

Figure S1. Enrichment of progression-associated differentially expressed genes for the CD4⁺ TEMRA-like **signature.** Barcode enrichment plot showing correlation of the progression-associated gene expression signature with the CD4⁺ TEMRA-like signature. Regions are ordered on the plot from right to left from most positive logFC to most negative when comparing progressors to non-progressors (x axis). Vertical bars indicate regions increased in CD4⁺ TEMRA cells in comparison with other T cell memory subtypes. Enrichment was assessed by fry gene set tests.