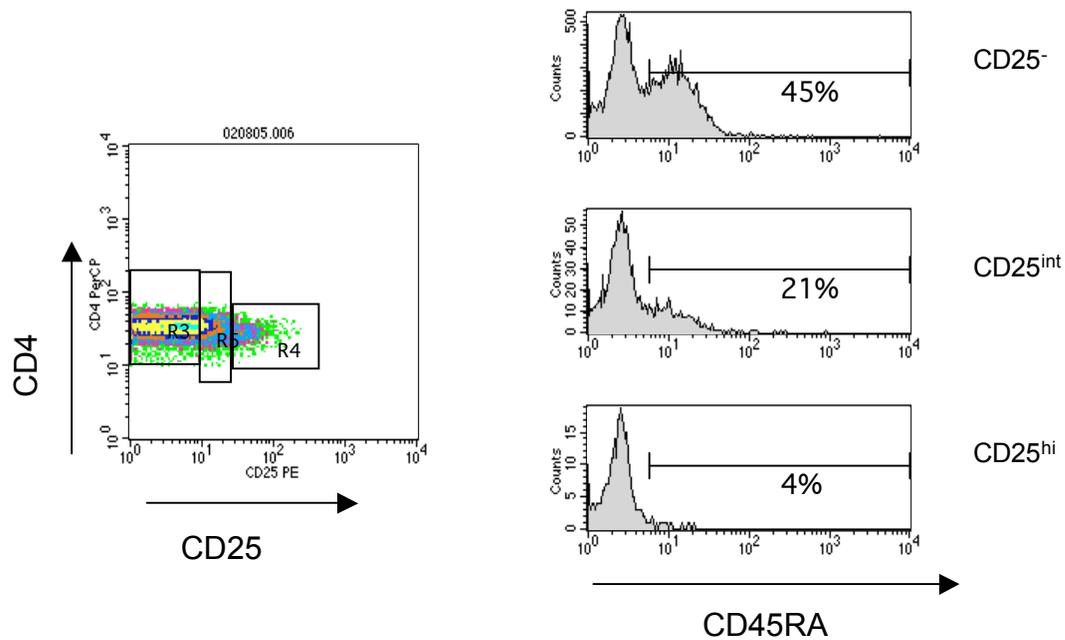
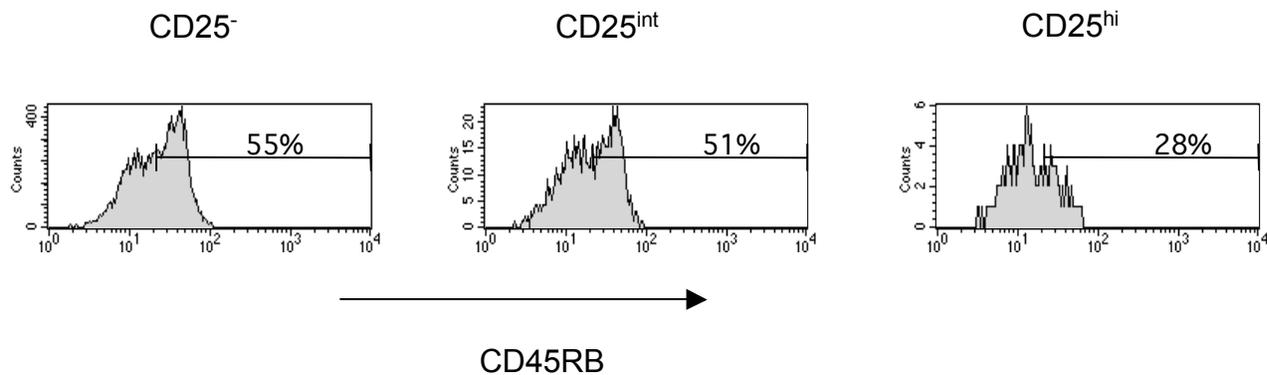


