Enhancing Virion Tethering of HIV by BST2 Sensitizes Productively and Latently Infected T cells to ADCC Mediated by Broadly Neutralizing Antibodies

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SUPPLEMENTARY INFORMATION

Supplementary Figure Legends

Fig. S1. Neutralization potency and ADCC potential of anti-HIV antibodies. CEM CD4⁺ T cells were infected with prototypic virus CCR5-tropic NL4.3.ADA.IRES.GFP WT strain and at 2 dpi infected cells were examined for (a) cell-surface Env expression and (b) susceptibility to ADCC using different concentrations of NAbs (in µg/ml). (c) Neutralizing antibodies were evaluated for their ability to neutralize cell-free virus. Extent of Env staining on GFP⁺ T cells over a range of concentrations was expressed as averaged median fluorescence intensity (MFI) units from 3 to 4 experiments. PBMCs from healthy donors were used as effector cells. Percentages of Ab-specific ADCC were determined following subtraction of cell lysis mediated by control IgG from that by test antibody. Mean ADCC activity from at least 4-7 donors is indicated. Serial dilutions of known concentrations of NAbs (or as control, PBS) were pre-incubated with prototypic CCR5-tropic NL4.3.ADA.IRES.GFP WT virus for 45 min at 37 °C. Luciferase-expressing HeLa TZMbl reporter cells were subsequently added to Ab-virus complexes. Forty-eight hours later, infected cells were lysed and relative light units enumerated on a luminometer. Data were analyzed using GraphPad Prism. Shown are mean IC₅₀ from 2 to 3 experiments; ND, not done.

Fig. S2. Cell-surface envelope, CD4 and BST2 expression profiles on HIV-infected CEM CD4⁺ T cells. Parental CEM CD4⁺ T cells were infected or mock-infected with CCR5-tropic NL4.3.ADA.IRES.GFP WT virus or derivatives lacking Vpu (U-), Nef (N-) or both (N-U-). The N-U-D368A virus has a mutation within the CD4-binding site of Env causing defective CD4-Env interactions. At 2 dpi, infected cells were analyzed by flow cytometry for (**a**) Env using indicated NAbs and (**b**) CD4 using anti-CD4 Ab. (**c**) Alternatively, CEM CD4⁺ T cells expressing (NT) or depleted (SH) of BST2 were infected with the indicated viruses as in Panel a and analyzed for cell-surface BST2 using anti-BST2 Ab or pre-immune serum (PI) as control. Shown in Panels a and c are expression levels on GFP⁺ cells presented as MFI units from a representative experiment. Panel b indicates mean CD4 expression from 3 experiments.

Fig. S3: Cell-surface envelope and BST2 expression profiles on HIV-infected

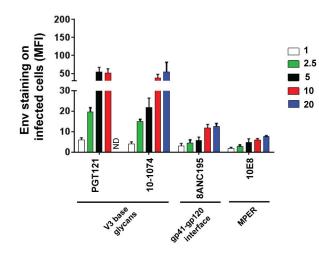
primary CD4⁺ T cells. Activated CD4⁺ T cells were infected or mock-infected with CCR5-tropic NL4.3.ADA.IRES.GFP WT virus or derivatives lacking Vpu (U-), Nef (N-) or both (N-U-). The N-U-D368A virus has a mutation within the CD4-binding site of Env causing defective CD4-Env interactions. At 3 dpi, infected GFP⁺ cells were analyzed by flow cytometry for (**a**) Env using indicated NAbs or (**b**) BST2 using anti-BST2 Ab. Shown in both Panels a and b are mean expression levels on GFP⁺ cells presented as MFI units from 3 different donors.

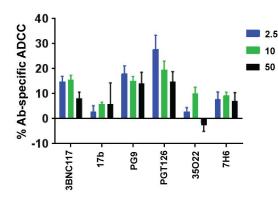
Fig. S4: Reactivated latent T cell lines are targets of ADCC and Vpu dampens

PGT121-mediated ADCC response. Jurkat-based T cells latently infected with CCR5tropic NL4.3.ADA.IRES.GFP WT or its mutant derivatives lacking Vpu (U-), Nef (N-) or both (N-U-) viruses were treated with TNF α and examined for their susceptibility to ADCC mediated by PGT121. PBMCs from healthy donors were used as effector cells. (a) Examples of ADCC response from 2 donors. (b) Compilation of PGT121-specific

ADCC. Boxes and whiskers graph shows median ADCC from 4 donors.

b

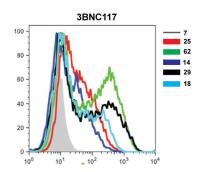


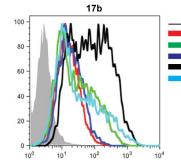


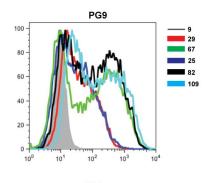
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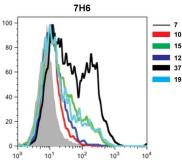
Antibody	Epitope (Env region)	IC ₅₀ ± S.D. (µg/ml)
3BNC117	CD4-bs (gp120)	0.07 ± 0.03
17b	CoR-bs (gp120)	1.30 ± 2.08
PG9	V1/V2 apex (gp120)	0.07 ± 0.02
PGT121	V3 base glycans (gp120)	0.04 ± 0.02
PGT126	V3 base glycans (gp120)	0.04 ± 0.01
35022	gp41-gp120 interface	17.78 ± 9.50
7H6	MPER (gp41)	0.26 ± 0.10

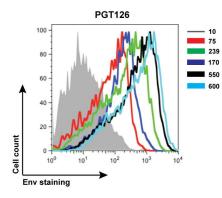


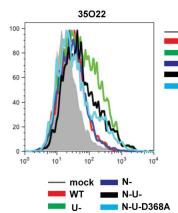


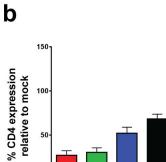












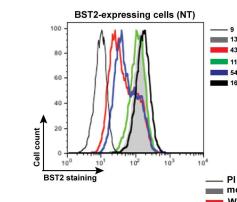
WT



N-U- N-U-D368A

N-

U-



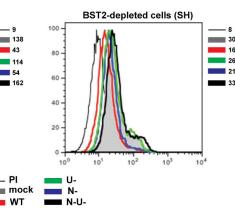


Fig. S2

