

Supplemental data

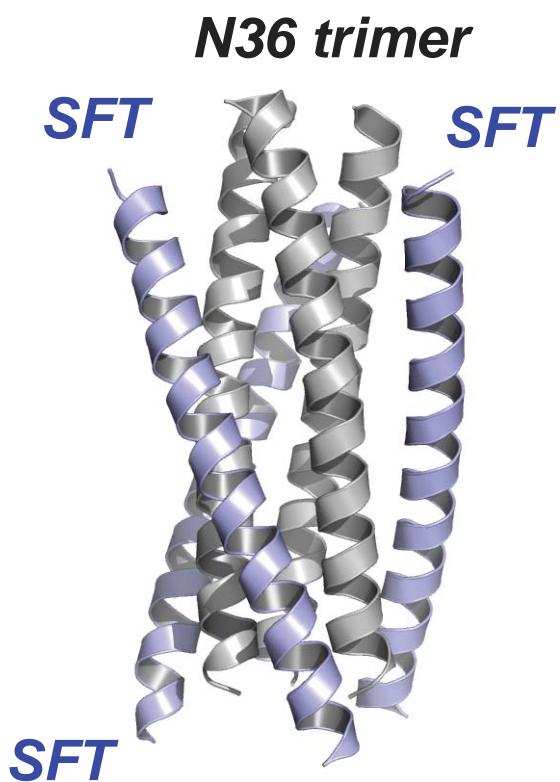
Figure S1. The 6-HB structure formed by SFT and N36 peptides. Ribbon model of 6-HB form by SFT and N36 (HIV gp41 NHR₃₅₋₇₀) in different views with the labels. (A) Side view of the 6-HB; (B) Top view of the 6-HB. The N36 peptides are colored in gray; the SFT peptides are colored in light blue.

Figure S2. Phylogenetic analysis of 43 HIV-1 Env sequences derived from the subtypes A, B and C that dominate the AIDS epidemics worldwide, and subtype B', CRF07_BC and CRF01_AE recombinants that are currently circulating in China.

Table S1. Sequence alignment of the NHR and CHR of HIV-1 gp41. Both NHR and CHR sequences show high variability.

