

Supplementary Information for: Rational Design and *in vivo* Characterization of mRNA-encoded Broadly Neutralizing Antibody Combinations Against HIV-1

Table S1. Neutralization antibody concentration inhibiting 80% of viral infection.

	IgG	scFv-Fc.r1	scFv-Fc.r2
PGDM1400			
Mean (SD)	3.75 (5.4)	21.05 (53.67)	31.76 (90.05)
Median (IQR)	1.17 (0.13, 6.75)	3.05 (0.38, 7.72)	1.64 (0.32, 12.56)
10-1074			
Mean (SD)	1.13 (2.46)	2.59 (7.45)	1.05 (2.14)
Median (IQR)	0.25 (0.07, 1.10)	0.14 (0.04, 0.45)	0.24 (0.07, 0.91)
PGT121			
Mean (SD)	3.67 (8.2)	2.72 (7.21)	2.54 (7.16)
Median (IQR)	0.13 (0.04, 0.56)	0.16 (0.07, 0.45)	0.12 (0.05, 0.42)
3BNC117			
Mean (SD)	3.23 (7.43)	13.15 (11.08)	3.58 (3.37)
Median (IQR)	0.55 (0.17, 1.61)	11.09 (1.02, 25.00)	2.21 (0.71, 5.97)
N6			
Mean (SD)	0.43 (0.51)		2.73 (3.00)
Median (IQR)	0.20 (0.05, 0.85)		1.33 (0.45, 5.24)
NIH45-46			
Mean (SD)	0.38 (0.41)		2.73 (3.00)
Median (IQR)	0.24 (0.06, 0.75)		1.33 (0.45, 5.24)

IQR, interquartile range (25% percentile, 75% percentile); SD, standard deviation.

Table S2. scFv-Sc Round 2 Variant Mutations.

scFv-Sc Round 2 Variants	Mutations
PGDM1400	
Var_7	Linker20 (GGGGS) 4
Var_8	L20 + H:K23R
Var_9	L20 + H:K23R + H:K94F
Var_10	L20 + H:K19R
Var_11	L20 + H:K19R + S82AN
Var_12	L20 + S39Y
Var_13	VH_VL + L20
10-1074	
Var_7	Linker20 (GGGGS)4
Var_8	L20 + H:S79Y
Var_9	L20 + H:S19R
Var_10	L20 + H:K81Q
Var_11	L20 + H:S18R + H:S79Y + H:K81Q
Var_12	L20 + H:S18R + H:S79Y + H:K81Q + H:V68T
Var_13	L20 + H:L80Y
Var_14	L20 + H:S23Q
Var_15	L20 + H:S40R
Var_16	L20 + H:K13R
Var_17	L20 + H:K13R + S23Q
Var_18	VH_VL + L20

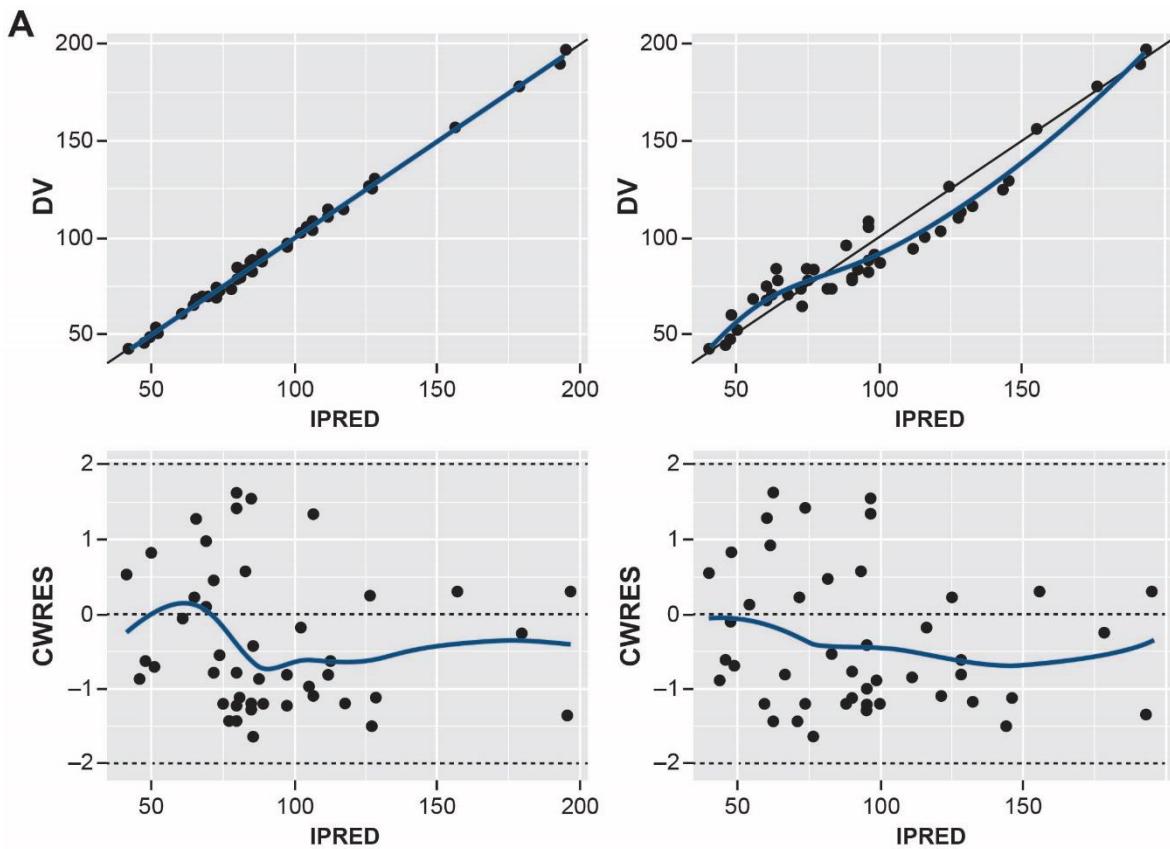
PGT121	
Var_7	Linker20 (GGGGS)4
Var_7a	L20 + 68N_Rev
Var_8	L20 + 68N_Rev + H:S79Y
Var_9	L20 + 68N_Rev + H:S19R
Var_10	L20 + 68N_Rev + H:S81Q
Var_11	L20 + 68N_Rev + H:S19R + H:S79Y + H:S81Q
Var_12	L20 + H:S19R + H:S79Y + H:S81Q
Var_13	L20 + 68N_Rev + H:S81I
Var_14	L20 + 68N_Rev + H:S23Q
Var_15	L20 + 68N_Rev + L:S63M
Var_16	VH_VL + L20
N6	
Var_2	Linker20 (GGGGS)4
Var_3	L20 + H:A10Y
Var_4	L20 + H:K12R
Var_5	L20 + H:D81Q
Var_6	L20 + H:D81Q + H:R82AY

