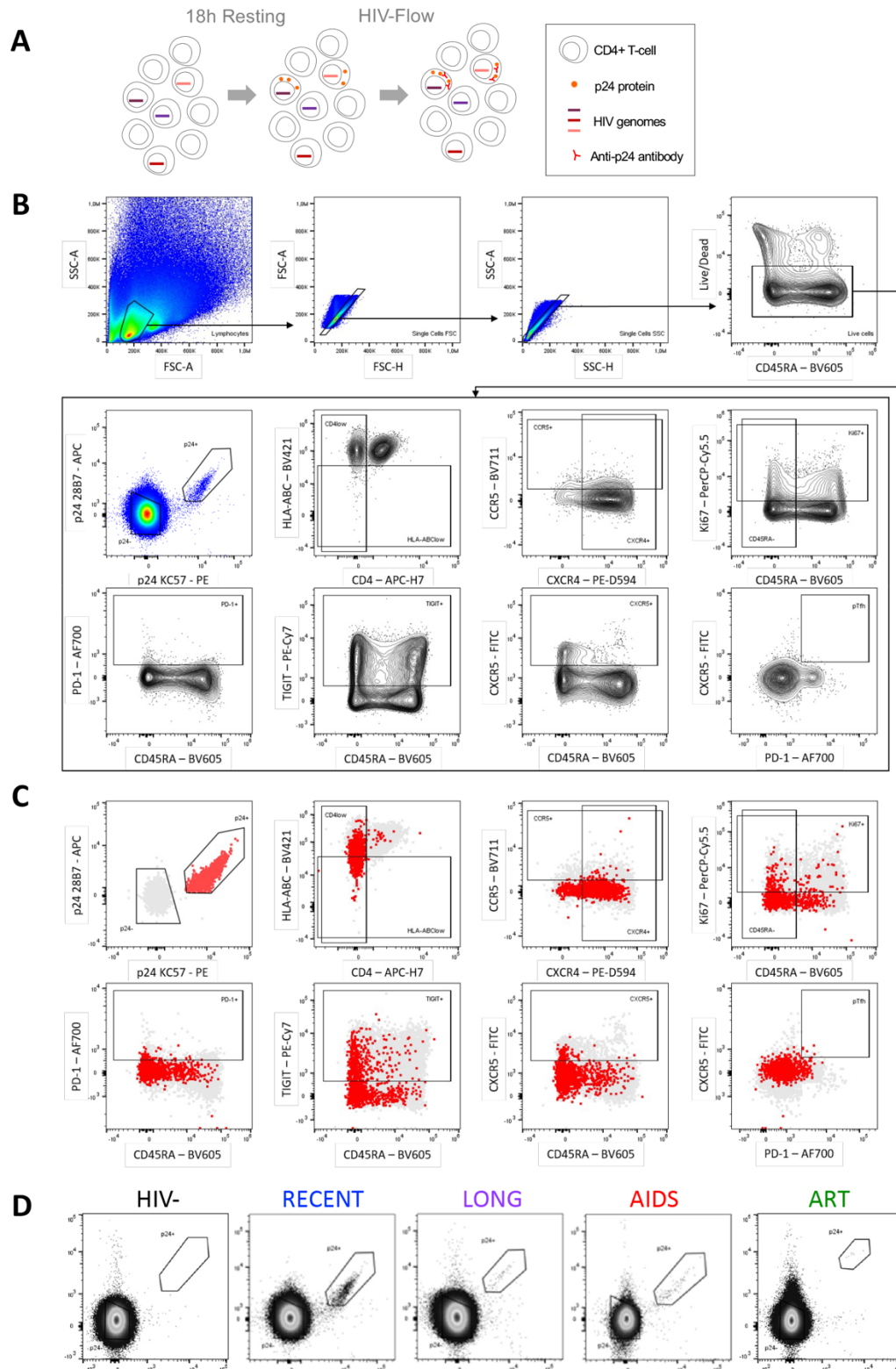
