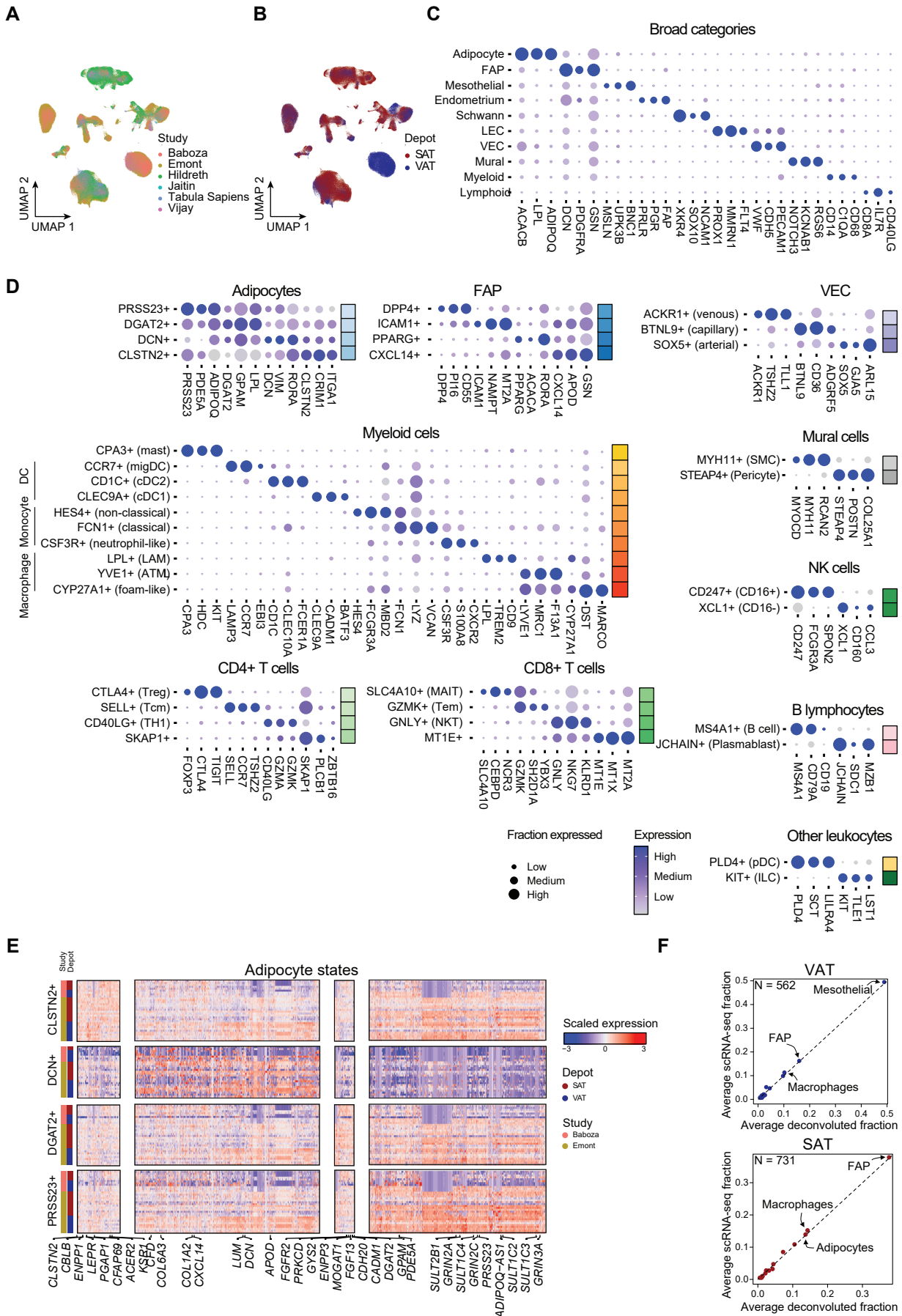


Supplementary Figure 5



Supplementary Figure 5: A-B) WATLAS UMAPs superimposed with (A) study and (B) depot annotations. **C-D)** Dotplots showing marker genes for broad cell type categories and subtypes, as indicated on the figure. **E)** Heatmap showing pseudo-bulk expression levels of genes identified as differentially expressed in the indicated adipocyte populations across all populations, depots, and studies. **F)** Scatterplot showing the linear relationship between average fraction of the cell types in WATLAS and the average deconvoluted fractions for VAT and SAT depots (left and right panel respectively).