Fig. S1

A



B

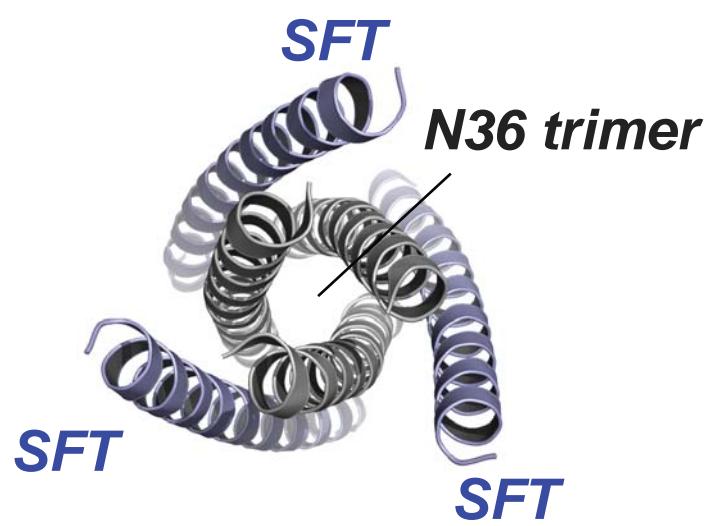
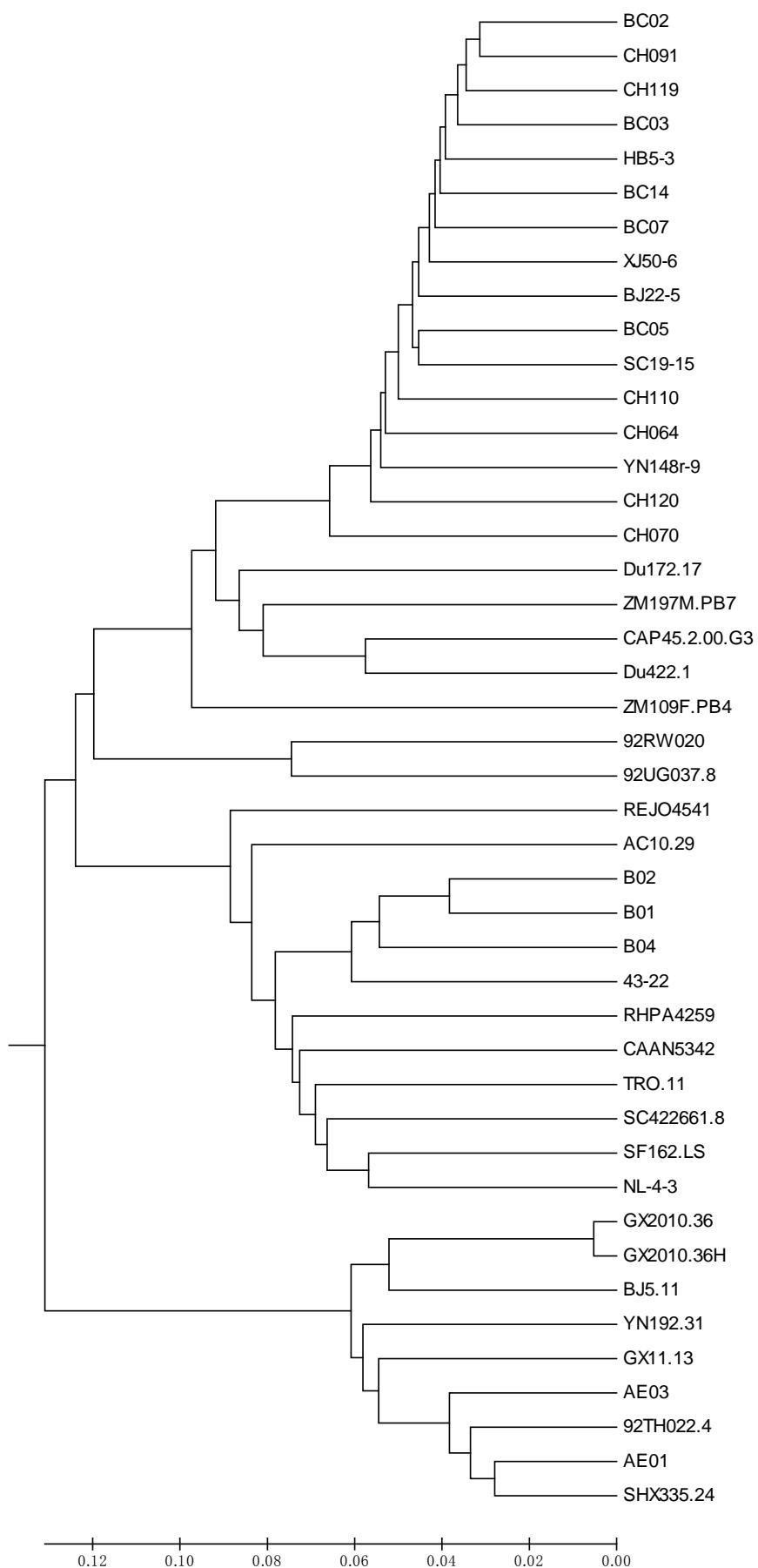


Fig. S2



Supplemental Table S1.