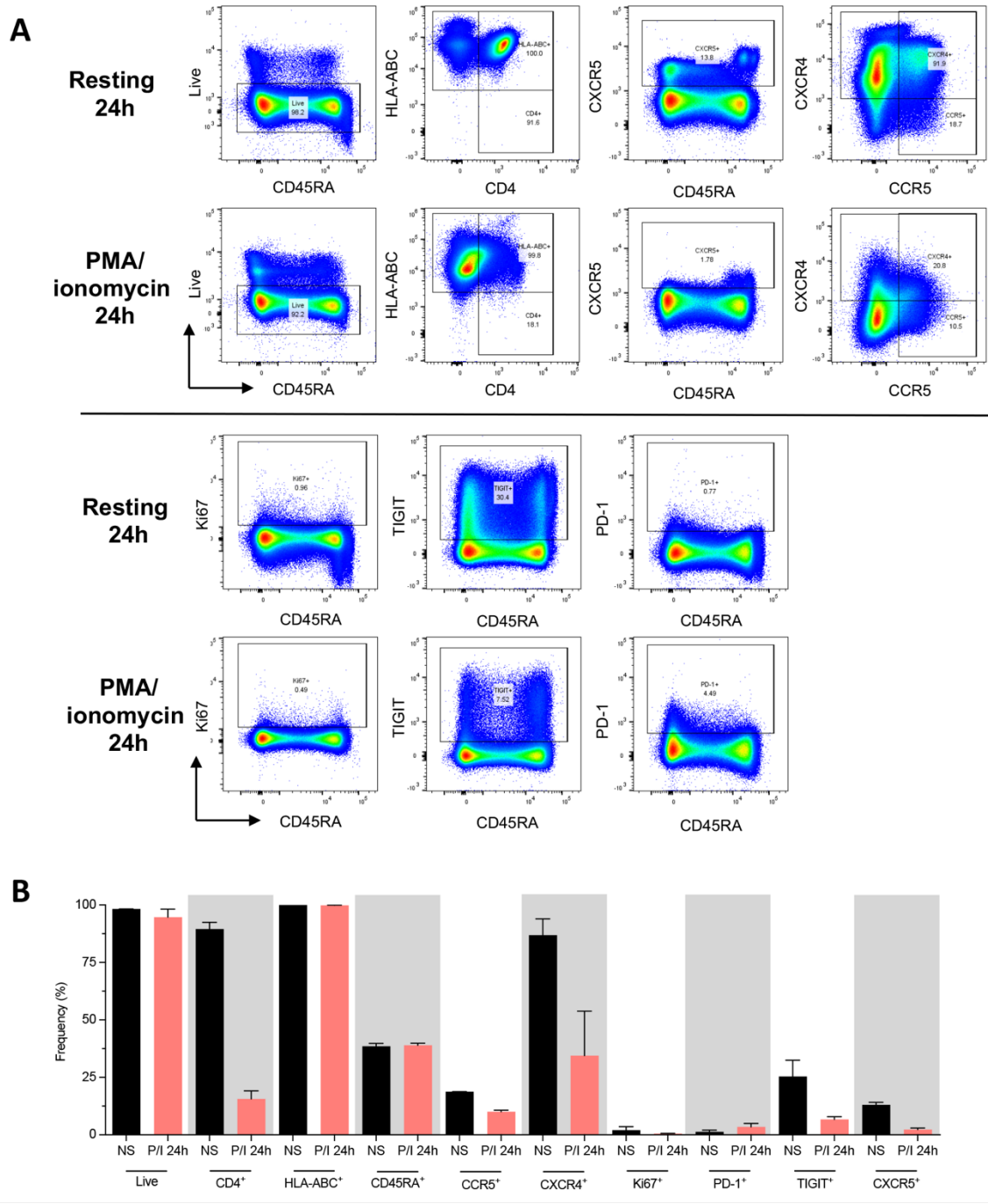


