categories "3BNC117 + 10-1074 combination therapy" and "3BNC117 or 10-1074 monotherapy") as additional fixed effect. Correlation between time points of a participant was modeled based on the order of measurements with the models shown in the column 'correlation'. Based on the linear mixed effects model analysis using the corSymm class of the R package nlme (version 3.1-131), we observed a significant difference between 3BNC117 + 10-1074 combination therapy and 3BNC117 or 10-1074 monotherapy ($P=0.00018$).

91C22

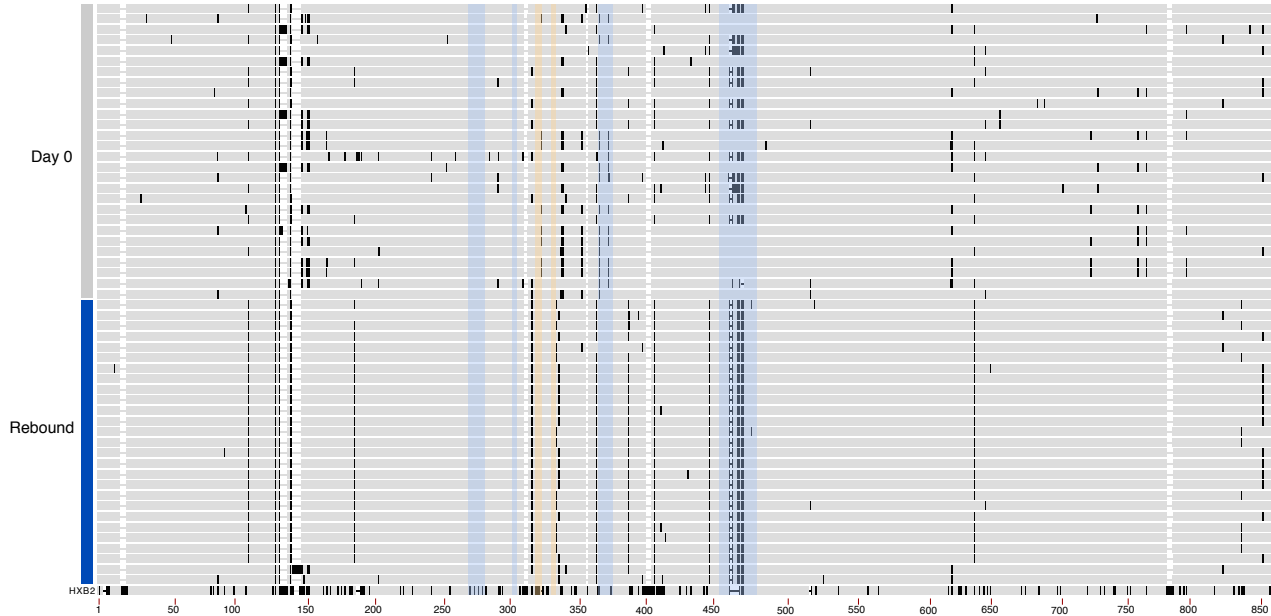


Supplementary Figure S6. Viral sequence analysis.

Multiple sequence alignment of Env amino acids obtained by SGA from plasma samples at day 0 and during viral rebound. Tick marks denote differences from day 0 consensus sequence. Blue shading depicts 3BNC117 binding sites and orange shading depicts 10-1074 binding sites. Sequences are numbered relative to HXB2. The bottom panel depicts the shaded areas in the upper panel. Dots depicts residues which are identical to the day 0 consensus sequence. Asterisks represents sequences that were used to generate pseudoviruses.

Supplementary Figure S6. Viral sequence analysis (continued)

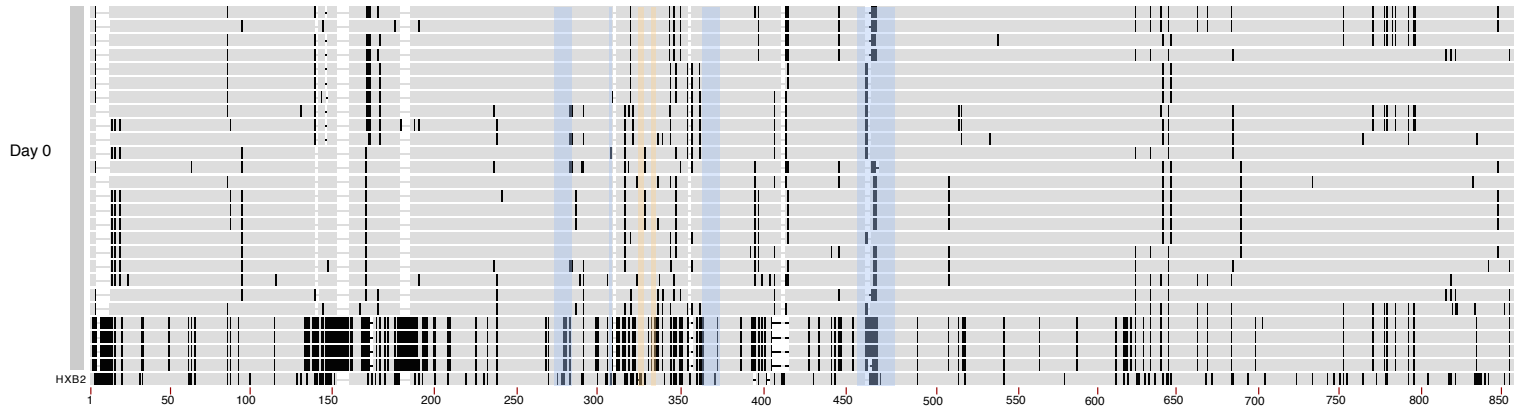
91C35



	274-283 (D loop)	308	364-374 (CD4 binding loop)	455-475 (B23 V5 loop)	324-327	332-334
HXB2	S V N F T D N A K T R	S S G G D P E I V T H	T R D G G N S - - - - N N E S E I F R P G G G D M	G N M R	N I S	
91C35_D0_consensus	S A N F S D N T K T T	H S G G D P E I V M H	T R D G G N A N T T E D G S N K N E T F R P G G G D M	G D I R	N I S	
91C35_D0_3HG-010517-D6_S46						
91C35_D0_3HG-010517-D11_S47		S	V			
91C35_D0_3HG-011317-C2_S29						
91C35_D0_3HG-011317-C7_S31		S	V			
91C35_D0_3HG-011317-D9_S32		S				
91C35_D0_3HG-011317-D10_S33						
★ 91C35_A1				S - S	S D	N T
★ 91C35_A2				S - S	S D	N T
★ 91C35_A3						
91C35_D0_3HG-031317P3-B9_S15				S - S	S D	N T
91C35_D0_3HG-031317P3-D2_S16						
★ 91C35_A4				S - S	S D	N T
★ 91C35_A5		P	V			
91C35_D0_3HG-031417P3-A10_S34		P	V			
★ 91C35_A7				S - S	S D	N T
91C35_D0_3HG-031417P3-F7_S40		S	V			
★ 91C35_A10		S	V			
91C35_D0_3HG-031417P3-F11_S42			E			
91C35_D0_3HG-031417P3-G11_S43				S - S	S D	N T
91C35_D0_3HG-031417P3-H1_S44		P	V			
★ 91C35_A12				S - S	S D	N T
91C35_D0_3HG-031617P1-F8_S63		P	V			
91C35_D0_3HG-031617P2-A3_S64		P	V			
91C35_D0_3HG-031617P2-C3_S65		P	M			
★ 91C35_B1		P	V			
91C35_D0_3HG-031617P2-D10_S67		P	V			
91C35_D0_3HG-031617P2-E9_S68		P	V			
91C35_D0_112216-A5_S26		P	V		S	R
★ 91C35_B6						
91C35_rebound_W3-3HG-012317-A8_S47						
91C35_rebound_W3-3HG-012317-B1_S49						
91C35_rebound_W3-3HG-012317-B8_S50						
91C35_rebound_W3-3HG-012317-C5_S53						
91C35_rebound_W3-3HG-012317-C9_S54						
91C35_rebound_W3-3HG-012317-C11_S56						
91C35_rebound_W3-3HG-012317-E4_S59						
★ 91C35_B3						
91C35_rebound_W3-3HG-012317-F7_S61						
91C35_rebound_W3-3HG-012317-G10_S63						
91C35_rebound_W3-3HG-012317-H8_S65						
91C35_rebound_W3-3HG-012317-H9_S66						
91C35_rebound_W3-3HG-030217P5-G12_S3						
91C35_rebound_W3-3HG-030617P3-A12_S14						
★ 91C35_B2						
91C35_rebound_W3-3HG-030617P3-F2_S16						
91C35_rebound_W3-3HG-030617P4-A3_S17						
★ 91C35_B5						
91C35_rebound_W3-3HG-030817P2-G6_S20						
91C35_rebound_W3-3HG-031317-A4_S21						
91C35_rebound_W3-3HG-031317-B3_S23						
91C35_rebound_W3-3HG-031317-B4_S24						
91C35_rebound_W3-3HG-031317-D8_S27						
91C35_rebound_W3-3HG-031317-E5_S28						
91C35_rebound_W3-3HG-031317-G9_S30						
91C35_rebound_W3-3HG-031317-G10_S31						

Supplementary Figure S6. Viral sequence analysis (continued)

91C33



	274-283 (D loop)	308	364-374 (CD4 binding loop)	455-475 (β23 V5 loop)	324-327	332-334
HXB2	S V N F T D N A K T	R	S S G G D P E V T H	T R D G G N - - - S N N E S E I F R P G G G D M	G N M R	N I S
91C33_D0_consensus	S E N F S N N A K I	H	H S G G D P E V V T H	T R D G G N - - - N S S A N E T F R P G G G D M	G D I R	N I S
91C33_D0_9-4-c4_S46 N H T
91C33_D0_9-6-d2b_S80 N H T
91C33_D0_9-6-e8b_S86 N H T
91C33_D0_abplate-a4b_S74 N H T
91C33_D0_9-6-d4b_S81 S G -
* 91C33_A3 S G -
91C33_D0_9-6-g8_S36 S G -
91C33_D0_23-8-h4b_S72	.	T S G -
91C33_D0_9-4-a10_S44 S G -
91C33_D0_9-6-e1b_S84	.	T S G -
91C33_D0_9-6-g7_S35	.	N S G -
91C33_D0_9-4-b10_S45	.	T - - - H T N -
91C33_D0_9-6-c1b_S79 - - - T N
91C33_D0_9-6-A5_S16 - - - T N
91C33_D0_9-6-f10b_S91 - - - T N
* 91C33_A5 - - - T N
* 91C33_A4 S G -
91C33_D0_9-6-f2_S28 - - - T N
91C33_D0_9-6-g3b_S93	.	T - - - T N
91C33_D0_9-6-g11_S38 - - - T N
91C33_D0_9-6-h3_S40 - - - N H T
91C33_D0_23-8-d10_S8 S G -
* 91C33_A6	.	A S H N	S M K T N T T E T F	T N
91C33_D0_23-8-g7_S10	.	A S H N	S M K T N T T E T F	T N
* 91C33_A9	.	A S H N	S M K T N T P E T F	T N
91C33_D0_9-6-f7_S30	.	A S H N	S M K T - - T E T F	T N

Supplementary Figure S6. *Viral sequence analysis (continued)*

91C34

Day 0

Week 6

Rebound

HXB2

1 50 100 150 200 250 300 350 400 450 500 550 600 650 700 750 800 850

274-283 (D loop)

HXB2 S N F T D N A K T

364-374 (CD4 binding loop)

S S G G D P E I V T H

455-475 (β23 V5 loop)