Var, variant.

Table S3. Kinetic–Pharmacodynamic Model Parameters.

Parameter (unit)	Estimate		
	N6 IgG	PGT121 scFv-Fc Var7B	PGDM1400 scFv-Fc Var7
m	0.862	0.566	0.438
CL _{ab} (mL/h)	0.00111	0.000719	0.000507
t _{1/2} (days)	26.0	40.2	57.0

CL_{ab}, the antibody clearance rate from the central compartment; Fc, crystallizable fragment; IgG, immunoglobulin G; M, linear slope parameter; scFv, single-chain variable fragment; t_{1/2}, half-life.



DV, Dependent variable; IPRED, Individual fit; PRED, Population Fit; CWRES, Conditional weighted residuals

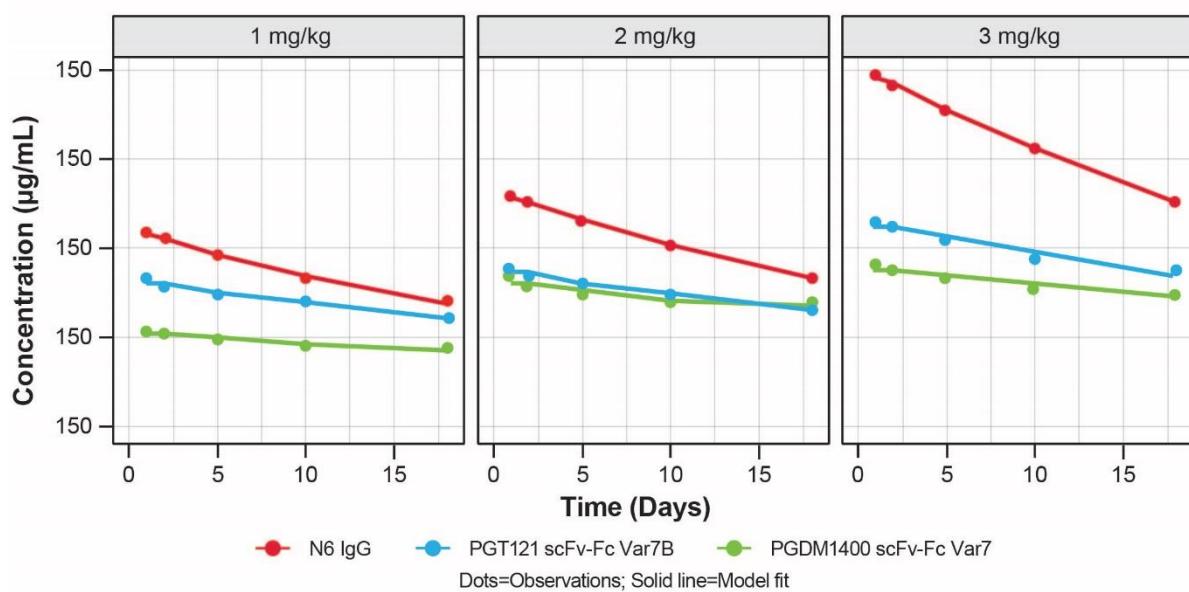


Figure S1. Kinetic–pharmacodynamic model diagnostics. (A) Goodness-of-fit plots and (B) fits for individual groups.