Supplemental Figure 1



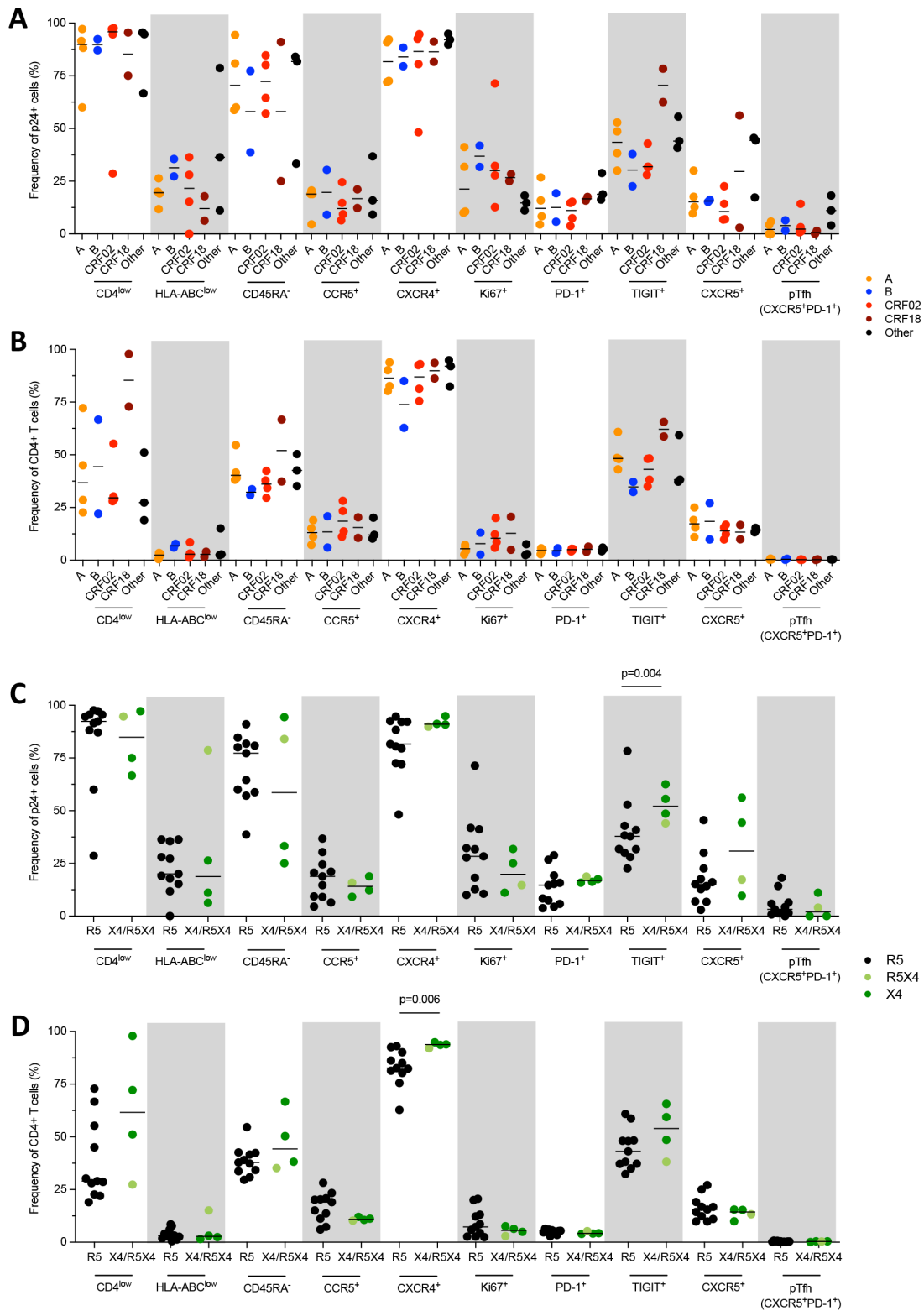
Supplementary Figure 1. Gating strategy for single cell sorting. A. Isolated CD4⁺ T cells were rested for 18 h and their p24 expression was analyzed by HIV-Flow. B. Gating strategy used for p24⁺ cell analysis with recording of CD4, HLA-ABC, CD45RA, CCR5, CXCR4, Ki67, PD-1, TIGIT and CXCR5 expression. C. Representative dot plots showing expression of markers described in B. in p24⁺ cells (in red), overlaid onto p24⁻ cells (in grey). D. Representative dot plots showing p24⁺ gating for one participant of each group (HIV⁻, recent infection, long-term infection, AIDS and ART-treated participants).

