



Figure 3-figure supplement 2. Sorting strategy of central memory CD4 T cell subsets and expression of candidate genes. a-b) The transcriptome of CD4 and CD8 T cell subsets purified from 4 to 5 CMV-seropositive donors was analysed by genome-wide mRNA expression arrays. Log2 expression values of 13,551 probes with a variance>0.01 corresponding to 10,699 unique genes were submitted to hierarchical clustering (**b**) and principal component (**c**) analyses.