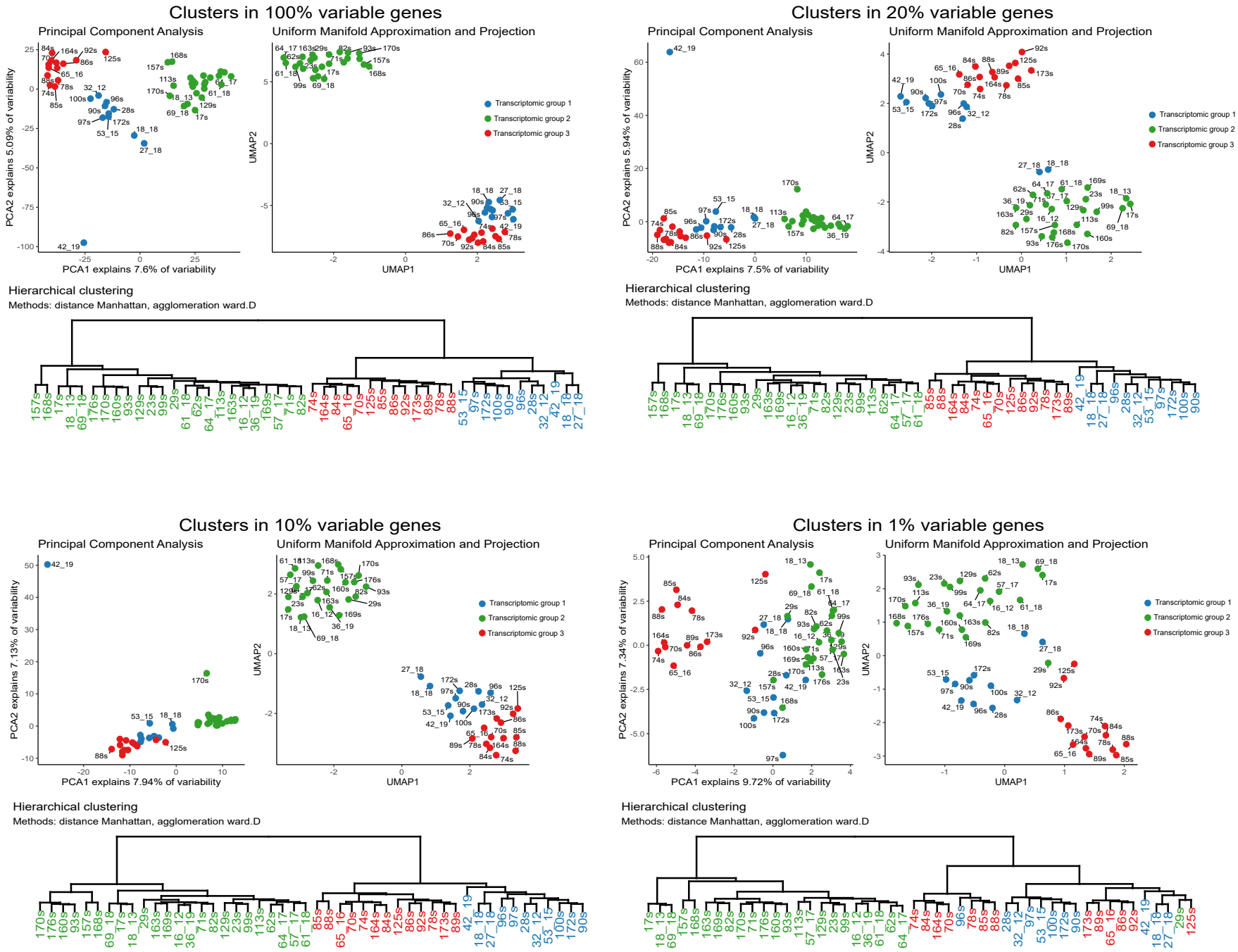

Figure S1. Overall gene expression profile in somatotroph tumors including Principal component analysis (PCA) and uniform manifold approximation and projection (UMAP) results and Hierarchical clustering based on the 100%, 20%, 10% and 1% of genes with most variable expression.