Supplemental Figure 2



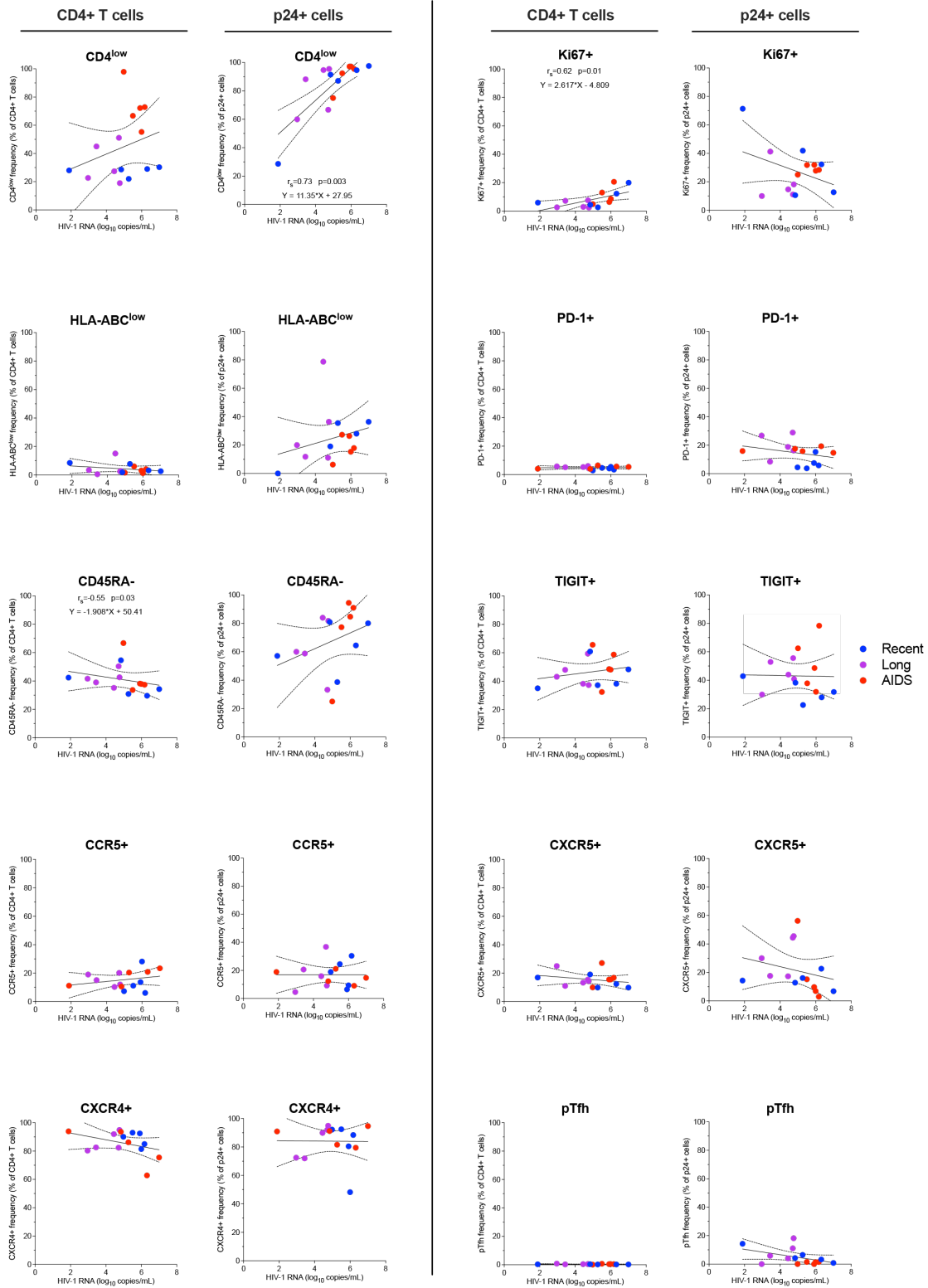
Supplementary Figure 2. Phenotype changes upon stimulation with PMA/ionomycin. A. Representative dot plot from one HIV- individual for each marker of interest in the absence or in presence of stimulation with BFA + PMA/ionomycin. **B.** Differential phenotypic expression for each marker. CD4, CCR5, CXCR4, TIGIT and CXCR5 were downregulated.

