

Arginine Insertion and Loss of N-linked Glycosylation Site in HIV-1 Envelope V3 Region Confer CXCR4-tropism

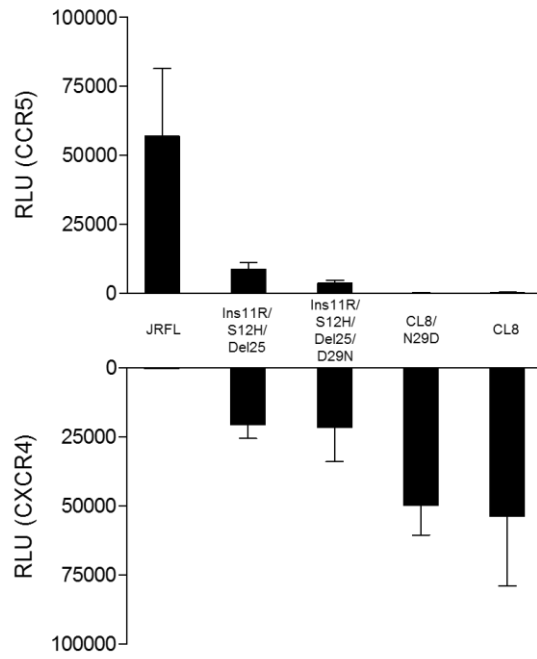
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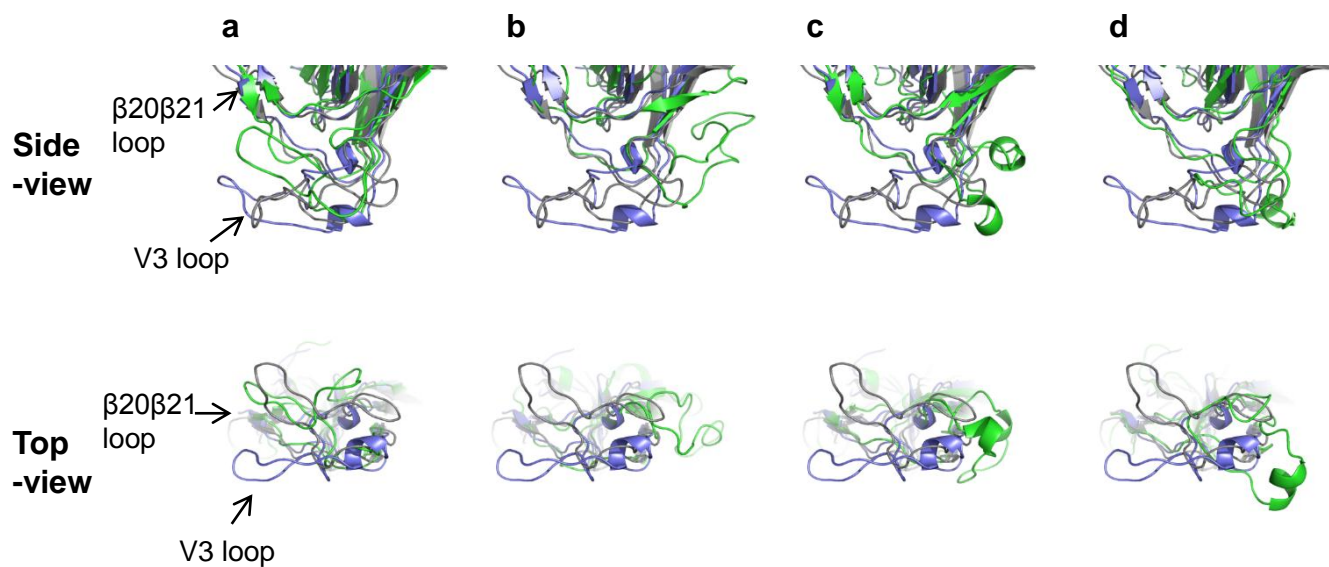
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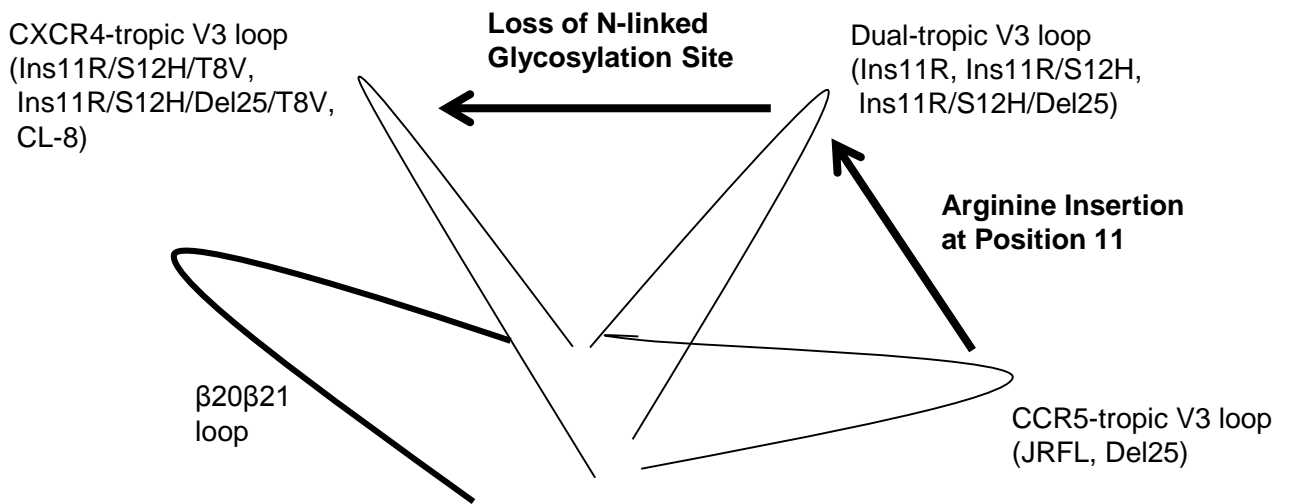
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Supplementary Figure S1. Effects of D29N in cell-cell fusion assay. Cell-cell fusion assay was performed using Env-expressing 293T cells and CD4⁺ and CCR5⁺/CXCR4⁺ COS-7 cells. Data are mean \pm SD values in relative luminescent unit (RLU) of six experiments (performed in duplicate and repeated three times).



Supplementary Figure S2. Structural models of V3 loops of Ins11R/S12H/Del25-derived mutants. MD simulations were performed for the HIV-1 Ins11R/S12H/Del25 gp120 outer domain with N5Y (a), K10I (b), Y22H (c), or V26G (d) substitution in V3 loop. Structures were constructed and superimposed as described in Fig. 3 caption. (a) Superimposition of N5Y (green), Ins11R/S12H/Del25 (gray), and CL8 (navy). (b) Superimposition of K10I (green), Ins11R/S12H/Del25 (gray), and CL8 (navy). (c) Superimposition of Y22H (green), Ins11R/S12H/Del25 (gray), and CL8 (navy). (d) Superimposition of V26G (green), Ins11R/S12H/Del25 (gray), and CL8 (navy).



Supplementary Figure S3. Schematic summary of V3 loop orientation and tropism. CCR5-tropic V3 loop was placed in the opposite direction from the β 20- β 21 loop. Arginine insertion at position 11 changed the V3 loop orientation and conferred dual-tropism. Loss of N-linked glycosylation site further changed the V3 loop orientation into the same direction with the β 20- β 21 loop and altered it CXCR4-tropic.