T R D G G N S N N E - - - S E I F R P G G G D M

324-327

G N M R N I S

332-334

G D I R N I T

91C34_D0_consensus

91C34_D0_YBAC10_S28

* 91C34_day0_#2

91C34_D0_YBBO3_S14

* 91C34_day0_#4

91C34_D0_YBCB3_S18

91C34_D0_YBCD3_S20

91C34_D0_YBCG11_S22

91C34_D0_YBD2_S24

91C34_D0_YBDA3_S1

91C34_D0_YBD4_S2

91C34_D0_YBDB1_S3

91C34_D0_YBDC12_S4

* 91C34_day0_#10

* 91C34_day0_#1

91C34_w6_Plat22C12_S36

91C34_w6_Plate1-D7_S28

91C34_w6_Plate1-D12_S29

91C34_w6_Plate2A2_S33

91C34_w6_Plate2A3_S34

91C34_w6_Plate2B8_S35

91C34_w6_Plate2E3_S38

91C34_w6_Plate2G11_S51

91C34_w6_Plate2H2_S42

* 91C34_W6#4

91C34_w6_Plate3A11_S44

91C34_w6_Plate3B1_S45

91C34_w6_Plate3B12_S46

91C34_w6_Plate3E3_S48

91C34_w6_Plate3F2_S49

91C34_w6_Plate3F3_S50

* 91C34_W6#6

91C34_w6_Plate4B7_S53

91C34_w6_Plate4E5_S54

91C34_w6_Plate4G8_S56

91C34_w6_Plate4H11_S57

91C34_w6_Plate4J2_S58

91C34_w6_Plate4A6_S59

91C34_w6_Plate5B1_S60

91C34_w6_Plate5C7_S61

91C34_w6_Plate5C8_S62

91C34_w6_Plate6A5_S63

91C34_w6_Plate6B4_S67

91C34_w6_Plate6C5_S68

91C34_w6_Plate6D6_S69

91C34_w6_Plate6D12_S72

91C34_w6_PlateF4_S30

91C34_w6_PlateG6_S31

91C34_w6_PlateH11_S32

91C34_w6_W6C87_S77

91C34_w6_W69C0A1_S73

91C34_w6_W69C0A8_S76

91C34_rebound_w12_11_S47

91C34_rebound_w12_24831-A3_S12

91C34_rebound_w12_24831-A5_S13

91C34_rebound_w12_24831-A12_S17

* 91C34_w12_#4

91C34_w12_#8

91C34_rebound_w12_24831-B10_S21

* 91C34_W12_#1

91C34_rebound_w12_24831-C2_S2

* 91C34_W12#2

* 91C34_w12_#7

91C34_rebound_w12_24831-d2_S7

91C34_rebound_w12_24831-d7_S9

* 91C34_w12_#5

91C34_rebound_w12_plate7-D1_S25

91C34_rebound_w12_w12-1-a3_S9

91C34_rebound_w12_w12-1-a6_S10

91C34_rebound_w12_w12-1-b3_S13

91C34_rebound_w12_w12-2-a5_S1

91C34_rebound_w12_w12-2-a11_S3

91C34_rebound_w12_w12-2-b1_S5

* 91C34_w12_#6

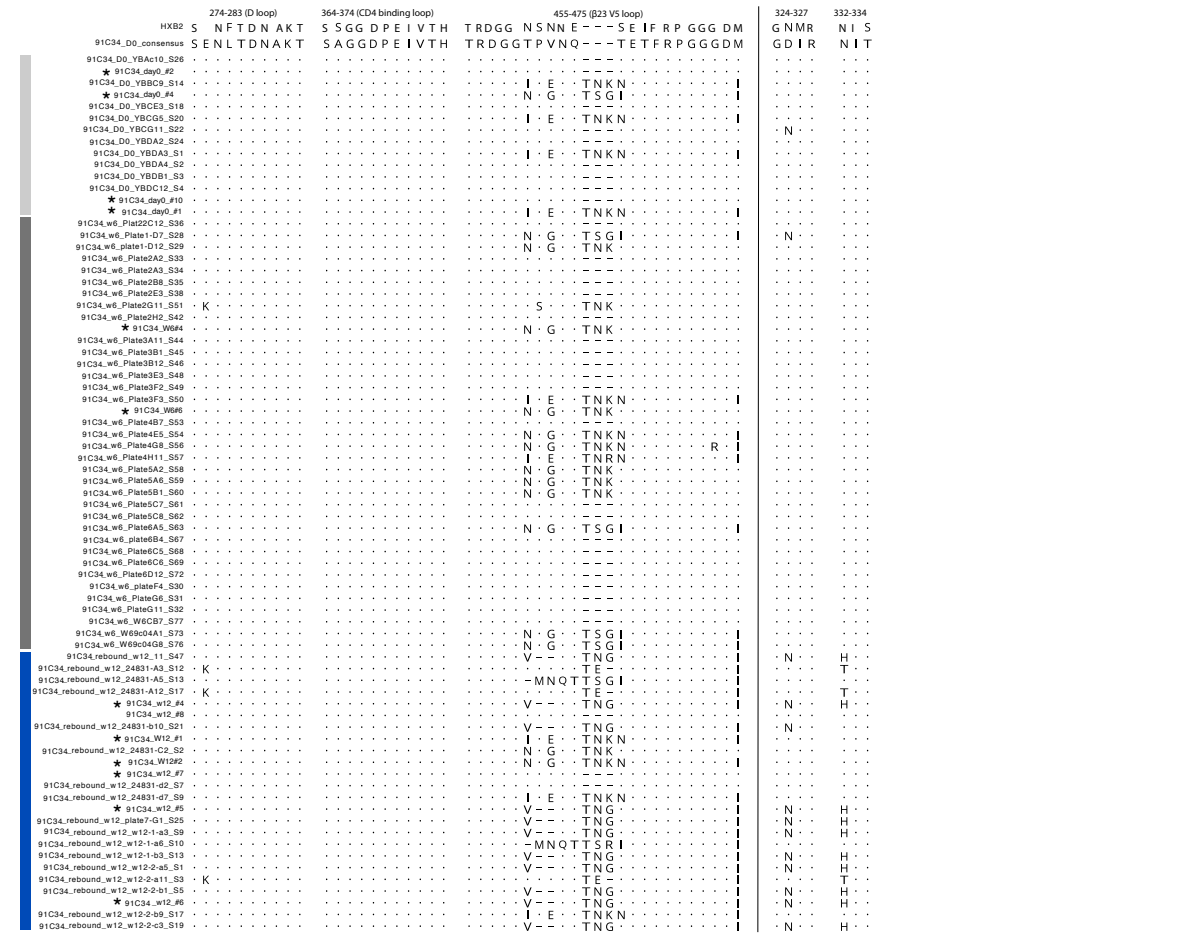
91C34_rebound_w12_w12-2-b6_S17

91C34_rebound_w12_w12-2-c3_S19

Day 0

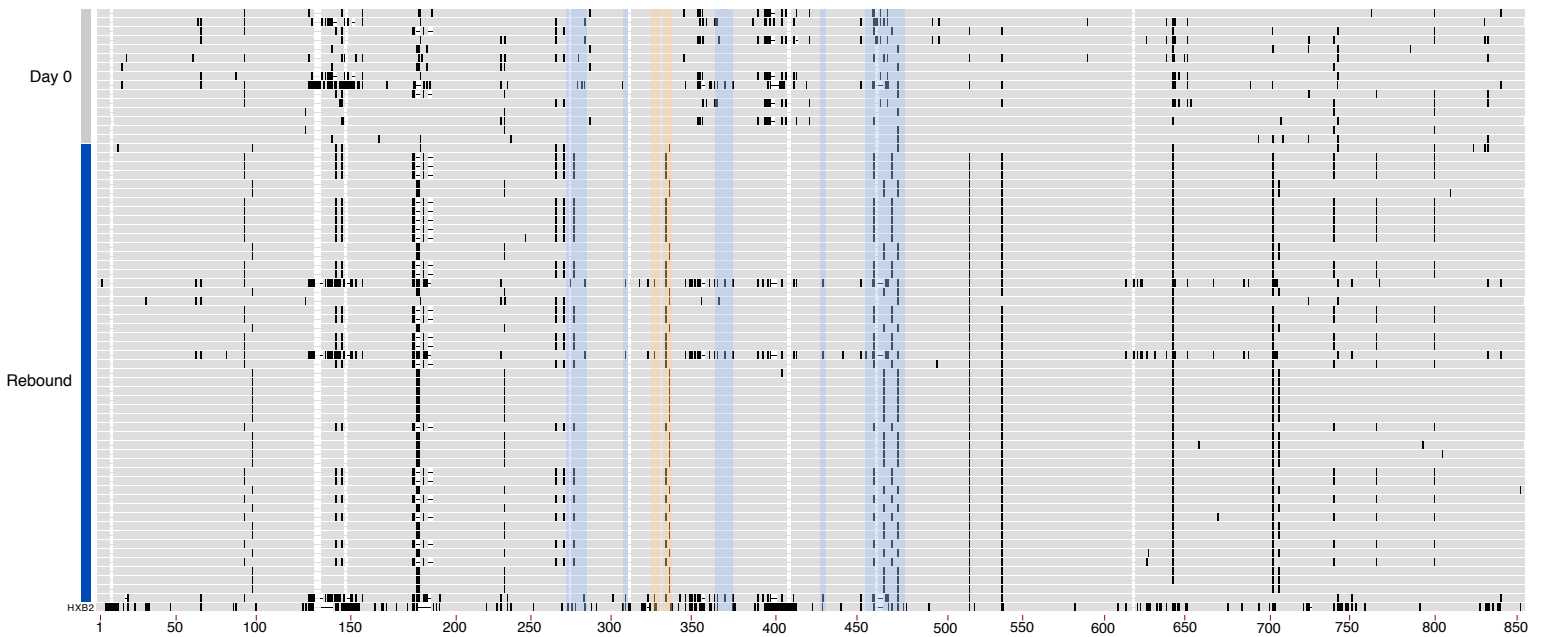
Week 6

Rebound



Supplementary Figure S6. Viral sequence analysis (continued)

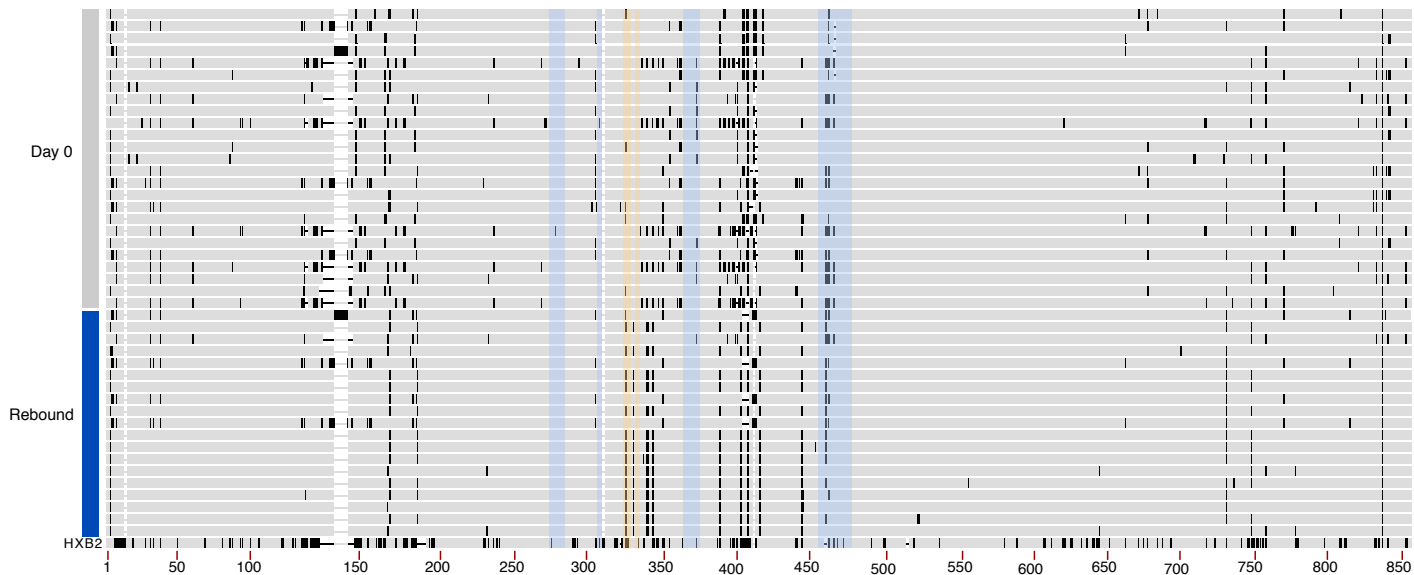
9341



	274-283 (D loop)	308	364-374 (CD4 binding loop)	428-430	455-475 (β23 V5 loop)	324-327	332-334
HXB2	S - VNF T DN AKT	R	S SGGDPEI VTH	QKV	TRDGGNS - - - - NNESEIFRP	GGGDM	GNMR NIS
9341_D0_consensus	- DNLTDNAKN	H	PSGGDPEI VRL	QEV	TRDGGNN - - - - NDTNNETETFR	PAGGNM	GDIR NIS
9341_D0_1-2812A3_S72DT - - V - - K -
9341_D0_1-2812A11_S74KGA - E - D - K -
*9341_A12T - - - - - - - - - I -
9341_D0_1-2812B11_S79T - - - - - - - - - E -
9341_D0_1A-1118E9_S96T - - - - - - - - - E -
9341_D0_1A-1118F5_S89T - - - - - - - - - K - G -
*9341_B2AT - - - - - - - - - K - G -
*9341_A5T - - - - - - - - - V - - -
*9341_B4T - - - - - - - - - V - - -
9341_D0_1A-1118H7_S82T - - - - - - - - - V - - -
9341_D0_1A-1118H8_S83T - - - - - - - - - V - - -
*9341_B5T - - - - - - - - - V - - -
9341_D0_1b-1118G7_S87T - - - - - - - - - - - - -
9341_D0_1b-1118G11_S68T - - - - - - - - - - - - -
9341_D0_1b-1118H5_S71T - - - - - - - - - - - - -
9341_w4_1-8-E2_S32T - - - - - - - - - - - - -N
9341_w4_1-8-E4_S33T - - - - - - - - - - - - -N
9341_w4_1-8-F2_S34T - - - - - - - - - - - - -N
9341_w4_aA2_S1T - - - - - - - - - - - - -N
9341_w4_aA4_S2T - - - - - - - - - - - - -N
9341_w4_aA7_S4T - - - - - - - - - - - - -N
9341_w4_aA8_S5T - - - - - - - - - - - - -N
9341_w4_aA10_S6T - - - - - - - - - - - - -N
9341_w4_aB5_S7T - - - - - - - - - - - - -N
9341_w4_ab10_S9T - - - - - - - - - - - - -N
9341_w4_aD7_S12T - - - - - - - - - - - - -N
9341_w4_aG7_S13T - - - - - - - - - - - - -N
9341_w4_BA3_S14T - - - - - - - - - - - - -N
9341_w4_BA4_S15T - - - - - - - - - - - - -N
9341_w4_BA6_S17T - - - - - - - - - - - - -N
9341_w4_BA7_S18T - - - - - - - - - - - - -N
9341_w4_BB1_S19T - - - - - - - - - - - - -N
9341_w4_BB2_S20T - - - - - - - - - - - - -N
*9341_B2T - - - - - - - - - - - - -N
9341_w4_BB6_S22T - - - - - - - - - - - - -N
9341_w4_BD9_S25T - - - - - - - - - - - - -N
9341_w4_CD6_S26T - - - - - - - - - - - - -N
9341_w4_CD8_S27T - - - - - - - - - - - - -N
9341_w4_CD9_S28T - - - - - - - - - - - - -N
9341_w4_CD3_S29T - - - - - - - - - - - - -N
9341_w4_CD2_S30T - - - - - - - - - - - - -N
9341_w4_CD8_S31T - - - - - - - - - - - - -N
9341_w5_1-8-C1_S50T - - - - - - - - - - - - -N
9341_w5_1-8-C3_S52T - - - - - - - - - - - - -N
9341_w5_1-8-C4_S53T - - - - - - - - - - - - -N
9341_w5_1-8-C6_S54T - - - - - - - - - - - - -N
9341_w5_1-8-C7_S55T - - - - - - - - - - - - -N
9341_w5_1-8-D1_S59T - - - - - - - - - - - - -N
9341_w5_1-8-D4_S61T - - - - - - - - - - - - -N
9341_w5_1-8-D6_S62T - - - - - - - - - - - - -N
9341_w5_1-8-E12_S68T - - - - - - - - - - - - -N
9341_w5_1-8-F5_S66T - - - - - - - - - - - - -N
9341_w5_1-8-F6_S70T - - - - - - - - - - - - -N
9341_w5_1-8-G3_S73T - - - - - - - - - - - - -N
9341_w5_1-8-G6_S74T - - - - - - - - - - - - -N
*9341_B7T - - - - - - - - - - - - -N
9341_w5_aD7_S37T - - - - - - - - - - - - -N
*9341_B5T - - - - - - - - - - - - -N
*9341_B6T - - - - - - - - - - - - -N
9341_w5_bb10_S41T - - - - - - - - - - - - -N
9341_w5_BE2_S42T - - - - - - - - - - - - -N
9341_w5_BE7_S44T - - - - - - - - - - - - -N
9341_w5_BE9_S45T - - - - - - - - - - - - -N
9341_w5_BE11_S46T - - - - - - - - - - - - -N
9341_w5_BE12_S47T - - - - - - - - - - - - -N
9341_w5_bI7_S49T - - - - - - - - - - - - -N

Supplementary Figure S6. Viral sequence analysis (continued)