Supplemental Figure 3



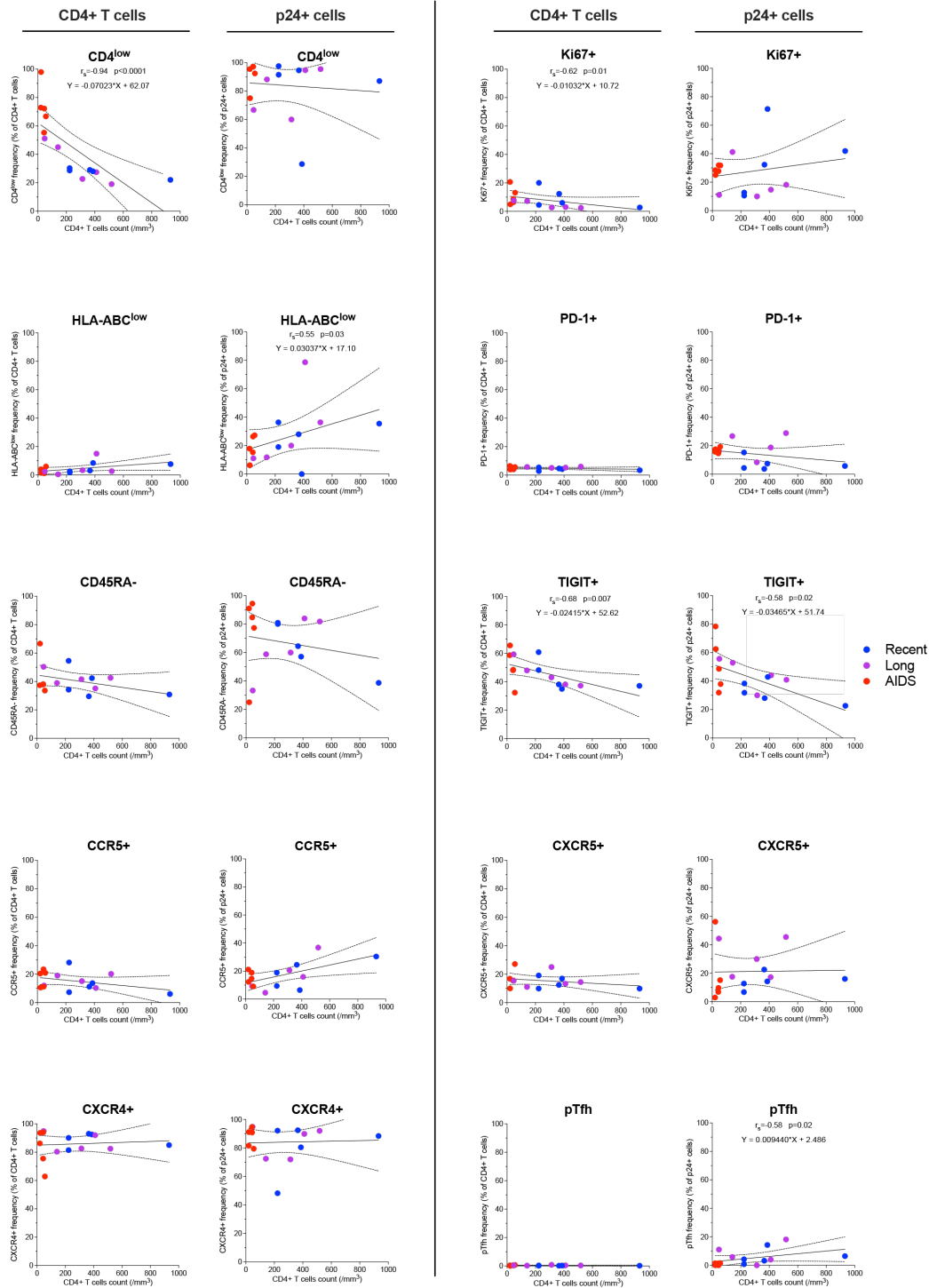
Supplementary Figure 3. Impact of HIV-1 subtype diversity and tropism on the phenotype of p24+ cells and CD4+ T cells. A-B. Participants were grouped according to their HIV-1 subtype (A, n=4; B, n=2; CRF02, n=4; CRF18, n=2, other, n=3) and the phenotype was displayed for p24+ cells (A.) and CD4+ T cells (B.). No statistical testing was possible due to limited numbers of participants per group. C-D. Participants were grouped according to the tropism of their HIV-1 strains (R5, n=11; R5/X4, n=1; X4, n=3) and the phenotype was displayed for p24+ cells (C.) and CD4+ T cells (D.). Significant changes are highlighted with the p-value (Mann-Whitney).