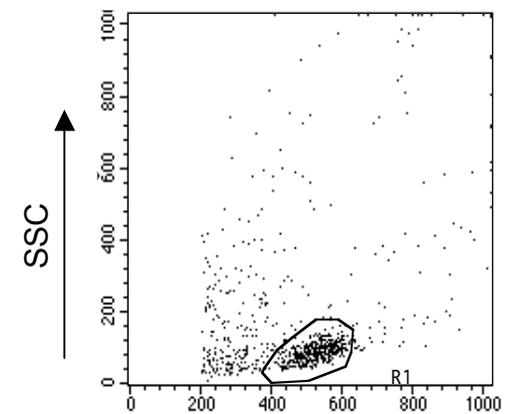
A



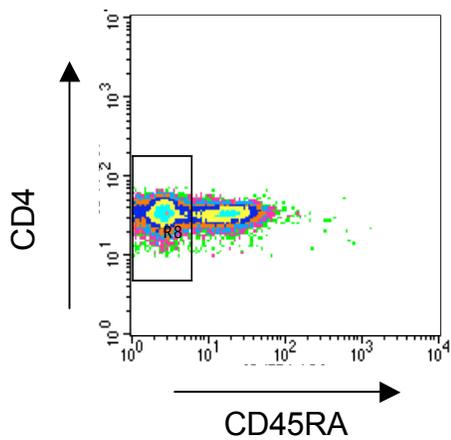
B



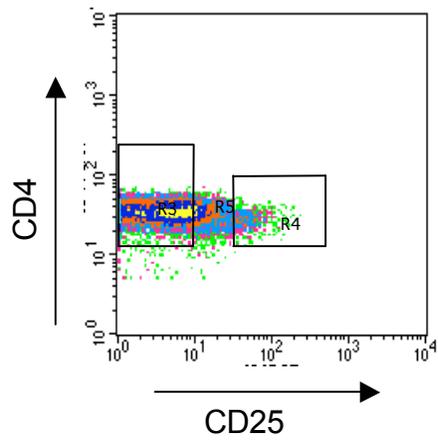
Supplemental figure 2



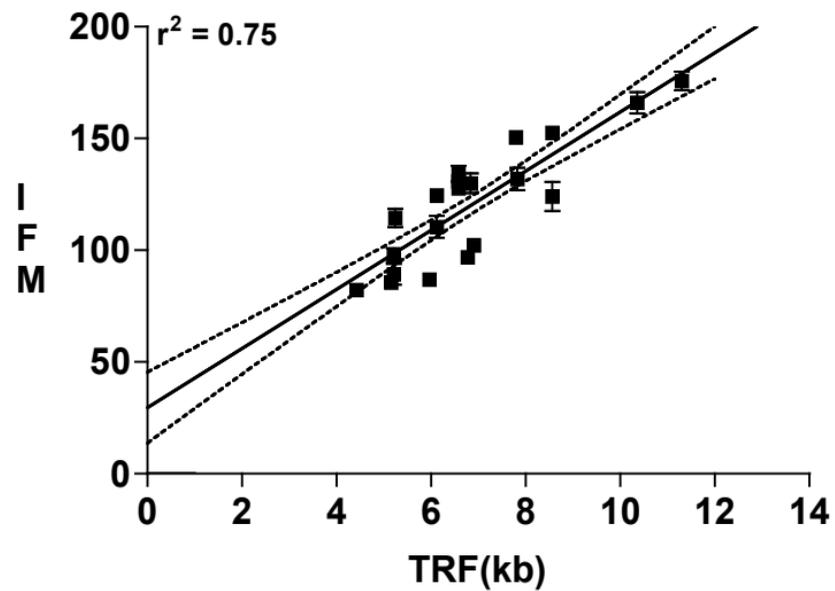
FSC



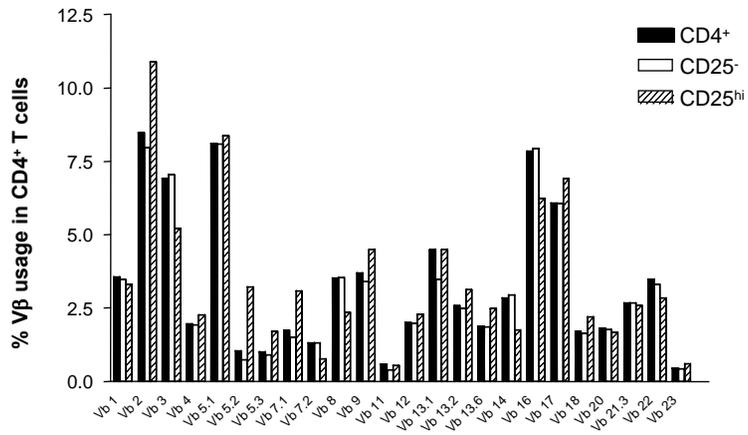
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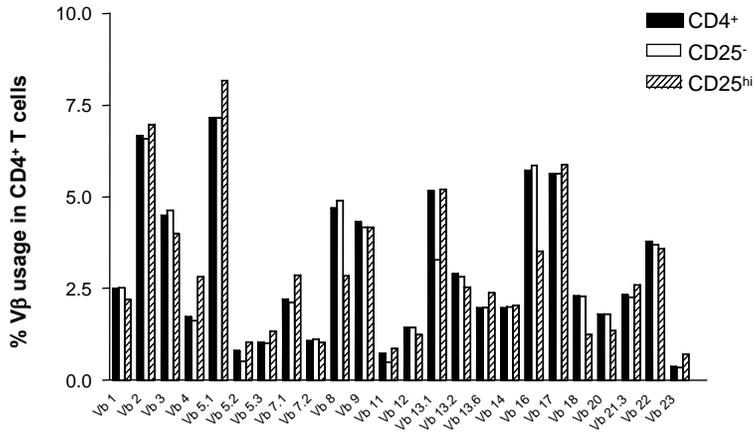
Flow FISH - Standard Curve