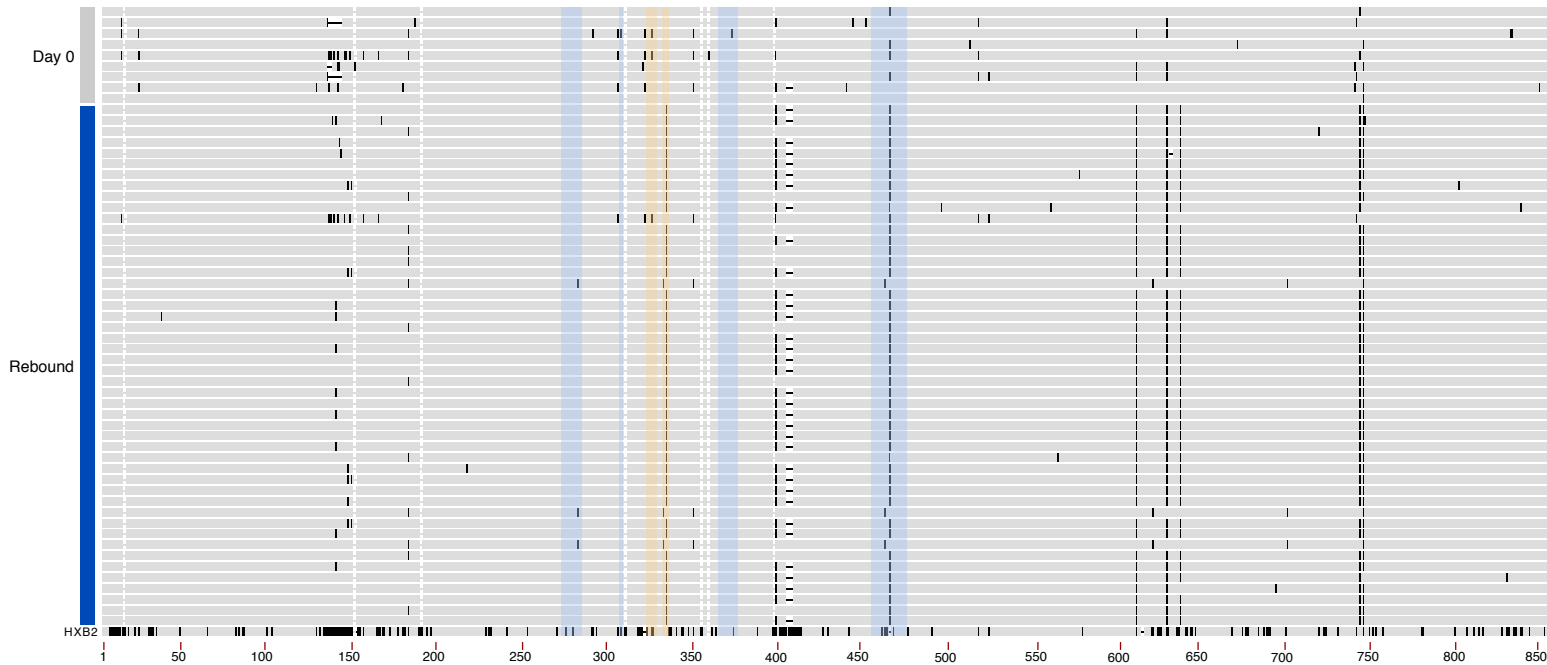
9342



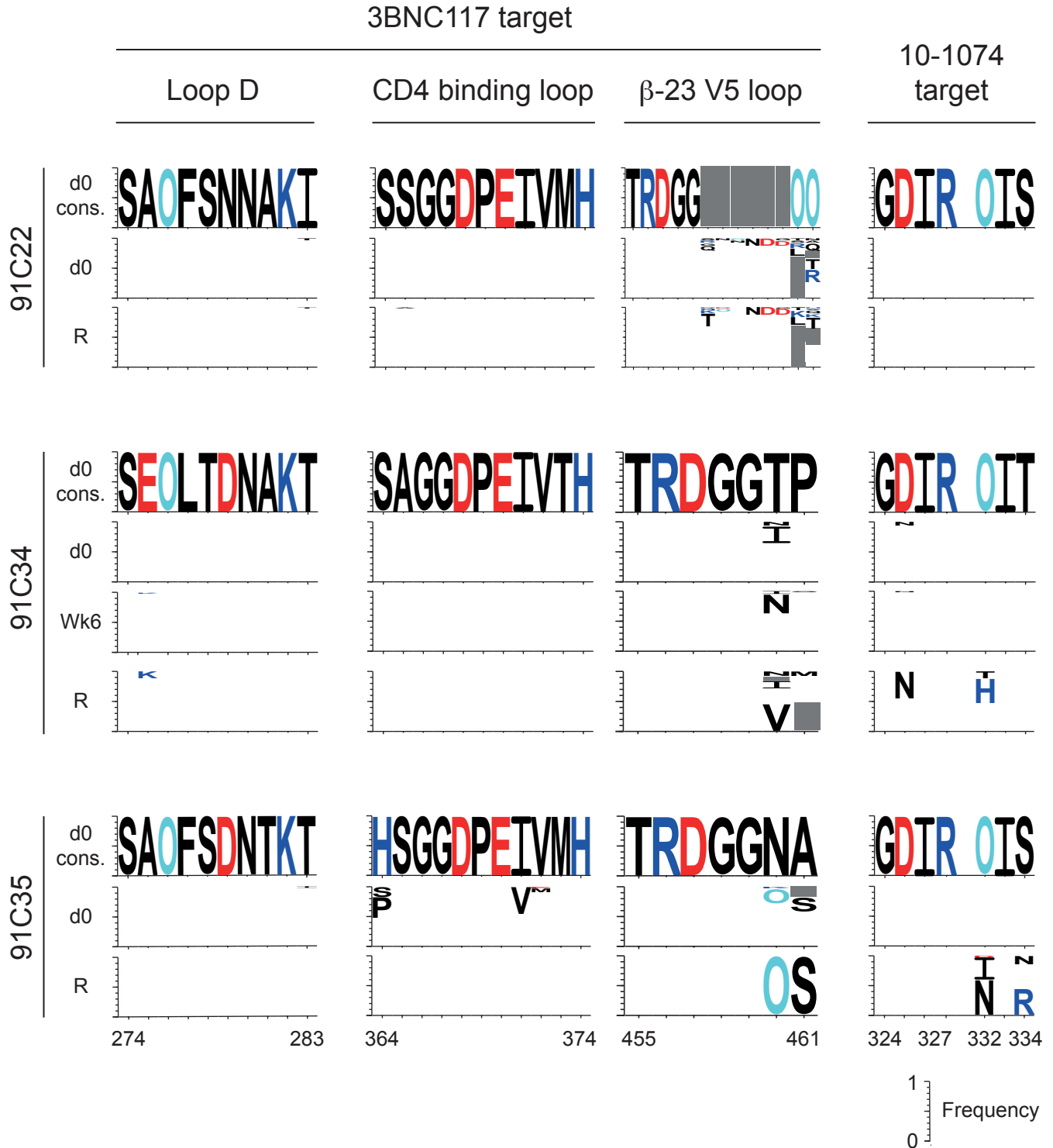
	274-283 (D loop)	308	364-374 (CD4 binding loop)	455-475 (β23 V5 loop)	324-327	332-334
HXB2	S V N F T D N A K T	R	S S G G D P E I V T H	T R D G G - N S N N E S E I F R P G G G D M	G N M R	N I S
9342_D0_consensus	S S N F T D N A K T	H	S S G G D P E I V M H	T R D G G G N N T N N N E T F R P T G G D M	G D I R	N L S
9342_D0_2-2812a6_S1	.	.	.	S	.	.
9342_D0_2-2812B3_S2	.	.	.	S	.	.
9342_D0_2-2812B9_S3	.	.	.	S	.	.
9342_D0_2-2812d3_S5
9342_D0_2-3012A1_S6	.	.	R	R G T S T T	.	.
9342_D0_2-3012A4_S7	.	.	.	S	.	.
9342_D0_2-3012A5_S8	.	.	G	.	.	.
9342_D0_2-3012A6_S9	.	.	G	R G T S T T	.	.
* 9342_B6	.	.	G	.	.	.
9342_D0_2-3012b10_S11	.	S	R	E G T S T T	.	.
9342_D0_2-3012b11_S12	.	.	G	.	.	.
9342_D0_2-3012b12_S13	.	.	G	.	N	.
9342_D0_2-3012c4_S14	.	.	G	.	.	.
* 9342_B7	.	.	.	S S	.	.
* 9342_B3	.	.	.	S S	.	.
* 9342_B5
* 9342_B8	N	.
9342_D0_2-3012D9_S20	.	.	.	S	N	.
* 9342_B2	.	K	.	E G T S T T	.	.
9342_D0_2-3012E7_S23	.	.	G	.	.	.
9342_D0_2-3012g6_S28	.	.	.	S S	.	.
* 9342_B1	.	.	R	R G T S T T	.	.
9342_D0_2-3012g12_S30	.	.	G	R G T S T T	.	.
9342_D0_2-3012h1_S31	N	.
9342_D0_2-3012h2_S32	.	.	.	E G T S T T	.	.
* 9342_A2	.	.	.	S S	G	.
9342_rebound_w20_1-3-18-a3_S49	.	.	.	V	N	.
* 9342_A1	.	.	G	R G T S T T	N	.
* 9342_A6	N	.
9342_rebound_w20_1-3-18-e2_S53	.	.	.	S S	.	.
9342_rebound_w20_1-3-18-h5_S57	.	.	.	V	K	.
9342_rebound_w20_1-3-18-g6_S56	.	.	.	V	K	.
* 9342_A3	.	.	.	S S	.	.
9342_rebound_w20_1-3-18-b8_S51	.	.	.	V	N	.
9342_rebound_w20_notfull-e11_S62	.	.	.	S S	.	.
9342_rebound_w20_notfull-f11_S64	.	.	.	V	N	.
9342_rebound_w20_full-e2_S65	.	.	.	V	N	.
9342_rebound_w20_full-e10_S66	.	.	.	V	N	.
9342_rebound_w20_full-f8_S67	.	.	.	V	N	.
* 9342_A8	.	.	.	V	N	.
9342_rebound_w20_notfull-e4_S61	.	.	.	S	N	.
9342_rebound_w20_1_S73	N	.
9342_rebound_w20_4_S76	.	.	.	V	N	.
9342_rebound_w20_7_S79	N	.

Supplementary Figure S6. Viral sequence analysis (continued)

9343



	274-283 (D loop)	308	364-374 (CD4 binding loop)	455-475 (beta23 V5 loop)	324-327	332-334
HXB2	S N F T N A K T R	S S G G D P E I V T H	T R D G G N S N N E S - E I F R P G G G D M	G N M R	N I S	
9343_D0_consensus	V D N	S S G G D P E I V M H	T R D G G N N N D T S A E V F R P G G G D M	G D I R	N I S	
9343_D0_3-2812-A2_S55	S E N F T N N A K T	.	.	T	.	
9343_D0_3-2812-A11_S59	
*9343_A1	
*9343_A6	
9343_D0_3-2812-d3_S63	.	.	.	T	.	
*9343_A3	
*9343_A5	.	.	.	T	.	
9343_D0_3B-1118-f3_S52	
*9343_A8	
9343_rebound_w7_-24-1-a11_S84	.	.	.	T	.	
9343_rebound_w7_-24-1-c12_S88	.	.	.	T	.	
9343_rebound_w7_-24-1-e6_S91	.	.	.	T	.	
9343_rebound_w7_-24-1-g7_S92	.	.	.	T	.	
9343_rebound_w7_A-25-1-b4_S13	.	.	.	T	.	
9343_rebound_w7_A-25-1-e8_S16	.	.	.	T	.	
9343_rebound_w7_A-25-1-f6_S17	.	.	.	T	.	
9343_rebound_w7_A-25-1-g10_S20	.	.	.	T	.	
9343_rebound_w7_A-25-1-h1_S22	.	.	.	T	.	
9343_rebound_w7_A-25-1-h4_S23	.	.	.	T	.	
9343_rebound_w7_b-25-1-A8_S1	.	.	.	T	.	
9343_rebound_w7_b-25-1-A11_S2	.	.	.	T	.	
9343_rebound_w7_b-25-1-A12_S3	.	.	.	T	.	
*9343_B5	.	.	.	T	.	
9343_rebound_w7_b-25-1-B9_S5	.	.	.	T	.	
9343_rebound_w7_b-25-1-d2_S7	.	.	.	T	.	
_rebound_w7_b-25-1-f12_S10	.	.	.	Y	.	
*9343_B9	.	E	.	.	T	
9343_rebound_w8_-24-1-c5_S67	.	.	.	T	.	
9343_rebound_w8_-24-1-d4_S71	.	.	.	T	.	
9343_rebound_w8_-24-1-d6_S72	.	.	.	T	.	
9343_rebound_w8_-24-1-d7_S73	.	.	.	T	.	
9343_rebound_w8_-24-1-e3_S74	.	.	.	T	.	
9343_rebound_w8_-24-1-e4_S75	.	.	.	T	.	
9343_rebound_w8_-24-1-f8_S77	.	.	.	T	.	
9343_rebound_w8_-24-1-g11_S81	.	.	.	T	.	
9343_rebound_w8_-25-1-A8_S25	.	.	.	T	.	
*9343_B6	.	.	.	T	.	
9343_rebound_w8_-25-1-A10_S27	.	.	.	T	.	
9343_rebound_w8_-25-1-A11_S28	.	.	.	T	.	
9343_rebound_w8_-25-1-b1_S29	.	.	.	T	.	
9343_rebound_w8_-25-1-b7_S31	.	.	.	T	.	
9343_rebound_w8_-25-1-b10_S33	.	.	.	T	.	
9343_rebound_w8_-25-1-b11_S34	.	.	.	T	.	
9343_rebound_w8_-25-1-c3_S36	.	.	.	T	.	
9343_rebound_w8_-25-1-c6_S37	.	.	.	T	.	
9343_rebound_w8_-25-1-c7_S38	.	.	.	T	.	
9343_rebound_w8_-25-1-c8_S39	.	.	.	T	.	
*9343_B4	.	E	.	Y	T	
9343_rebound_w8_-25-1-d3_S41	.	.	.	T	.	
*9343_B1	.	E	.	Y	T	
*9343_B2	.	.	.	T	.	
9343_rebound_w8_-25-1-f4_S53	.	.	.	T	.	
9343_rebound_w8_-25-1-f5_S54	.	.	.	T	.	
9343_rebound_w8_-25-1-g6_S60	.	.	.	T	.	
9343_rebound_w8_-25-1-g10_S61	.	.	.	T	.	
9343_rebound_w8_-25-1-g12_S62	.	.	.	T	.	
9343_rebound_w8_-25-1-h2_S63	.	.	.	T	.	



