



Supplemental_Fig_S12: Unsupervised hierarchical clustering of Type 2 diabetic single cell transcriptomes separates samples by cell type.

Circular dendrogram of Type 2 diabetic single cell transcriptomes illustrates similar clustering by cell type as observed in non-diabetic samples. 1908 highly expressed genes with $\log_2(\text{CPM}) > 10.5$ were used in the analysis. Color labels are the same as shown in Figure 3B.