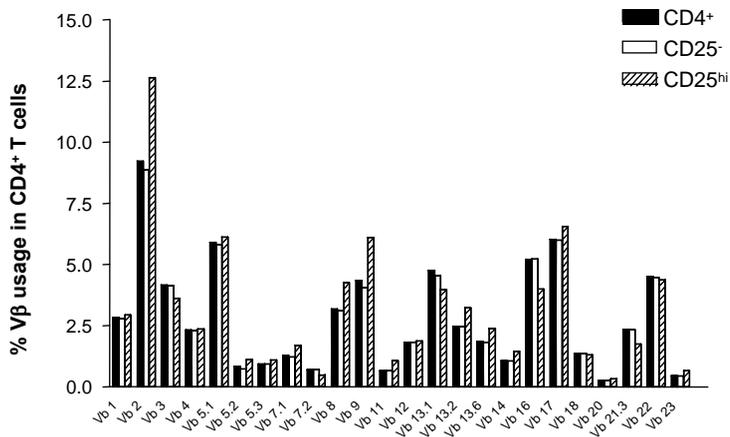
Donor A



Donor B



Donor C



Supplementary Figure 1. CD4⁺CD25^{hi} T cells have a highly differentiated memory phenotype. Freshly isolated PBMCs from young and old donors were stained with CD4-PerCP, CD25-PE and CD45RB-FITC and CD45RA-APC (**A**). Based on CD25 expression CD4 population is subdivided into CD25⁻, CD25^{int} and CD25^{hi} (left panel) as described in Materials and Methods. Histograms illustrate the CD45RA expression in each population and the percentage of CD45RA⁺ T cells is indicated. In (**B**), the histograms illustrate CD45RB expression in the 3 subsets. CD4⁺ cells were first gated on the basis of CD45RA expression so that only CD45RA⁻ (CD45RO⁺) memory cells were analysed (as described in Materials and Methods). Percentage of CD45RB^{hi} cells in each subset is indicated.

Supplementary Figure 2. Gating strategy used for FACS sorting of CD25⁻, CD25^{int} and CD25^{hi} CD4 populations. Purified CD4⁺ T cells were stained with anti-CD4 (PerCP), anti-CD45RA (FITC) and anti-CD25 (PE). Both CD4⁺CD25^{hi} and CD4⁺CD25⁻ populations were collected from the CD45RA⁻ fraction and are therefore designated as CD45RO⁺ for clarity. The CD4⁺CD25^{int} population was not collected to ensure stringency of the sorted CD25⁻ and CD25^{hi} populations. The same gating strategy was implemented for phenotypic analysis of freshly isolated PBMCs.

Supplementary Figure 3. Flow-Fish Standard curve. To construct the standard curve PBMC were collected from 20 donors of various ages. Half of each sample was used to isolate CD4⁺ T cells and telomere length was measured by Southern blot. The other half of the sample was used to measure telomeres by Flow Fish, gating on CD4⁺ T cells (in triplicate). All samples for the standard curve were run at the same time. TRF length (in kb) was plotted against mean fluorescence intensity (mean±SEM of triplicate samples) and a linear regression line was fitted using GraphPad Prism software. Dashed line indicates 95% confidence intervals.

Supplementary Figure 4. Comparison of TCR V β repertoires of CD4⁺CD25^{hi} and CD4⁺CD25⁻ cell populations. Freshly isolated PBMC from three young donors were stained with anti-CD4 and anti-CD25 and 24 different anti-TCR mAb

from the IOTest Beta Mark kit according to manufacturer's recommendations (Beckman Coulter). The TCR V β usage of the CD4⁺ (black bars), CD4⁺CD25⁻ (open bars) and CD4⁺CD25^{hi} (cross-hatched bars) T cell subsets was analysed on a FACSCalibur.