Supplementary Figure 7. *HIV-1 env* sequence logo plots

Logo plots of SGA-derived plasma HIV-1 Env amino acid frequencies at and around 3BNC117 and 10-1074 contact residues over time during 3BNC117/10-1074 combination therapy and compared to the day 0 consensus sequence (d0 cons.). Frequency is shown by the height of the letter. d0 indicates sequences obtained on day 0 pre-infusion, Wk6 indicates sequences obtained six weeks after the first antibody infusion and R indicates sequences obtained at viral rebound. White space indicates the lack of a change compared to the day 0 consensus. Single-letter amino acids highlight a change compared to day 0 and are colored for acidic (red) and basic (blue) amino acids. Turquoise “O” indicates an asparagine that acts as potential glycosylation site. Grey boxes indicate gaps. Logo plots were generated using LASSIE.

3BNC117 target

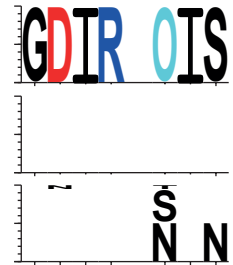
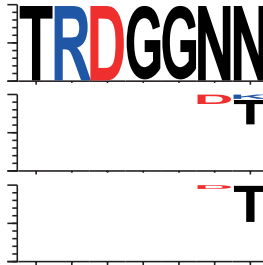
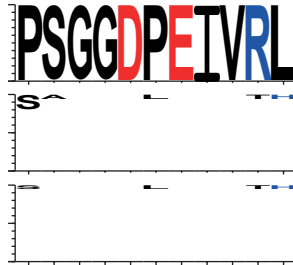
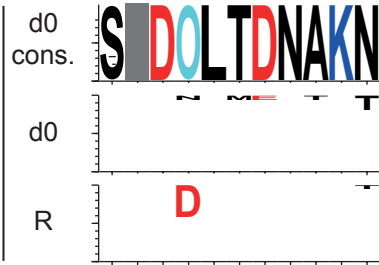
10-1074 target

Loop D

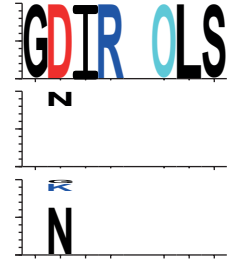
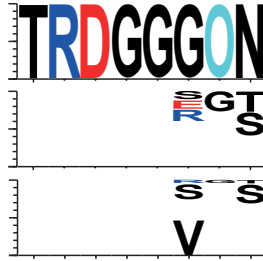
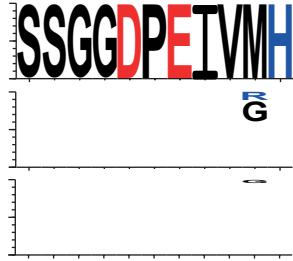
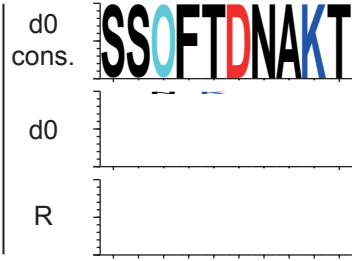
CD4 binding loop

β -23 V5 loop

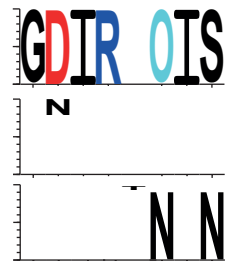
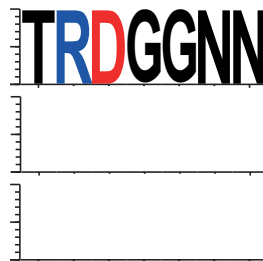
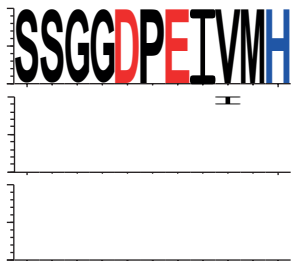
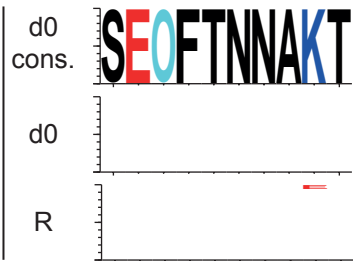
9341



9342



9343



274

283

364

374

455

461

324

327

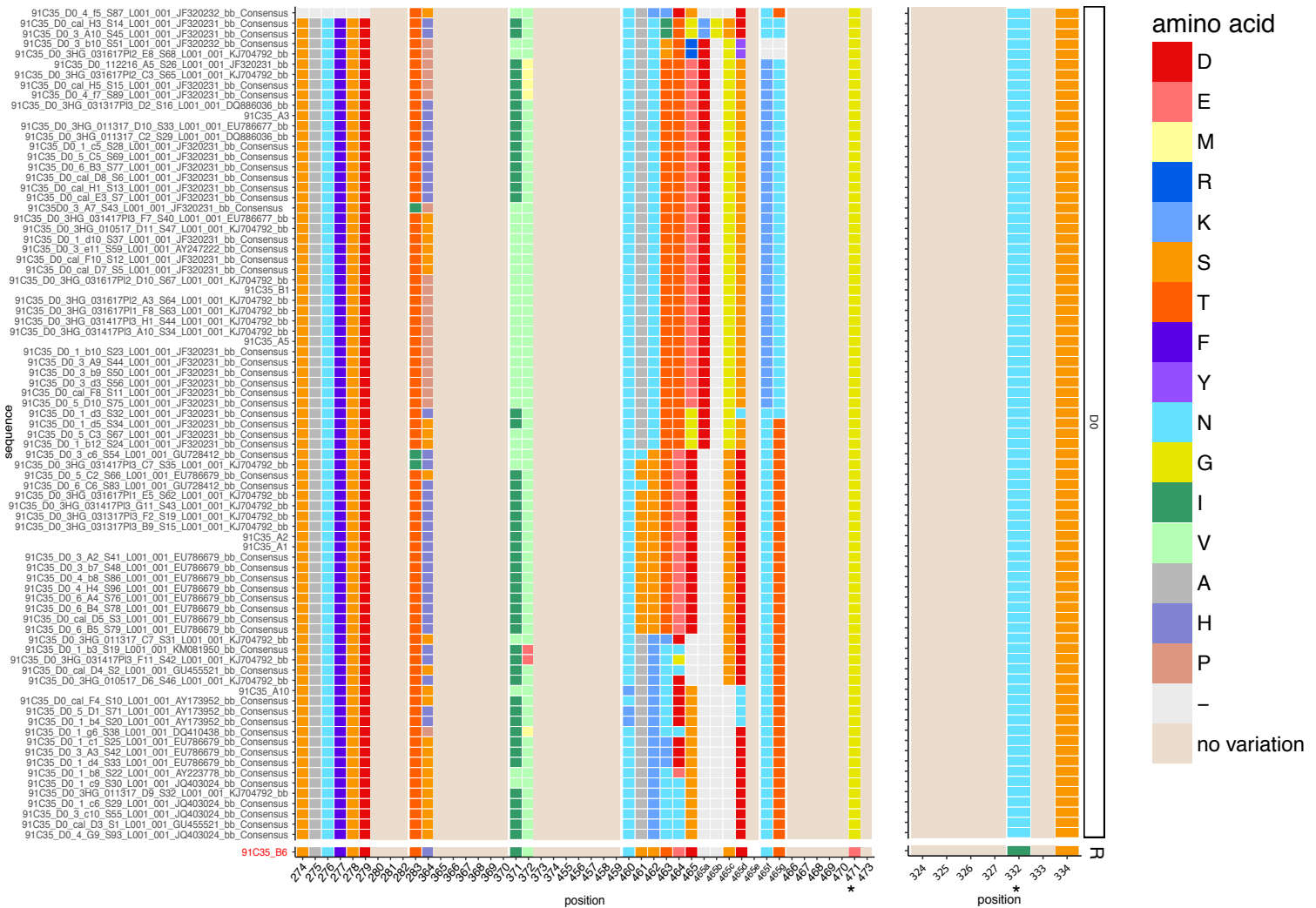
332

334

1
0
Frequency

Supplementary Figure 7. HIV-1 env sequence logo plots (continued)

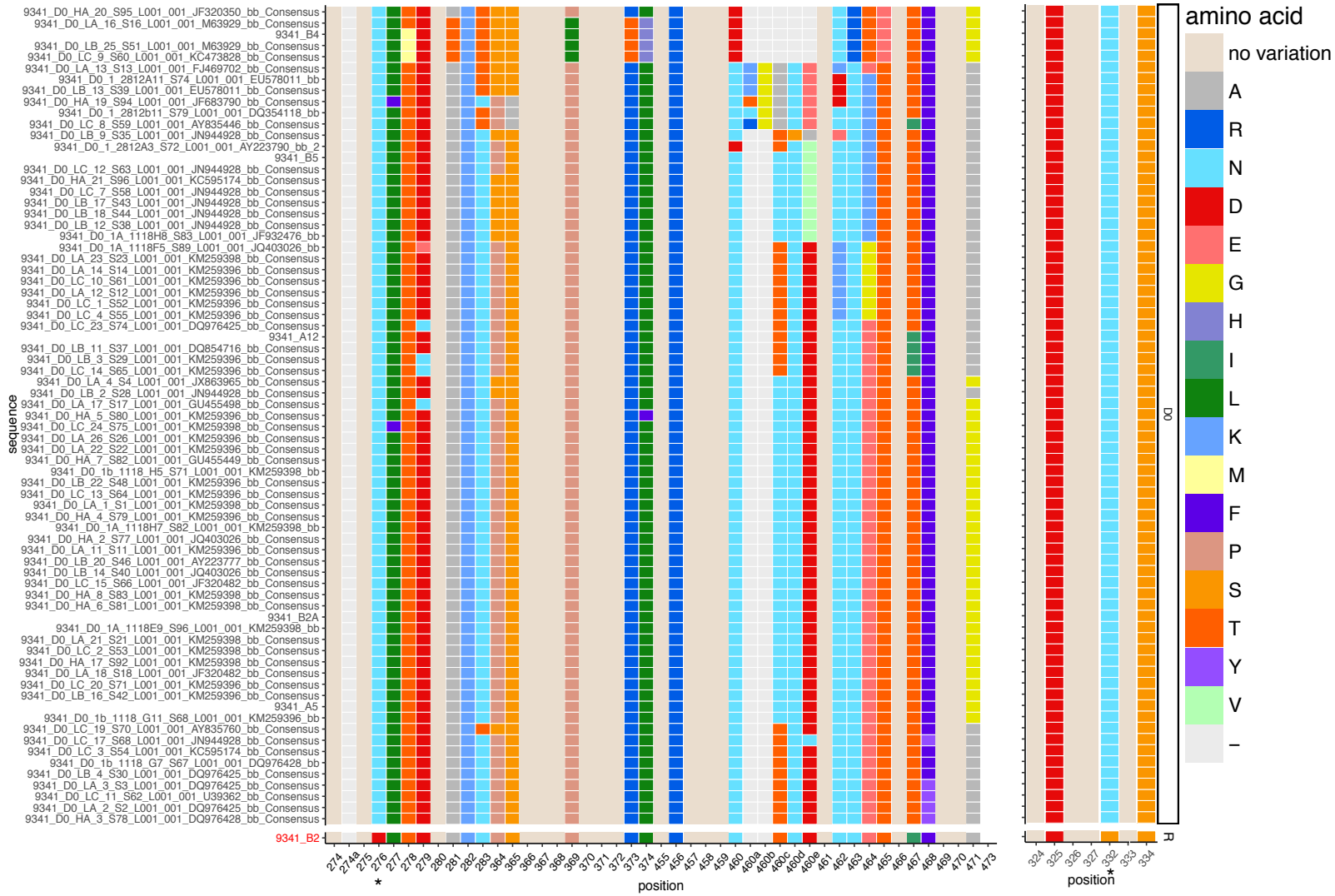
91C35



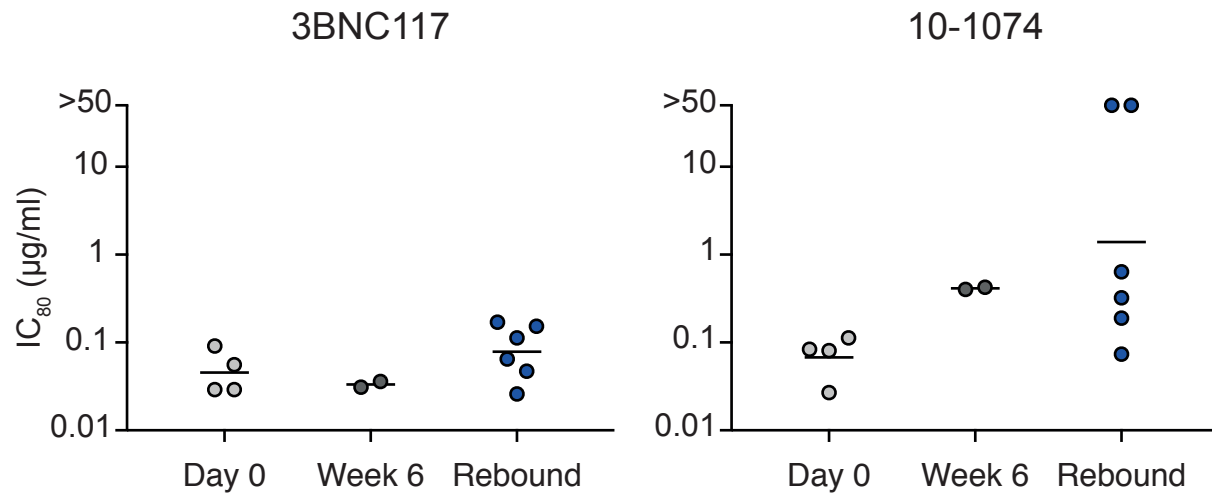
Supplementary figure S8. In-depth analysis of 3BNC117 and 10-1074 binding site mutations in participants 91C35 and 9341

Amino acids in and around known 3BNC117 and 10-1074 contact residues in Env sequences isolated by SGS from participants 91C35 and 9341. Amino acids are numbered according to HXB2. Upper panel shows sequences isolated from day 0 viruses and the bottom row depicts Env amino acids in resistant rebound virus. * represent instances in which an amino acid was found in the resistant rebound virus but not in day 0 sequences. The resistant rebound virus is highlighted in red.

