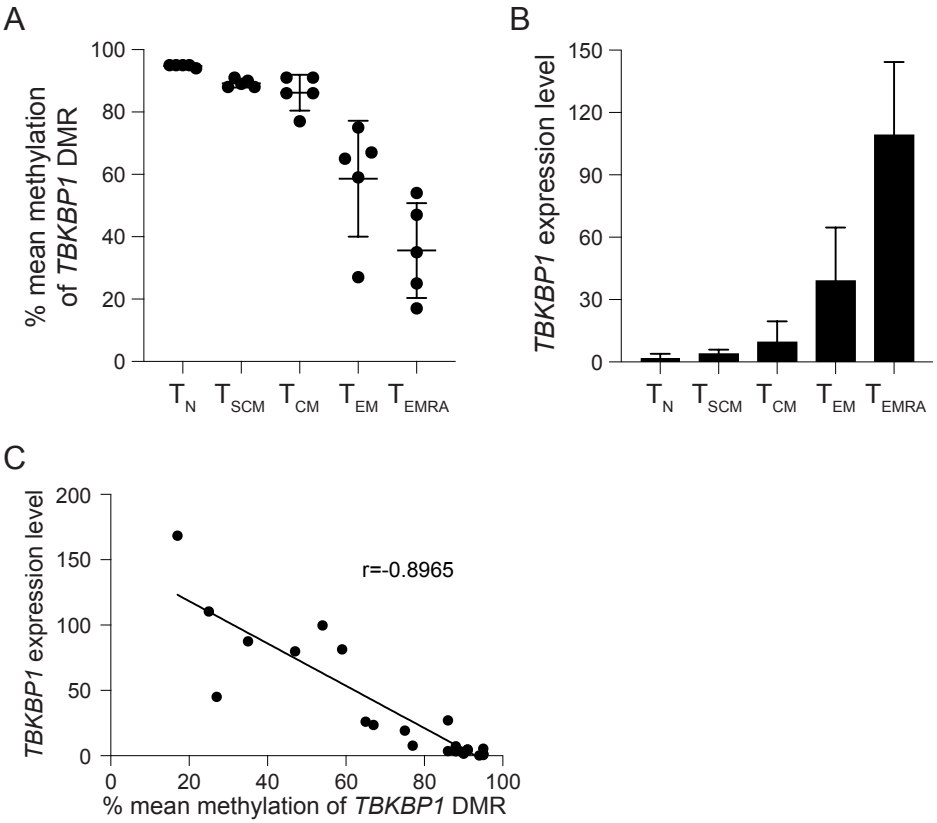

Supplementary Figure 3: Correlation between methylation level of *TBKBP1* DMR and *TBKBP1* expression in CD8⁺ T cells from CMV-seronegative donors. (A) Mean methylation level of *TBKBP1* DMR in sorted CD8⁺ T cell subpopulations T_N, T_{SCM}, T_{CM}, T_{EM}, and T_{EMRA}. Samples from five CMV-seronegative donors are shown. **(B)** *TBKBP1* expression in indicated CD8⁺ T cell subsets from five CMV-seronegative donors were analyzed by RT-PCR. **(C)** Correlation analysis visualize the relation between methylation status of *TBKBP1* DMR (x-axis) and associated *TBKBP1* gene expression (y-axis) including the linear regression line and the Pearson correlation coefficient (r).