Table S1. Alignment of the NHR and CHR sequences of HIV-1 gp41

Pseudovirus	Accession no.	Clade	NHR sequence	CHR sequence
NL4-3	AF324493_8	B	QARQLLSDIV QQQNLLRAI EAQQHLLQLT VVGIKQLQAR ILAVERYLKD QQLL	WMEWDREINN YTSLIHSILIE ESQNQQEKNE QELLELDKWA SLWNWF
SF162.LS	EU123924	BG.E..D. .N..YT.....
AC10. 0.29	AY835446	B	...L..G.Q....S.YT.....
TR0. 11	AY835445	B	...L..G.M....	...E..D. ..D.YI.L. K. I.....
REJ04541	AY835449	BG. ...S....	.R.EK..D. ..D.YT.. K.....
RHPA4259	AY835447	B	...L..G. ...S....	..Q.E..D. ..D.YN.L.
CAAN5342	AY835452	B	...L..G.EK..D. ..GI.YN.L.
SC422661.8	AY835441	B	...L..G.V. I.....
Du172.17	DQ411853	CG. ...S....M....	...V. I.....
Du422.1	DQ411854.1	CG. ...S....T. V. I.....
ZM197M.PB7	DQ388515.1	CG. ...S....M....	...T. V. I.....
ZM109P.PB4	AY424138.2	CG. ...S....V. I....Q.
CAP45.2.00.G3	DQ435682.1	CG. ...S....M....	...V. I.....
92RW020	AY669706.1	AG. ...S....M.K.V.
92UG037.8	U09127.1	A	.V.K...G. ...S....K.V.R.
92TH022.4	AY669778.1	EG. ...S....X..M....	...V.KF.
CH064	EF117254	B/C	H.....G. ...S....T. V. I.....
CH070	EF117255	B/CG.K.V. I.....
CH091	EF117256	B/CG. ...S....T. V. I.....
CH110	EF117257	B/CG. ...S....T. V. I.....
CH119	EF117261	B/CG. ...S....T. V. I.....
CH120	EF117260	B/CG. ...S..K.T. V. I.....
SC19-15	ADT65148	B/CG. ...S....T. V.
BJ22-5	ADT65144	B/CG.M....	...T. V. I.....

BC07	ABY71530	B/CG.S.....H.....T. V..I.....EQ.E...S. ...DT.YR.L. D.....R. KD..A..S.N N....
BC14	ABY71537	B/CG.S..K.T. V..I.....	...Q..K.... .DI.YR.L. ...S...R. KD..A..S.K N..S..
YN148r-9	ADT65158	B/CG.S.....V..M.....	...Q..K..S. ...NT.YR.L. D.....R. KD..A..S.Q N....
XJ50-6	ADT65154	B/CG.S.....T. V..I.....	...Q.EE..S. ...T.YR.L. KD..A..S.K N....
HB5-3	ADT65143	B/CG.S.....T. V..I....Q.Q..K..S. ...YT.YR.L. KD..A..S.K N....
BC02	ABY71525	B/CG.S.....T. V..I.....	...Q..K..S. ...NT.YR.L. D.....R. KD..A..S.K N....
BC03	ABY71526	B/CG.S.....M....T. V..I....R.Q..K..S. ...KT.YQ.L. D..Y...R. KD..A..S.K N..S..
BC05	ABY71528	B/CG.S.....T. V..M.....	...Q....SK ..NT.YG.L. D.....R. KD..A..S.K N..T..
B02	ABY71519	B'G.R.....V.....	...E..D. ..RE.YT. L.....
43-22	CHH	B'G.R.....V.....	...E..A. ..KE.YT. . Q. L..... S..
B01	ABY71518	B'M.G.R.....V.....	...Q.E.....RE.YA. L.....
B04	ABY71521	B'G.R.....V.....	...Q.E..D. ..KE.YT. LD..K..E.
SHX335.24	ADD83190	A/EG.S.....M....V.....T..	.I..EK..S. ...Q.YKILT ...D..DR. KD.....
YN192.31	GU475046	A/EG.S.....V.....Q. .KF.	.T..E..S. ...NK.YG.LTDR. KD.....
AE03	ABY71544	A/EG.S.....V.....V.....KF.	.I..E..S. ...NQ.YEILTD.. KD.....T N....
GX2010.36	ADD83183	A/EG.S.....K.....V.....N. .KF.	.T..E..S. ...DT.YQ.LTD.. KD..... N....
GX11.13	GU475042	A/EFG.S.....V.....V.....KF.	.T..E..S. ...NQ.YEILTDR. KD.....
GX2010.36H	GU475026.1	A/EG.S.....K.....V.....N. .KF.	.T..E..S. ...NT.YQ.LTD.. KD.....
AE01	ABY71542	A/EG.S.....V.....KF.	.I..E..S. ...Q.YEILTDR. KD.....
BJ5.11	ADD83181	A/EG.S.....M....V.....KF.	.I..EK..S. ...NK.YD.LTR. KD.....