9341



Supplementary figure S8. In-depth analysis of 3BNC117 binding site mutations in patients 91C35 and 9341 (continued)



Supplementary Figure 9. *Pseudovirus sensitivities of participant 91C34*

IC₈₀s of pseudoviruses constructed from SGA-derived *env* sequences (day 0, *n*=4; week 6, *n*=2; rebound, *n*=6) obtained at the indicated time points against 3BNC117 (left) and 10-1074 (right) determined in TZM-bl neutralization assay. Line indicates geometric mean IC₈₀ and each circle indicates a single pseudovirus.

Supplementary Table 1. BNab sensitivities of bulk culture-derived viruses from viremic individuals

a Pre-screening individual results

Trial ID	Pre-Screen ID	3BNC117		10-1074		Pre-Screen ID	3BNC117		10-1074		µg/ml
		IC ₅₀	IC ₈₀	IC ₅₀	IC ₈₀		IC ₅₀	IC ₈₀	IC ₅₀	IC ₈₀	
91C22	B321	0.084	0.187	0.028	0.087	0536	0.130	0.491	>20	>20	>20
91C33	0167	0.875	2.592	0.346	0.801	0537	1.115	4.103	>20	>20	< 0.5
91C34	0456	0.114	0.393	0.087	0.235	0541	1.855	6.025	0.148	0.412	0.5 - 1.0
91C35	B315	1.018	3.382	0.037	0.097	0549	6.662	19.690	1.097	3.828	1.0 - 2.0
9341	B450	1.059	3.679	1.244	4.445	0561	4.108	15.937	0.324	3.192	2.0 - 5.0
9342	0648	0.529	1.359	0.140	0.355	0615	0.295	0.799	>20	>20	5.0 - 20.0
9343	0787	0.125	0.567	0.058	0.191	0771	>25	>25	>25	>25	20
	0034	1.341	3.704	0.029	0.069	0775	0.397	1.298	0.365	0.947	
	0039	5.238	>20	>20	>20	0779	>20	>20	>20	>20	
	0277	1.066	2.592	5.652	13.627	0782	>20	>20	11.543	>20	
	0338	0.125	0.385	0.044	0.130	0795	>25	>25	>25	>25	
	0355	4.192	13.086	0.097	0.214	0799	1.900	6.120	5.720	19.510	
	0359	1.036	3.807	4.903	>20	0801	2.230	6.260	>25	>25	
	0403	>20	>20	0.900	3.598	0806	>25	>25	0.480	1.920	
	0422	0.318	1.070	0.003	0.008	0821	0.650	2.100	2.760	10.530	
	0423	0.987	3.306	>20	>20	B270	0.720	2.791	>20	0.043	
	0428	1.650	6.583	0.295	3.185	B312	>20	>20	0.861	2.047	
	0438	0.761	3.165	0.016	0.060	B331	1.531	7.140	>20	>20	
	0440	1.489	4.112	>20	>20	B340	0.461	1.917	>20	>20	
	0464	0.313	1.099	8.282	20.000	B367	2.073	6.230	1.870	6.582	
	0472	0.083	0.383	0.140	0.373	B394	1.013	5.932	0.230	1.211	
	0473	0.530	1.861	0.025	0.076	B407	0.044	0.193	0.017	0.057	
	0480	0.492	2.037	0.041	0.165	B411	2.018	5.255	1.357	2.958	
	0508	>20	>20	>20	>20	B434	>20	>20	0.454	1.431	
	0527	>20	>20	>20	>20	B461	0.250	1.161	6.029	16.394	
	0535	0.334	0.989	0.187	0.704	B465	0.325	2.481	0.305	1.712	

b Pre-screening summary

Antibody	IC ₅₀ <2 µg/ml	
	n	%
3BNC117	35	67.3%
10-1074	30	57.7%
3BNC117 and 10-1074	21	40.4%

Table indicates IC₅₀s and IC₈₀s of 3BNC117 and 10-1074 against bulk CD4⁺ T cell outgrowth culture-derived viruses obtained from viremic HIV-1-infected individuals. (a) Individual culture results. (b) Summary of all (n=52) tested cultures.

Supplementary Table 2. Individual participant baseline demographics

Study ID	Dose / Regimen	Age (years)	Gender	Race	Years since HIV-1 diagnosis	ART at enrollment*	HIV-1 Clade	HIV-1 RNA at day 0 (copies/ml)	CD4+ T cell count at day 0 (cells/μl)	Pre-Screen Sensitivity (μg/ml)**			
										3BNC117		10-1074	
										IC50	IC80	IC50	IC80
1A12	10 mg/kg (ea.) day 0	53	F	American Indian/Hisp	23	EFV, TDF, FTC	n.d.	<20	809		-		
1A27	10 mg/kg (ea.) day 0	47	M	Black	16	RPV, TDF, FTC	n.d.	<20	648		-		
1A29	10 mg/kg (ea.) day 0	45	M	Black	16	RPV, TDF, FTC	n.d.	<20	558		-		
1A71	10 mg/kg (ea.) day 0	55	M	Black	10	EFV, TDF, FTC	n.d.	<20	1,033		-		
1B05	30 mg/kg (ea.) day 0	34	M	Black	17	EFV, TDF, FTC	n.d.	<20	711		-		
1B44	30 mg/kg (ea.) day 0	51	M	Black	18	ATV/r, TDF, FTC	n.d.	<20	754		-		
1B55	30 mg/kg (ea.) day 0	56	F	Black	25	RAL, ETR, ABC, 3TC, ZDV	n.d.	<20	814		-		
1B61	30 mg/kg (ea.) day 0	64	M	Black	19	DTG, ABC, 3TC	n.d.	<20	602		-		
91C22	30 mg/kg (ea.) day 0	58	F	Black	4	- (naïve)	B	730	586	0.084	0.187	0.028	0.087
91C33	30 mg/kg (ea.) day 0	28	M	White	5	- (naïve)	B	1,750	750	0.875	2.592	0.346	0.801
91C34	30 mg/kg (ea.) day 0	32	M	White	8	- (naïve)	B	65,800	520	0.114	0.393	0.087	0.235
91C35	30 mg/kg (ea.) day 0	26	M	Mult/Hisp	4	- (experienced)†	B	53,540	320	1.018	3.382	0.037	0.097
9341	30 mg/kg (ea.) day 0 - 14 - 28	48	F	Black	22	- (experienced)‡	B	23,610	682	1.059	3.679	1.244	4.445
9342	30 mg/kg (ea.) day 0 - 14 - 28	41	M	White	15	- (naïve)	B	2,550	780	0.529	1.359	0.140	0.355
9343	30 mg/kg (ea.) day 0 - 14 - 28	42	M	White	0.2	- (naïve)	B	97,800	350	0.125	0.567	0.058	0.191

* RPV - rilpivirine, TDF - tenofovir disoproxil fumarate, FTC - emtricitabine, EFV - efavirenz, ATV - atazanavir, /r - ritonavir-boosted, RAL - raltegravir, ETR - etravirine, ABC - abacavir, 3TC - lamivudine, ZDV - zidovudine.

** Pre-screening of virus obtained from CD4+ T cell bulk outgrowth culture for antibody sensitivity by TZM-bl assay.

† 91C35 initiated ART (elvitegravir, cobicistat, TDF, FTC) within six months of presumed HIV-1 exposure. This regimen was maintained for four years until he chose to discontinue it. At the time of sensitivity screening and on the day of the first infusion, participant had been off ART for six and eight months, respectively.

‡ 9341 initiated ART within a year of diagnosis and was discontinued shortly thereafter. Since then, she had multiple periods of intermittent ART and reports receiving multiple different ART regimens. At the time of enrollment, she had been off ART for at least 12 months.

Table indicates baseline parameters of ART-treated and viremic study participants.

Supplementary Table 3. HIV-1 RNA levels, CD4⁺ T cell counts and bNAb serum levels

Viremic Single infusion, day 0 3BNC117 + 10-1074 (30 mg/kg each i.v.)																	
91C22	HIV-1 RNA		CD4 ⁺ T cells		3BNC117		10-1074		91C33	HIV-1 RNA		CD4 ⁺ T cells		3BNC117		10-1074	
	cp/ml	Δlog ₁₀	cells/μl	%	ELISA	TZM-bi	ELISA	TZM-bi		cp/ml	Δlog ₁₀	cells/μl	%	ELISA	TZM-bi	ELISA	TZM-bi
Study visit					Q769.d22	Du422.1	X2088_c9	Du422.1	Study visit					Q769.d22	X2088_c9	Du422.1	Du422.1
	cp/ml	Δlog ₁₀	cells/μl	%	μg/ml					cp/ml	Δlog ₁₀	cells/μl	%	μg/ml			
Screen	1,430	-	488	21.0	-	-	-	-	Screen	279	-	900	20	-	-	-	-
Wk -2 (+7d)	640	-	617	20.0	-	-	-	-	Wk -2 (+3)	421	-	800	21	-	-	-	-
Day 0 baseline	730	-	586	20.0	<0.51	<0.46	<0.14	<0.08	Day 0 baseline	1,750	-	750	25	<0.51	<0.46	<0.14	<0.10
Day 0 end 3BNC117	-	-	-	-	996.19	860.92	<0.14	<0.08	Day 0 end 3BNC117	-	-	-	-	800.29	618.63	<0.14	<0.08
Day 0 end 10-1074	-	-	-	-	994.78	1,443.37	1,561.17	5,123.32	Day 0 end 10-1074	-	-	-	-	473.37	1,307.44	1,321.43	1,917.72
Day 2	720	-0.01	-	-	184.24	192.19	853.61	854.44	Day 2 (+1d)	839	-0.32	-	-	155.27	165.99	695.70	455.56
Wk 1	<20	-1.58*	-	-	92.64	108.93	450.08	711.26	Wk 1	575	-0.48	-	-	90.31	66.24	465.64	282.67
Wk 2	<20	-1.58*	249	17.8	29.05	32.06	152.75	140.71	Wk 2	1,040	-0.23	830	20	47.33	48.02	189.58	171.74
Wk 3	<20D	-1.58*	-	-	14.92	12.21	103.63	60.15	Wk 3	1,250	-0.15	-	-	30.09	63.23	180.44	113.08
Wk 4	<20	-1.58*	456	19.0	6.31	7.48	54.22	30.21	Wk 4 (+3d)	699	-0.40	690	20	17.34	22.03	114.45	57.44
Wk 6	<20D	-1.58*	-	-	1.31	1.38	13.51	10.32	Wk 6	961	-0.26	-	-	6.92	5.80	62.45	27.08
Wk 8	<20	-1.58*	576	24.0	<0.51	<0.46	8.31	3.34	Wk 8	1,520	-0.06	790	18	2.12	1.36	27.61	11.14
Wk 12 (-4d)	70	-1.02	763	24.6	<0.51	<0.46	1.50	0.69	Wk 12 (-6d)	249	-0.85	800	19	<0.51	<0.46	14.12	3.46
Wk 16	1,590	0.34	692	24.7	<0.51	<0.46	<0.14	<0.1	Wk 16 (-1d)	732	-0.38	630	21	<0.51	<0.46	2.18	0.63
Wk 20 (+1d)	740	0.01	533	20.5	<0.51	<0.46	<0.14	<0.1	Wk 20 (+2d)	614	-0.45	860	20	<0.51	<0.46	0.58	<0.1
Wk 24 (+3d)	1,020	0.15	349	16.6	<0.51	<0.46	<0.14	<0.1	Wk 24 (+2d)	722	-0.38	860	23	<0.51	<0.28	<0.14	<0.08

91C34	HIV-1 RNA		CD4 ⁺ T cells		3BNC117		10-1074		91C35	HIV-1 RNA		CD4 ⁺ T cells		3BNC117		10-1074	
	cp/ml	Δlog ₁₀	cells/μl	%	ELISA	TZM-bi	ELISA	TZM-bi		cp/ml	Δlog ₁₀	cells/μl	%	ELISA	TZM-bi	ELISA	TZM-bi
Study visit					Q769.d22	Du422.1	X2088_c9	Du422.1	Study visit					Q769.d22	X2088_c9	Du422.1	Du422.1
	cp/ml	Δlog ₁₀	cells/μl	%	μg/ml					cp/ml	Δlog ₁₀	cells/μl	%	μg/ml			
Screen	33,500	-	490	36	-	-	-	-	Screen	22,760	-	345	27.0	-	-	-	-
Wk -2 (+3)	33,900	-	500	33	-	-	-	-	Wk -2 (+7d)	15,120	-	382	29.0	-	-	-	-
Day 0 baseline	65,800	-	520	36	<0.51	<0.46	<0.14	<0.1	Day 0 baseline	53,540	-	320	26.7	<0.51	<0.46	<0.14	<0.08
Day 0 end 3BNC117	-	-	-	-	735.93	905.74	<0.14	<0.08	Day 0 end 3BNC117	-	-	-	-	611.21	716.79	<0.14	<0.08
Day 0 end 10-1074	-	-	-	-	665.80	945.12	889.07	1,242.07	Day 0 end 10-1074	-	-	-	-	739.37	1,440.54	1,224.26	1,154.49
Day 2	65,600	0.00	-	-	180.98	447.79	723.69	613.03	Day 2	-	-	-	-	N/A	N/A	N/A	N/A
Wk 1 (+1d)	2,520	-1.42	-	-	104.13	164.31	365.53	270.02	Wk 1	1,400	-1.58	-	-	94.36	82.52	385.14	265.15
Wk 2 (+2d)	1,850	-1.55	530	36	56.59	50.85	258.52	200.97	Wk 2	5,210	-1.01	469	27.6	42.87	41.12	253.06	172.42
Wk 3 (-1d)	1,430	-1.66	-	-	41.00	55.96	208.91	181.26	Wk 3 (+1d)	166,310	0.49	-	-	23.93	33.70	168.68	107.55
Wk 4	1,320	-1.70	510	34	23.04	88.32	182.26	85.10	Wk 4	56,850	0.03	356	25.4	16.62	31.86	127.75	63.31
Wk 6 (-1d)	888	-1.87	-	-	12.68	46.87	105.43	43.24	Wk 6	46,800	-0.06	-	-	6.42	5.47	51.06	29.46
Wk 8	810	-1.91	420	35	5.68	5.01	84.27	29.26	Wk 8 (+2d)	78,420	0.17	414	28.0	2.56	2.23	23.98	8.13
Wk 12 (+3d)	10,100	-0.81	340	41	1.56	<0.46	55.46	5.51	Wk 12	33,400	-0.20	459	35.3	<0.51	1.59	7.43	2.62
Wk 16	14,800	-0.65	320	36	<0.51	<0.46	3.52	1.16	Wk 16	N/A	-	N/A	N/A	N/A	N/A	N/A	N/A
Wk 20	14,100	-0.67	400	26	<0.51	<0.46	0.53	<0.1	Wk 20	N/A	-	N/A	N/A	N/A	N/A	N/A	N/A
Wk 24	24,800	-0.42	300	33	<0.51	<0.28	<0.14	<0.08	Wk 24	N/A	-	N/A	N/A	N/A	N/A	N/A	N/A

* HIV-1 RNA < 20 copies/ml. For calculation of log-change, a value of 19 copies/ml was set.
 < 20D: HIV-1 RNA detected but below limit of quantification.
 Green shaded area indicates HIV-1 RNA nadir.
 Grey shaded area indicates time on ART.

