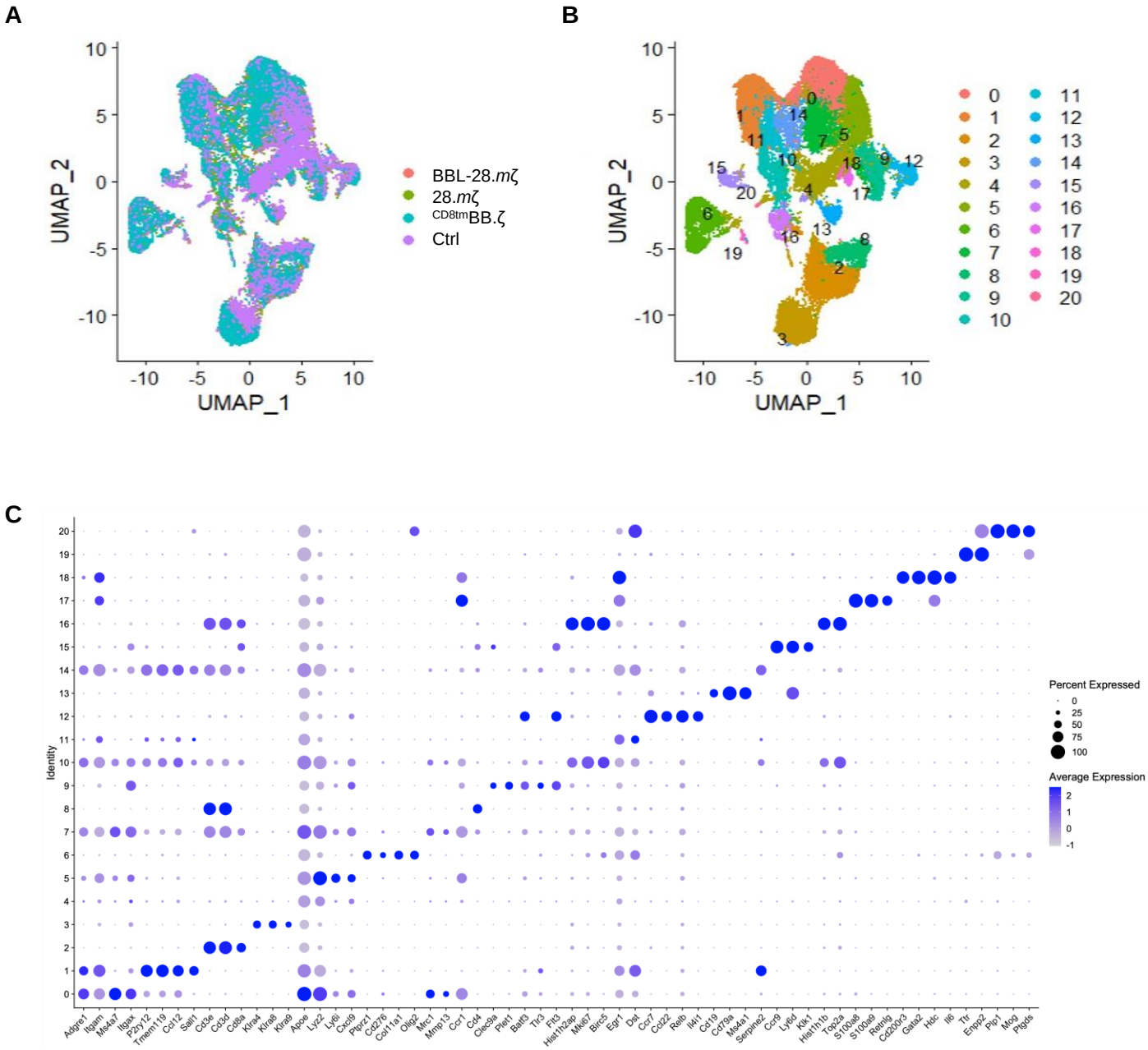

Supplementary Fig. S7: Analysis of scRNAseq data using 10 dimensionality reduction method. **(A)** UMAP visualization of major immune cell clusters per treatment group. **(B)** UMAP plot of single cells from four treatment groups colored by their distribution into 21 major cell types. **(C)** Dot plot visualizing the canonical cell-type marker expression within each cell cluster.