Supplemental Figure 4



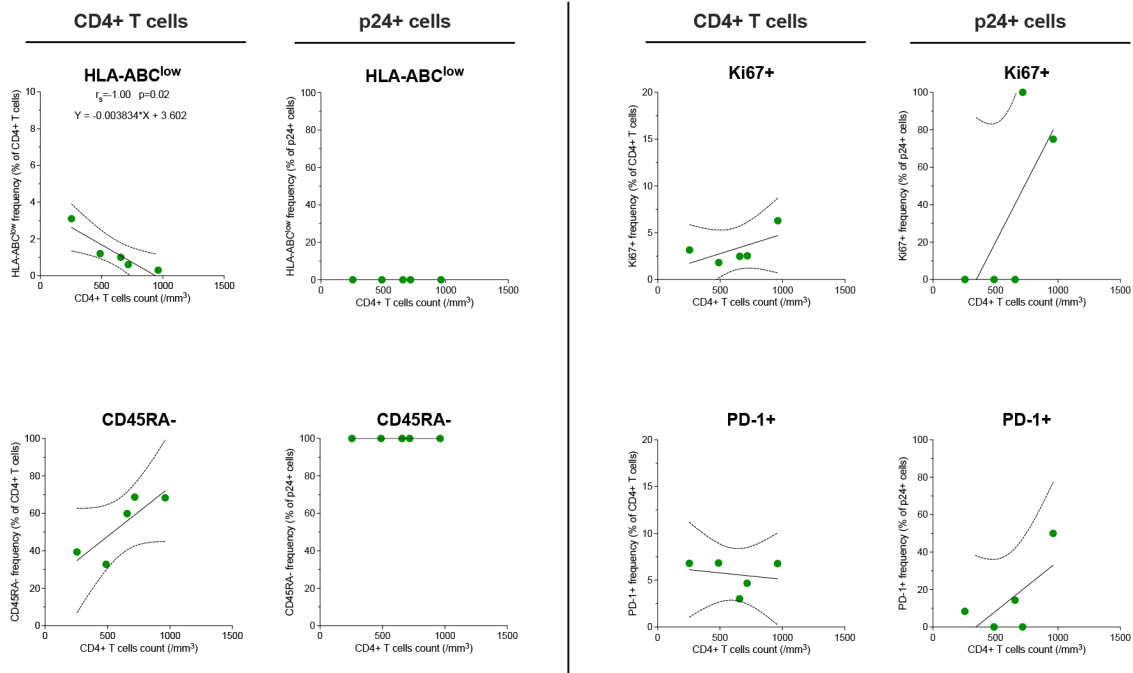
Supplementary Figure 4. Correlation between markers expression and plasma viral load for viremic participants. Correlations between plasma viremia and the frequency of total CD4⁺ T cells (left) and p24⁺ cells (right) expressing or not one the following markers: CD4^{low}, HLA-ABC^{low}, CD45RA⁻, CCR5⁺, CXCR4⁺, Ki67⁺, PD-1⁺, TIGIT⁺, CXCR5⁺, pTfh cells (CXCR5⁺PD-1⁺). The regression line was plotted with 95%CI. Significant associations are highlighted with the regression equation and correlation values (Spearman).

Supplemental Figure 5



Supplementary Figure 5. Correlation between markers expression and CD4+ T cells count for viremic participants. Correlations between CD4+ T cells count and the frequency of total CD4+ T cells (left) and p24+ cells (right) expressing or not one the following markers: CD4low, HLA-ABClow, CD45RA-, CCR5+, CXCR4+, Ki67+, PD-1+, TIGIT+, CXCR5+, pTfh cells (CXCR5+PD-1+). The regression line was plotted with 95%CI. Significant associations are highlighted with the regression equation and correlation values (Spearman).

Supplemental Figure 6



Supplemental Figure 6. Correlation between markers expression and CD4+ T cells count for ART-treated participants. Correlations between CD4+ T cells count and the frequency of total CD4+ T cells (left) and p24+ cells (right) expressing or not one the following markers: HLA-ABC^{low}, CD45RA-, Ki67+, PD-1+. The regression line was plotted with 95%CI. Significant associations are highlighted with the regression equation and correlation values (Spearman).