Table indicates clinical laboratory parameters and antibody serum concentrations as determined by ELISA and TZM-bi neutralization assay (using the indicated pseudoviruses) in viremic and ART-treated study participants.

Supplementary Table 3. HIV-1 RNA levels, CD4⁺ T cell counts and bNAb serum levels (continued)

Viremic
Three infusions, day 0 / 14 / 28
3BNC117 + 10-1074 (30 mg/kg each i.v.)

9341	HIV-1 RNA		CD4 ⁺ T cells		3BNC117		10-1074	
	cp/ml	$\Delta\log_{10}$	cells/ μ l	%	ELISA	TZM-bl	ELISA	TZM-bl
					Q769.d22	Q769.d22	X2088_c9	Du422.1
Study visit	μ g/ml		μ g/ml		μ g/ml			
Screen	43,760	-	570	29	-	-	-	-
Wk -2 (+2d)	47,430	-	834	28	-	-	-	-
Day 0 baseline	23,610	-	682	28	<0.51	<0.46	<0.14	<0.10
Day 0 end 3BNC117	-	-	-	-	591.87	691.29	<0.14	<0.08
Day 0 end 10-1074	-	-	-	-	514.55	760.43	964.42	1,205.59
Day 2 (-1d)	56,930	0.38	-	-	175.43	184.83	1,172.69	705.70
Week 1	3,670	-0.81	-	-	65.12	65.50	467.19	354.60
Week 2	1,140	-1.32	783	26	28.08	34.29	230.89	142.84
Week 2 End 3BNC117	-	-	-	-	605.54	522.54	251.10	69.74
Week 2 End 10-1074	-	-	-	-	464.40	710.17	1,618.91	956.97
Day 16 (-1d)	1,280	-1.27	-	-	206.52	192.74	1,134.66	802.17
Week 3	1,970	-1.08	-	-	79.20	65.73	704.27	351.66
Week 4 (+2d)	20,910	-0.05	655	27	25.90	37.38	272.58	161.28
Week 4 End 3BNC117	-	-	-	-	778.41	662.16	238.07	87.72
Week 4 End 10-1074	-	-	-	-	616.46	908.73	1,608.18	1,164.13
Day 30 (+2d)	27,210	0.06	-	-	158.75	176.71	1,085.61	686.97
Week 5	33,530	0.15	-	-	103.16	129.10	864.85	414.15
Week 6 (+1d)	30,860	0.12	889	34	41.43	92.51	372.33	208.83
Week 7	41,480	0.24	-	-	23.49	44.78	200.72	114.37
Week 8 (+1d)	34,130	0.16	-	-	12.36	15.36	132.56	66.99
Week 10 (+1d)	43,670	0.27	-	-	3.92	2.99	57.63	22.16
Week 12	28,070	0.08	815	29	1.17	1.47	29.98	9.12
Week 16 (-5d)	41,270	0.24	525	31	<0.51	0.64	5.44	2.08
Week 20	38,220	0.21	529	25	<0.51	0.48	1.48	0.42
Week 28	38,300	0.21	707	27	<0.51	<0.28	<0.14	<0.08

9342	HIV-1 RNA		CD4 ⁺ T cells		3BNC117		10-1074	
	cp/ml	$\Delta\log_{10}$	cells/ μ l	%	ELISA	TZM-bl	ELISA	TZM-bl
					Q769.d22	Q769.d22	X2088_c9	Du422.1
Study visit	μ g/ml		μ g/ml		μ g/ml			
Screen	12,600	-	780	37	-	-	-	-
Wk -2 (+6d)	6,510	-	1,150	39	-	-	-	-
Day 0 baseline	2,550	-	780	37	<0.51	<0.46	<0.14	<0.1
Day 0 end 3BNC117	-	-	-	-	906.30	598.56	<0.14	<0.08
Day 0 end 10-1074	-	-	-	-	640.24	1,387.82	1,899.04	991.55
Day 2 (+1d)	3,650	0.16	-	-	150.29	157.32	1,070.88	383.33
Week 1	331	-0.89	-	-	95.67	99.11	709.76	350.58
Week 2	39	-1.82	1,040	39	51.07	143.38	377.23	115.40
Week 2 End 3BNC117	-	-	-	-	822.47	1,187.98	378.85	192.30
Week 2 End 10-1074	-	-	-	-	685.45	1,524.62	1,859.48	648.58
Day 16 (+1d)	61	-1.62	-	-	197.35	248.52	1,357.52	500.25
Week 3	45	-1.75	-	-	109.53	141.15	836.22	364.62
Week 4	25	-2.01	1,140	41	66.94	59.73	555.68	255.74
Week 4 End 3BNC117	-	-	-	-	907.46	612.74	570.14	346.48
Week 4 End 10-1074	-	-	-	-	751.33	1,604.27	2,281.04	3,298.12
Day 30 (+1d)	55	-1.67	-	-	212.77	396.80	1,456.90	1,948.69
Week 5	<20 D	-2.13*	-	-	119.35	145.15	906.38	666.54
Week 6	24	-2.03	870	39	76.63	148.56	454.16	499.36
Week 7	<20 D	-2.13*	-	-	47.02	91.86	393.01	400.05
Week 8	24	-2.03	-	-	32.11	29.37	323.64	285.06
Week 10	32	-1.90	-	-	15.28	15.71	197.13	87.75
Week 12	<20	-2.13*	800	39	7.61	3.25	130.64	56.27
Week 16	55	-1.67	900	44	4.86	<0.462	30.80	6.13
Week 20	645	-0.60	1,066	41	<0.78	<0.28	8.86	1.60
Week 28 (-1d)	3,780	0.17	900	40	<0.51	<0.26	<0.41	<0.08

9343	HIV-1 RNA		CD4 ⁺ T cells		3BNC117		10-1074	
	cp/ml	$\Delta\log_{10}$	cells/ μ l	%	ELISA	TZM-bl	ELISA	TZM-bl
					Q769.d22	Q769.d22	X2088_c9	Du422.1
Study visit	μ g/ml		μ g/ml		μ g/ml			
Screen	95,400	-	410	34	-	-	-	-
Wk -2 (+5d)	91,000	-	380	35	-	-	-	-
Day 0 baseline	97,800	-	350	39	<0.51	<0.46	<0.14	<0.1
Day 0 end 3BNC117	-	-	-	-	708.31	704.45	<0.14	<0.08
Day 0 end 10-1074	-	-	-	-	646.99	1,302.26	1,854.69	1,948.61
Day 2	60,100	-0.21	-	-	140.54	270.87	852.16	705.63
Week 1	1,880	-1.72	-	-	89.38	106.08	546.67	632.45
Week 2 (-2d)	1,200	-1.91	400	40	51.85	61.30	352.90	256.46
Week 2 End 3BNC117	-	-	-	-	723.18	524.50	436.11	168.51
Week 2 End 10-1074	-	-	-	-	644.86	1,457.60	1,639.91	1,511.34
Day 16 (-2d)	808	-2.08	-	-	185.87	398.94	1,271.33	771.96
Week 3 (-3d)	359	-2.44	-	-	118.57	189.64	944.48	878.39
Week 4 (+1d)	335	-2.47	350	39	61.31	70.56	487.90	257.37
Week 4 End 3BNC117	-	-	-	-	898.03	689.28	478.73	285.73
Week 4 End 10-1074	-	-	-	-	556.75	1,976.30	2,051.36	1,486.18
Day 30	373	-2.42	-	-	215.08	786.85	1,398.07	2,797.41
Week 5	265	-2.57	-	-	134.37	97.78	1,008.78	531.42
Week 6	660	-2.17	520	39	112.40	51.56	677.90	235.62
Week 7	3,660	-1.43	-	-	72.81	19.32	467.60	159.35
Week 8 (-1d)	12,000	-0.91	-	-	55.72	23.49	305.69	151.33
Week 10 (+1d)	22	-	-	-	20.52	15.88	223.08	59.75
Week 12	<20D	-	370	40	10.84	4.94	107.82	37.16
Week 16	<20	-	444	39	4.64	1.16	51.12	7.88
Week 20	<20	-	430	38	0.82	<0.26	13.67	2.59
Week 28	<20	-	350	35	<0.51	<0.26	2.36	0.62

* HIV-1 RNA < 20 copies/ml. For calculation of log-change, a value of 19 copies/ml was set.
 < 20D: HIV-1 RNA detected but below limit of quantification.
 Green shaded area indicates HIV-1 RNA nadir.
 Grey shaded area indicates time on ART.

Supplementary Table 3. HIV-1 RNA levels, CD4⁺ T cell counts and bNAbs serum levels (continued)

On ART
Single infusion, day 0
3BNC117 + 10-1074 (10 mg/kg each i.v.)

1A12	HIV-1 RNA	CD4 ⁺ T cells		3BNC117		10-1074	
				ELISA	TZM-bl	ELISA	TZM-bl
				Q769.d22	Q769.d22	X2088_c9	Du422.1
Study visit	cp/ml	cells/μl	%	μg/ml			
Screen	<20	505	42.0	-	-	-	-
Wk -2 (pre-infusion)	<20	556	43.0	-	-	-	-
Day 0 baseline	<20	558	42.9	<0.51	<0.88	0.43*	<0.22
Day 0 end 3BNC117	-	-	-	306.43	253.08	<0.41	<0.27
Day 0 end 10-1074	-	-	-	260.33	192.17	474.87*	281.05
Day 2	<20	-	-	74.25	59.90	323.39*	205.32
Wk 1	<20	-	-	33.12	18.00	191.53*	102.06
Wk 2	<20	691	43.2	20.35	12.33	139.84*	48.48
Wk 3	<20	-	-	14.01	9.02	91.62*	47.17
Wk 4	<20	558	42.9	10.20	5.44	65.02*	21.68
Wk 6	<20	-	-	4.39	3.01	40.93*	17.25
Wk 8	<20	440	44.0	2.38	1.22	18.04*	7.45
Wk 12	<20	534	44.5	0.88	<0.91	6.93*	1.48
Wk 16	<20	566	43.5	<0.51	<0.91	bkg*	<0.25
Wk 20	<20	639	42.6	<0.51	<0.91	bkg*	<0.25
Wk 24	<20	538	44.8	<0.51	<0.91	<0.41	<0.25

1A27	HIV-1 RNA	CD4 ⁺ T cells		3BNC117		10-1074	
				ELISA	TZM-bl	ELISA	TZM-bl
				Q769.d22	Q769.d22	X2088_c9	Du422.1
Study Visit	cp/ml	cells/μl	%	μg/ml			
Screen	<20	780	39.0	-	-	-	-
Wk -2 (pre-infusion)	<20	967	40.3	-	-	-	-
Day 0 baseline	<20	1,033	39.8	1.17*	<0.88	<0.14	<0.22
Day 0 end 3BNC117	-	-	-	197.25*	199.52	<0.14	<0.27
Day 0 end 10-1074	-	-	-	127.91*	84.55	217.43	179.35
Day 2	<20D	-	-	39.51*	29.84	151.20	126.12
Wk 1	<20	-	-	22.15*	14.05	97.38	68.12
Wk 2	<20	971	42.2	15.60*	7.89	76.31	38.78
Wk 3	<20	-	-	8.48*	5.79	46.90	29.10
Wk 4	<20	1,210	43.0	5.54*	2.77	31.86	18.99
Wk 6	<20	-	-	bkg*	<1.11	20.08	11.10
Wk 8	<20	844	40.2	bkg*	<0.91	11.62	2.87
Wk 12	<20	1,350	42.2	bkg*	<0.91	3.82	1.23
Wk 16	30	1,405	43.9	bkg*	<0.91	1.10	0.53
Wk 20	<20	1,592	41.9	bkg*	<0.91	0.54	<0.25
Wk 24	<20	1,229	45.5	<0.51	<0.91	<0.14	<0.25

1A29	HIV-1 RNA	CD4 ⁺ T cells		3BNC117		10-1074	
				ELISA	TZM-bl	ELISA	TZM-bl
				Q769.d22	Q769.d22	X2088_c9	Du422.1
Study Visit	cp/ml	cells/μl	%	μg/ml			
Screen	<20	516	32.4	-	-	-	-
Wk -2 (pre-infusion)	<20	388	35.3	-	-	-	-
Day 0 baseline	<20	648	36.0	<0.51	<0.88	<0.14	<0.22
Day 0 end 3BNC117	-	-	-	228.08	184.64	<0.14	<0.28
Day 0 end 10-1074	-	-	-	187.44	127.30	326.19	246.88
Day 2	<20	-	-	39.82	32.06	140.18	109.65
Wk 1	<20	-	-	26.01	12.10	116.08	59.57
Wk 2	<20	794	36.1	17.91	8.79	93.34	56.12
Wk 3	<20	-	-	12.67	9.52	63.90	47.62
Wk 4	<20	741	34.0	8.55	4.71	41.98	22.84
Wk 6	<20	-	-	5.55	2.51	31.46	15.91
Wk 8	<20	596	33.1	2.71	1.22	18.52	8.87
Wk 12	<20	667	35.1	1.02	<0.91	14.89	4.16
Wk 16	<20	676	35.6	<0.51	<0.91	4.64	1.64
Wk 20	<20	892	34.3	<0.51	<0.91	1.91	0.74
Wk 24	<20D	956	35.4	<0.51	<0.91	0.99	0.39

1A71	HIV-1 RNA	CD4 ⁺ T cells		3BNC117		10-1074	
				ELISA	TZM-bl	ELISA	TZM-bl
				Q769.d22	Q769.d22	X2088_c9	Du422.1
Study Visit	cp/ml	cells/μl	%	μg/ml			
Screen	<20	478	36.8	-	-	-	-
Wk -2 (pre-infusion)	<20	552	34.5	-	-	-	-
Day 0 baseline	<20	358	35.8	<0.51	<0.88	<0.14	<0.22
Day 0 end 3BNC117	-	-	-	169.68	164.67	<0.14	<0.28
Day 0 end 10-1074	-	-	-	117.29	89.49	382.39	201.53
Day 2	<20	-	-	49.08	34.85	270.88	166.39
Wk 1	<20	-	-	24.48	14.14	164.63	56.31
Wk 2	<20	461	38.0	16.12	9.47	138.58	44.15
Wk 3	<20	-	-	12.11	8.02	102.34	30.65
Wk 4	<20	650	43.3	7.11	3.08	62.63	18.28
Wk 6	<20	-	-	5.91	1.95	46.15	15.37
Wk 8	<20	418	38.0	2.81	<0.91	28.89	7.23
Wk 12	<20	467	38.9	1.38	<0.91	10.95	3.24
Wk 16	<20D	458	41.6	<0.51	<0.91	4.58	1.34
Wk 20	<20	458	38.2	<0.51	<0.91	1.89	0.52
Wk 24	<20	429	39.0	<0.51	<0.91	0.66	<0.25

* Number in red indicates background levels detected by ELISA at baseline that are subtracted from all subsequent values.
 bkg indicates time points with a detectable signal within a 3-fold range of the baseline background level.
 < 20D: HIV-1 RNA detected but below limit of quantification.
 Grey shaded area indicates time on ART.

Supplementary Table 3. HIV-1 RNA levels, CD4⁺ T cell counts and bNAb serum levels (continued)

On ART
Single infusion, day 0
3BNC117 + 10-1074 (30 mg/kg each i.v.)

1B05	HIV-1 RNA	CD4 ⁺ T cells		3BNC117		10-1074	
				ELISA	TZM-bl	ELISA	TZM-bl
				Q769.d22		X2088_c9 Du422.1	
Study Visit	cp/ml	cells/μl	%	μg/ml		μg/ml	
Screen	<20	830	35.0	-	-	-	-
Wk -2 (pre-infusion)	<20	843	34.0	-	-	-	-
Day 0 baseline	<20	711	37.4	<0.51	<0.88	<0.14	<0.22
Day 0 end 3BNC117	-	-	-	1,298.73	1,045.69	<0.14	<0.28
Day 0 end 10-1074	-	-	-	481.98	323.18	915.08	952.08 500.20
Day 2	<20	-	-	281.88	208.81	1,486.91	781.03
Wk 1	<20	-	-	147.17	47.28	663.37	396.49
Wk 2	<20	830	34.6	118.27	41.48	557.30	258.98
Wk 3	<20	-	-	61.36	37.63	354.79	147.58
Wk 4	<20	963	34.4	40.15	14.82	210.45	82.94
Wk 6	<20	-	-	21.00	10.58	120.48	60.10
Wk 8	<20	727	34.6	10.28	5.30	72.39	26.64
Wk 12	30	854	35.6	3.01	0.95	29.58	5.72
Wk 16	<20	865	34.6	1.00	<0.91	7.91	1.23
Wk 20	<20D	902	35.0	<0.51	<0.91	2.05	0.30
Wk 24	<20D	966	34.5	<0.51	<0.91	0.62	<0.25

1B44	HIV-1 RNA	CD4 ⁺ T cells		3BNC117		10-1074	
				ELISA	TZM-bl	ELISA	TZM-bl
				Q769.d22		X2088_c9 Du422.1	
Study Visit	cp/ml	cells/μl	%	μg/ml		μg/ml	
Screen	<20D	625	28.0	-	-	-	-
Wk -2 (pre-infusion)	<20D	432	27.0	-	-	-	-
Day 0 baseline	<20	754	31.4	<0.51	<0.88	<0.14	<0.22
Day 0 end 3BNC117	-	-	-	1,062.42	791.00	<0.14	<0.28
Day 0 end 10-1074	-	-	-	837.40	532.67	1,107.92	2,189.62 613.42
Day 2	<20	-	-	228.66	242.26	807.98	914.41
Wk 1	<20D	-	-	100.12	45.24	367.80	412.46
Wk 2	30	653	27.2	44.02	30.24	209.61	150.12
Wk 3	<20	-	-	27.16	12.69	149.04	65.75
Wk 4	<20	737	27.3	13.66	7.03	91.36	48.47
Wk 6	60	-	-	5.87	1.95	42.68	15.71
Wk 8	60	620	29.5	1.97	<0.91	17.87	4.13
Wk 12	20	511	28.4	<0.51	<0.91	4.75	1.09
Wk 16	<20D	637	27.7	<0.51	<0.91	2.07	0.57
Wk 20	<20D	647	30.8	<0.51	<0.91	<0.41	<0.25
Wk 24	<20D	592	29.6	<0.51	<0.91	<0.14	<0.25

1B55	HIV-1 RNA	CD4 ⁺ T cells		3BNC117		10-1074	
				ELISA	TZM-bl	ELISA	TZM-bl
				Q769.d22		X2088_c9 Du422.1	
Study Visit	cp/ml	cells/μl	%	μg/ml		μg/ml	
Screen	<20	575	34.0	-	-	-	-
Wk -2 (pre-infusion)	<20D	763	35.0	-	-	-	-
Day 0 baseline	<20	634	37.3	2.68*	<0.88	<0.14	<0.22
Day 0 end 3BNC117	-	-	-	1282.13*	955.36	<0.14	<0.28
Day 0 end 10-1074	-	-	-	1316.73*	775.93	1,110.10	1,070.39 954.18
Day 2	<20	-	-	199.22*	154.53	688.89	645.71
Wk 1	<20	-	-	140.52*	61.74	488.65	327.15
Wk 2	<20	409	34.1	80.01*	31.14	318.25	179.07
Wk 3	<20	-	-	51.36*	28.72	221.56	142.75
Wk 4	<20	560	35.0	31.38*	10.52	171.94	80.41
Wk 6	<20	-	-	17.93*	7.29	102.91	47.29
Wk 8	<20	680	37.8	10.06*	3.22	57.89	24.78
Wk 12	<20	707	39.3	bkg*	ND	22.65	ND
Wk 16	<20	760	38.0	bkg*	<0.91	8.14	2.15
Wk 20	<20	425	35.4	bkg*	<0.91	3.03	0.79
Wk 24	<20	905	43.1	bkg*	ND	1.25	ND

1B61	HIV-1 RNA	CD4 ⁺ T cells		3BNC117		10-1074	
				ELISA	TZM-bl	ELISA	TZM-bl
				Q769.d22		X2088_c9 Du422.1	
Study Visit	cp/ml	cells/μl	%	μg/ml		μg/ml	
Screen	<20D	449	34.5	-	-	-	-
Wk -2 (pre-infusion)	<20D	619	32.6	-	-	-	-
Day 0 baseline	<20	602	35.4	<0.51	<0.88	<0.14	<0.22
Day 0 end 3BNC117	-	-	-	987.11	754.17	<0.14	<0.28
Day 0 end 10-1074	-	-	-	653.74	707.29	984.32	713.26 456.89
Day 2	<20	-	-	182.35	149.24	706.38	709.00
Wk 1	<20	-	-	96.75	66.41	402.61	336.79
Wk 2	<20	285	28.5	69.99	25.70	284.87	178.69
Wk 3	<20	-	-	46.82	26.44	204.69	115.93
Wk 4	<20	686	36.1	28.47	13.46	156.18	64.06
Wk 6	<20	-	-	20.73	9.13	110.30	47.55
Wk 8	<20	492	32.8	8.08	2.99	70.46	27.80
Wk 12	30	679	37.7	1.27	<0.91	34.42	12.78
Wk 16	<20D	566	35.4	<0.51	<0.91	16.65	3.97
Wk 20	<20	515	34.3	<0.51	<0.91	6.20	1.27
Wk 24	<20	675	35.5	<0.51	<0.91	2.52	0.93

* Number in red indicates background levels detected by ELISA at baseline that are subtracted from all subsequent values.

bkg indicates time points with a detectable signal within a 3-fold range of the baseline background level.

< 20D: HIV-1 RNA detected but below limit of quantification.

Grey shaded area indicates time on ART.

ND: Not determined because of detectable activity (ID₅₀ > 20) against MuLV negative control.

Supplementary Table 4. Adverse events reported during study follow-up

Adverse Event (AE)	AEs (no.)	Severity Grade			Participants of n=15 (no.)
		Mild (no.)	Moderate (no.)	Severe (no.)	
Related to 3BNC117 + 10-1074					
Nausea	1	1	0	0	1
Transaminitis	1	1	0	0	1
Upper respiratory tract infection	1	1	0	0	1
Total related	3	3	0	0	3
Not related to 3BNC117 + 10-1074					
Gonorrhea	2	0	2	0	2
Upper respiratory tract infection	2	1	0	0	2
Abdominal distension	1	1	0	0	1
Agitation	1	1	0	0	1
Altered mental status*	1	0	0	1	1
Common cold	1	1	0	0	1
Headache	1	1	0	0	1
Hypoglycemia	1	1	0	0	1
Increase in direct bilirubin**	1	n/a	n/a	2	1
Mycoplasma urethritis	1	0	1	0	1
Sore throat	1	1	0	0	1
Vomiting	1	0	1	0	1
Worsening of tinea pedis	1	0	1	0	1
Total Not related	15	7	5	3	6
Total	18	10	5	3	9

* One participant experienced altered mental status due to drug intoxication with gamma-hydroxybutyric acid (GHB).

** One participant noted to have transient elevation to 0.3 mg/dL one week after first mAb infusions, and 0.4 mg/dL 24 weeks after last mAb infusions. These were graded as severe according to the DAIDS Toxicity table (version 2.0)

Table indicates adverse events observed during the study observation period (24 weeks after the last antibody infusion) in the 15 enrolled participants. Adverse events were graded according to the Division of AIDS (DAIDS) Table for Grading the Severity of Adult and Pediatric Adverse Events (version 2.0).

Supplementary Table 5. Pharmacokinetics of 3BNC117 and 10-1074

a Summary

Antibody	ART	Dose	Regimen	Method (n)	$t_{1/2}$ (days)*			AUC _{last} ($\mu\text{g} \times \text{h/ml}$)			C _{max} ($\mu\text{g/ml}$)		
					Mean	SD	Range	Mean	SD	Range	Mean	SD	Range
3BNC117	Off	30 mg/kg	wk 0	ELISA (4)	10.1	3.9	6.0-15.1	3,155.5	667.7	2,612.5-4,039.2	817.9	122.5	735.9-996.2
				TZM-bl (4)	7.9	3.0	6.3-12.5	5,022.7	1,650.6	3,183.3-6,643.2	1,284.1	234.7	945.1-1,443.4
	Off	30 mg/kg	wk 0-2-4	ELISA (3)	12.3	4.1	8.1-16.2	7,226.3	1,833.3	5,331.7-8,991.5	861.3	71.9	778.4-907.5
				TZM-bl (2)**	9.6	2.5	7.8-11.4	10,603.8	5,705.9	6,568.4-14,637.8	1,256.6	491.8	908.7-1,604.3
	All viremic	30 mg/kg		ELISA (7)	11.1	3.8	6.0-16.2						
				TZM-bl (6)**	8.5	2.8	6.3-12.5						
	On	10 mg/kg	wk0	ELISA (4)	16.4	4.3	10.2-19.8	923.2	262.6	579.0-1,197.2	225.4	59.1	169.7-306.4
TZM-bl (4)				12.1	2.1	9.4-14.4	562.0	167.9	387.2-775.0	200.5	37.9	164.7-253.1	
On	30 mg/kg	wk0	ELISA (4)	12.5	3.3	9.4-16.7	3,971.3	969.9	2,817.1-4,878.8	1,166.2	166.4	987.1-1,316.7	
			TZM-bl (4)	10.9	2.1	7.7-12.0	2,353.6	318.5	1976.2-2754.5	886.6	137.5	754.2-1,045.7	
All ART-treated			ELISA (8)	14.5	4.1	9.4-19.8							
			TZM-bl (8)	11.5	2.1	7.7-14.4							
10-1074	Off	30 mg/kg	wk 0	ELISA (4)	11.9	3.5	7.9-15.1	12,924.1	2,507.2	10,009.6-16,136.0	1,249.0	278.6	889.1-1,561.2
				TZM-bl (4)	11.5	1.6	9.5-13.3	10,880.4	2,459.1	9,260.1-14,514.9	2,359.4	1,873.9	1,154.5-5,123.3
	Off	30 mg/kg	wk 0-2-4	ELISA (3)	12.7	1.5	11.7-14.4	44,769.0	9,803.1	35,626.4-55,120.4	1,983.8	336.2	1,618.9-2,281.0
				TZM-bl (3)	11.5	0.5	11.0-12.1	31,505.9	9,067.2	21,143.8-37,984.6	2,433.7	1,092.7	1,205.6-3,298.1
	All viremic	30 mg/kg		ELISA (7)	12.2	2.7	7.9-15.1						
				TZM-bl (7)	11.5	1.2	9.5-13.3						
	On	10 mg/kg	wk0	ELISA (4)	19.9	3.4	16.0-23.2	5,083.5	1,513.2	3,153.8-6,485.4	350.2	107.7	217.4-474.9
TZM-bl (4)				20.0	5.5	11.9-23.9	2,469.8	423.3	1,934.7-2,916.2	227.2	45.6	179.4-281.1	
On	30 mg/kg	wk0	ELISA (4)	18.1	2.3	15.4-20.5	16,519.6	5,125.0	10,951.7-23,366.6	1,172.3	217.8	984.3-1,486.9	
			TZM-bl (4)	16.8	2.8	13.9-20.6	10,418.1	706.7	9,751.7-11,377.6	1,231.3	655.9	713.3-2,189.6	
All ART-treated			ELISA (8)	19.0	2.8	15.4-23.2							
			TZM-bl (8)	18.4	4.4	11.9-23.9							

* Calculated terminal elimination half-lives.

** One participant not included because of adjusted $R^2 < 0.8$.

AUC - Area under the curve, SD - standard deviation.

Table indicates pharmacokinetic parameters for the administered antibodies derived from serum concentrations determined by ELISA and TZM-bl assay. (a) Summary of results grouped by antibody and treatment status. (b) and (c) Results for 3BNC117 and 10-1074, respectively, obtained in individual study participants.

Supplementary Table 5. Pharmacokinetics of 3BNC117 and 10-1074 (continued)

b 3BNC117

ID	ART	Dose	Regimen	Method	C _{max} (µg/ml)	Adjusted R ²	Estimated t _{1/2} (days)	λ _z lower (days)	λ _z upper (days)	AUC (INF_pred)	t _{last} (days)	C _{last} (µg/ml)	AUC_% Extrap_pred
91C22	Off	30 mg/kg	wk 0	ELISA	996.2	0.999	6.0	21	42	2,623.7	42	1.3	0.4
					1,443.4	0.988	6.3	14	42	3,196.2	42	1.4	0.4
91C35	Off	30 mg/kg	wk 0	ELISA	739.4	0.998	11.1	22	58	4,078.9	58	2.6	1.0
					1,440.5	0.904	12.5	7	84	6,660.3	84	1.6	0.3
91C33	Off	30 mg/kg	wk 0	ELISA	800.3	1.000	8.2	31	56	2,691.2	56	2.1	0.9
					1,307.4	0.999	6.3	21	56	4,109.7	56	1.4	0.3
91C34	Off	30 mg/kg	wk 0	ELISA	735.9	0.996	15.1	28	87	3,337.3	87	1.6	1.0
					945.1	0.857	6.7	28	56	6,226.7	56	5.0	1.0
9341	Off	30 mg/kg	wk 0-2-4	ELISA	778.4	0.999	8.1	43	84	5,345.9	84	1.2	0.3
					908.7	0.864	11.4	32	140	6,570.7	140	0.5	0.0
9342	Off	30 mg/kg	wk 0-2-4	ELISA	907.5	0.933	16.2	35	112	9,068.4	112	4.9	0.8
					1,604.3	0.967	7.8	42	84	14,677.4	84	3.2	0.3
9343	Off	30 mg/kg	wk 0-2-4	ELISA	898.0	0.983	12.8	42	55	8,362.9	55	55.7	12.0
					1,976.3	0.779	4.2	29	55	11,893.3	55	23.5	0.5
1A12	On	10 mg/kg	wk 0	ELISA	306.4	0.994	18.3	42	84	1,220.1	84	0.9	1.9
					253.1	0.992	12.8	7	56	798.6	56	1.2	3.0
1A27	On	10 mg/kg	wk 0	ELISA	197.3	0.986	10.2	7	28	657.6	28	5.5	11.9
					199.5	0.973	9.4	7	28	425.9	28	2.8	9.1
1A29	On	10 mg/kg	wk 0	ELISA	228.1	0.996	17.4	21	84	1,056.6	84	1.0	2.3
					184.6	1.000	14.4	29	57	631.1	57	1.2	4.0
1A71	On	10 mg/kg	wk 0	ELISA	169.7	0.973	19.8	14	84	920.6	84	1.4	3.9
					164.7	0.937	12.0	7	43	511.2	43	1.9	6.1
1B05	On	30 mg/kg	wk 0	ELISA	1,298.7	0.998	16.7	56	112	4,682.2	112	1.0	0.5
					1,045.7	0.995	12.0	42	84	2,381.6	84	1.0	0.7
1B44	On	30 mg/kg	wk 0	ELISA	1,062.4	0.995	9.4	15	57	2,844.0	57	2.0	0.9
					791.0	0.999	7.7	22	43	1,998.1	43	1.9	1.1
1B55	On	30 mg/kg	wk 0	ELISA	1,316.7	0.978	13.5	14	43	5,208.4	43	17.9	6.3
					955.4	0.957	11.9	7	56	2,807.3	56	3.2	1.9
1B61	On	30 mg/kg	wk 0	ELISA	987.1	1.000	10.4	42	84	3,549.8	84	1.3	0.5
					754.2	0.952	11.8	21	56	2,374.9	56	3.0	2.3

Supplementary Table 5. Pharmacokinetics of 3BNC117 and 10-1074 (continued)

c 10-1074

ID	ART	Dose	Regimen	Method	C _{max} (µg/ml)	Adjusted R ²	Estimated t _{1/2} (days)	λ _z lower (days)	λ _z upper (days)	AUC (INF_pred)	t _{last} (days)	C _{last} (µg/ml)	AUC_% Extrap_pred
91C22	Off	30 mg/kg	wk 0	ELISA	1,561.2	0.983	9.8	14	80	10,029.1	80	1.5	0.2
					5,123.3	0.996	9.5	28	80	14,523.8	80	0.7	0.1
91C35	Off	30 mg/kg	wk 0	ELISA	1,224.3	1.000	15.1	42	84	12,900.2	84	7.4	1.3
					1,154.5	0.991	11.2	7	84	9,295.1	84	2.6	0.4
91C33	Off	30 mg/kg	wk 0	ELISA	1,321.4	0.996	14.6	21	142	12,824.0	142	0.6	0.1
					1,917.7	1.000	13.3	56	111	9,512.7	111	0.6	0.1
91C34	Off	30 mg/kg	wk 0	ELISA	889.1	0.961	7.9	87	140	16,141.1	140	0.5	0.0
					1,242.1	0.996	12.1	56	112	10,266.9	112	1.2	0.2
9341	Off	30 mg/kg	wk 0-2-4	ELISA	1,618.9	0.988	12.1	43	140	35,646.9	140	1.5	0.1
					1,205.6	0.990	12.1	71	140	21,150.4	140	0.4	0.0
9342	Off	30 mg/kg	wk 0-2-4	ELISA	2,281.0	0.996	14.4	84	140	55,298.8	140	8.9	0.3
					3,298.1	0.991	11.3	42	140	38,008.2	140	1.6	0.1
9343	Off	30 mg/kg	wk 0-2-4	ELISA	2,051.4	0.997	11.7	30	55	48,842.1	55	305.7	10.8
					2,797.4	0.817	11.0	35	55	37,411.5	55	151.3	5.4
1A12	On	10 mg/kg	wk 0	ELISA	474.9	0.990	16.0	7	84	6,213.0	84	6.9	2.3
					281.1	1.000	11.9	42	84	2,941.4	84	1.5	0.9
1A27	On	10 mg/kg	wk 0	ELISA	217.4	0.995	18.2	20	140	3,166.0	140	0.5	0.4
					179.4	1.000	23.0	56	112	1,952.3	112	0.5	0.9
1A29	On	10 mg/kg	wk 0	ELISA	326.2	1.000	22.0	112	161	4,657.4	161	1.0	0.7
					246.9	1.000	23.9	112	161	2,682.7	161	0.4	0.5
1A71	On	10 mg/kg	wk 0	ELISA	382.4	1.000	23.2	111	176	6,507.3	176	0.7	0.3
					201.5	1.000	21.2	84	140	2,375.3	140	0.5	0.7
1B05	On	30 mg/kg	wk 0	ELISA	1,486.9	0.998	17.2	112	175	23,382.3	175	0.6	0.1
					952.1	0.995	13.9	21	147	11,382.8	147	0.3	0.0
1B44	On	30 mg/kg	wk 0	ELISA	1,107.9	0.997	15.4	57	105	10,996.5	105	2.1	0.4
					2,189.6	0.979	16.6	57	105	10,071.2	105	0.6	0.1
1B55	On	30 mg/kg	wk 0	ELISA	1,110.1	0.999	19.4	21	168	16,131.3	168	1.2	0.2
					1,070.4	0.999	16.2	29	139	10,502.1	139	0.8	0.2
1B61	On	30 mg/kg	wk 0	ELISA	984.3	0.999	20.5	112	168	15,734.5	168	2.5	0.5
					713.3	0.986	20.6	21	168	9,772.0	168	0.9	0.2

Supplementary Table 6. Antibody sensitivity of plasma SGA-derived pseudoviruses

Response	ID	Seq ID	Time point	3BNC117		10-1074		PGDM1400		PGT121		µg/ml	
				IC50	IC80	IC50	IC80	IC50	IC80	IC50	IC80		
91C22	91C22 A2 91C22 A5 91C22 A6 91C22 A10	Day 0	0.003	0.013	0.003	0.008	0.089	0.572	0.002	0.007			
			0.010	0.033	0.004	0.010	0.123	0.721	0.004	0.014			
			0.002	0.006	0.003	0.008	0.104	0.638	0.002	0.007			
			0.005	0.013	0.003	0.008	0.212	0.833	0.003	0.008			
	91C22 A11 91C22 B1 91C22 B3 91C22 B4	Week 14	0.006	0.016	0.050	0.179	0.070	0.324	0.027	0.103			
			0.008	0.022	0.058	0.189	0.096	0.326	0.028	0.078			
			0.006	0.018	0.003	0.008	0.159	1.055	0.003	0.016			
			0.006	0.016	0.002	0.008	0.118	0.946	0.003	0.009			
	91C22 B7	Week 16	0.005	0.014	0.003	0.007	0.155	0.933	0.004	0.013			
	91C34	91C34 day 0_1 91C34 day 0_2 91C34 day0_4 91C34 day0_10	Day 0	0.020	0.091	0.017	0.081	> 50	> 50	0.089	0.417		
				0.009	0.029	0.025	0.084	> 50	> 50	0.070	0.321		
				0.016	0.056	0.026	0.113	> 50	> 50	0.091	0.305		
0.007				0.029	0.006	0.027	NT	NT	NT	NT			
91C34 W6_4 91C34 w6_6		Week 6	0.009	0.031	0.062	0.403	> 50	> 50	0.137	0.923			
			0.011	0.036	0.073	0.427	> 50	> 50	0.123	0.529			
91C34 w12_1 91C34 w12_2 91C34 w12_4 91C34 w12_5 91C34 w12_7 91C34 w12_8		Week 12	0.014	0.047	0.094	0.640	> 50	> 50	1.210	> 50			
			0.035	0.113	0.075	0.323	> 50	> 50	0.045	0.151			
			0.056	0.153	> 50	> 50	> 50	> 50	> 50	> 50			
			0.060	0.170	> 50	> 50	> 50	> 50	> 50	> 50			
	0.009		0.065	0.017	0.189	NT	NT	NT	NT				
	0.008		0.026	0.021	0.074	NT	NT	NT	NT				
9342	9342 B1 9342 B2 9342 B3 9342 B5 9342 B6 9342 B7 9342 B8	Day 0	0.015	0.052	0.016	0.054	3.254	10.811	0.010	0.036			
			0.013	0.039	0.030	0.084	6.452	18.102	0.022	0.083			
			0.033	0.101	0.028	0.085	0.837	2.801	0.025	0.107			
			0.052	0.181	0.019	0.055	0.981	5.296	0.009	0.030			
	9342 A1 9342 A2 9342 A3 9342 A6 9342 A8	Week 20	0.010	0.031	0.016	0.037	1.904	9.166	0.010	0.031			
			0.027	0.099	> 50	> 50	1.173	5.389	> 50	> 50			
			0.042	0.116	0.057	0.192	1.473	4.911	0.023	0.108			
			0.062	0.216	> 50	> 50	3.942	16.337	> 50	> 50			
9343	9343 A1 9343 A3 9343 A5 9343 A6 9343 A8	Day 0	0.016	0.045	0.014	0.038	> 50	> 50	0.010	0.035			
			0.005	0.018	0.005	0.018	> 50	> 50	0.003	0.009			
			0.013	0.031	0.013	0.031	> 50	> 50	0.005	0.017			
			0.011	0.033	0.011	0.033	> 50	> 50	0.013	0.100			
	9343 B6 9343 B9	Week 7	0.030	0.218	> 50	> 50	NT	NT	NT	NT			
			0.034	0.179	> 50	> 50	> 50	> 50	> 50	> 50			
	9343 B1 9343 B2 9343 B4 9343 B5	Week 8	0.094	0.428	> 50	> 50	> 50	> 50	> 50	> 50			
			0.053	0.178	> 50	> 50	NT	NT	NT	NT			
			0.043	0.143	> 50	> 50	NT	NT	NT	NT			
			0.049	0.188	> 50	> 50	NT	NT	NT	NT			

NT: Not tested.

Table indicates sensitivity of pseudoviruses against 3BNC117, 10-1074, PGDM1400 and PGT121. Pseudoviruses were constructed using *env* sequences obtained by SGA on plasma samples of the indicated time points. Sensitivity was determined in the TZM-bl neutralization assay. Participants are grouped based on the response to the 3BNC117 and 10-1074 infusions. Seq ID refers to virus sequence ID shown in **Fig. 3** and **Supplementary Fig. 6**.

Supplementary Table 6. Antibody sensitivity of plasma SGA-derived pseudoviruses (continued)

	ID	Seq ID	Time point	3BNC117		10-1074		PGDM1400		PGT121		µg/ml
				IC50	IC80	IC50	IC80	IC50	IC80	IC50	IC80	
Short-term response	91C35	91C35 A1	Day 0	0.386	1.721	<0.001	0.003	0.040	0.204	<0.001	0.001	< 0.01 0.01 - 0.1 0.1 - 0.5 0.5 - 5 5 - 50 > 50
		91C35 A2		0.304	1.490	0.001	0.003	0.048	0.626	<0.001	0.002	
		91C35 A3		0.125	0.442	0.002	0.006	1.814	> 50	<0.001	0.005	
		91C35 A4		0.234	1.233	<0.001	0.002	0.170	1.214	<0.001	0.001	
		91C35 A5		0.255	0.861	<0.001	0.004	0.044	0.208	<0.001	0.003	
		91C35 A7		0.421	1.851	0.002	0.004	0.257	16.039	0.001	0.003	
		91C35 A10		0.073	0.235	0.002	0.008	1.484	38.424	0.001	0.006	
		91C35 A12		0.516	2.335	0.001	0.005	0.090	0.527	<0.001	0.003	
		91C35 B1		0.348	1.436	0.002	0.007	0.028	0.092	0.001	0.004	
		91C35 B2		0.339	2.152	> 50	> 50	0.063	0.560	30.773	> 50	
	91C35 B3	0.219	1.419	> 50	> 50	0.072	0.694	> 50	> 50			
	91C35 B5	0.295	1.918	> 50	> 50	0.057	0.438	2.507	> 50			
	91C35 B6	41.958	> 50	> 50	> 50	0.073	0.509	11.548	> 50			
	9341	9341 A5	Day 0	0.035	0.104	0.149	0.424	0.602	3.104	0.151	0.568	
		9341 A12		0.097	0.350	0.205	0.721	0.567	2.867	0.240	1.130	
		9341 B2A		0.057	0.160	0.156	0.710	0.417	2.121	0.344	1.643	
9341 B4		0.007		0.019	0.067	0.292	0.073	0.347	0.038	0.165		
9341 B5		0.040		0.113	0.132	0.475	0.519	2.482	0.109	0.505		
9341 B2		Week 4	42.305	> 50	> 50	> 50	0.173	0.888	> 50	> 50		
9341 B7		Week 5	0.040	0.213	> 50	> 50	NT	NT	NT	NT		
9341 B6	0.046		0.358	> 50	> 50	0.216	1.636	> 50	> 50			
9341 B5	0.060		0.298	> 50	> 50	0.271	2.157	> 50	> 50			
No response	91C33	91C33 A3	Day 0	0.027	0.075	0.078	0.273	4.433	43.039	0.181	0.891	
		91C33 A4		0.023	0.110	0.053	0.203	40.565	> 50	0.089	0.517	
		91C33 A5		0.029	0.082	0.077	0.219	4.210	46.952	0.193	0.939	
		91C33 A6		> 50	> 50	> 50	> 50	> 50	> 50	> 50	> 50	
		91C33 A9		> 50	> 50	> 50	> 50	NT	NT	NT	NT	

NT: Not tested.