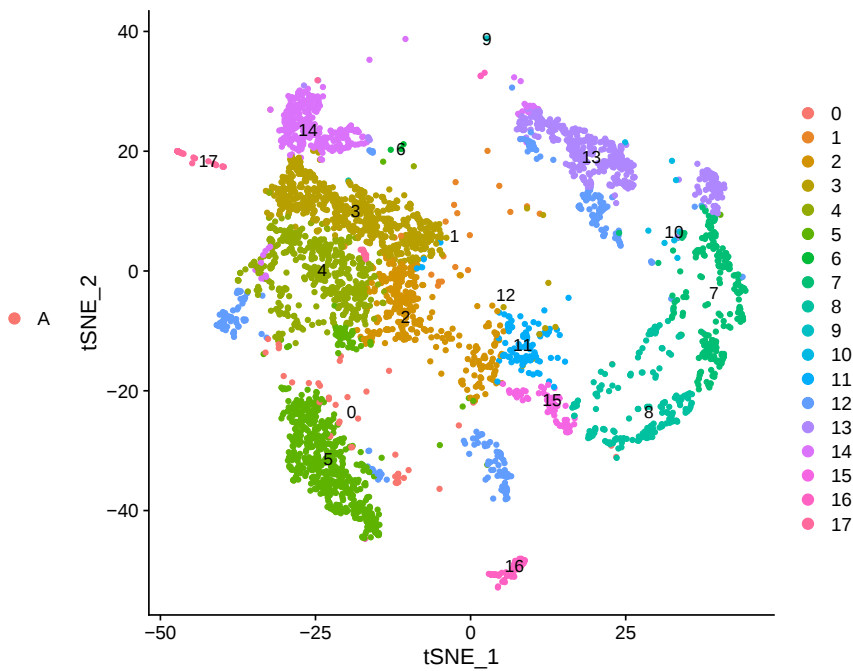
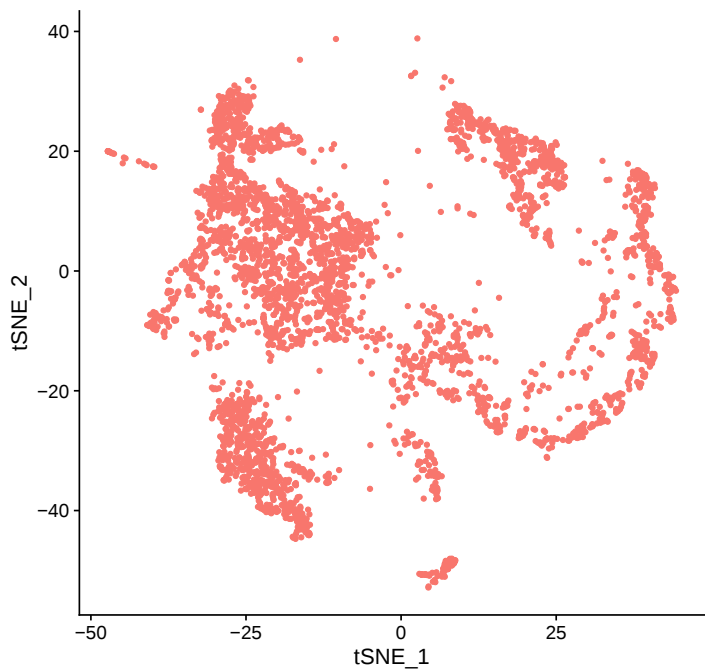
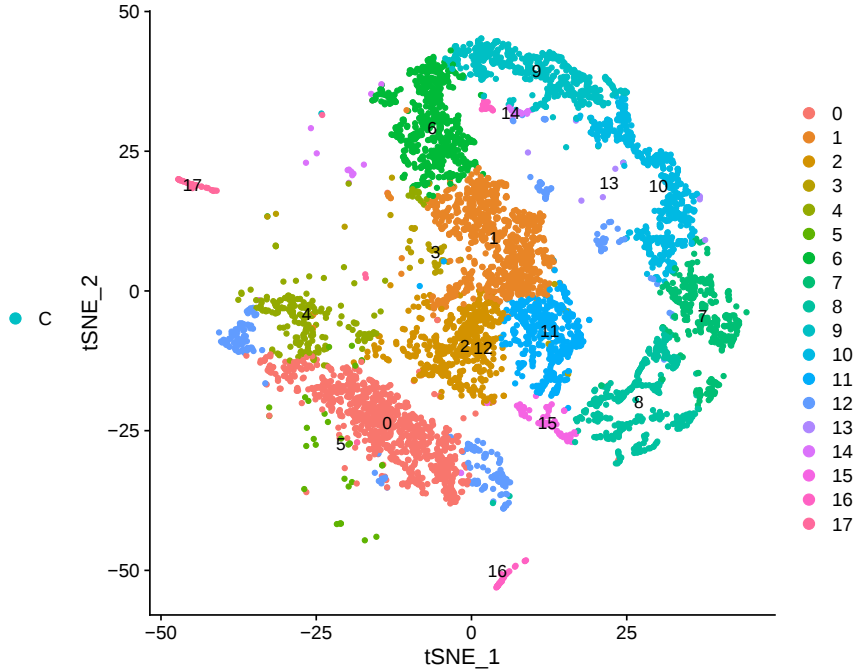
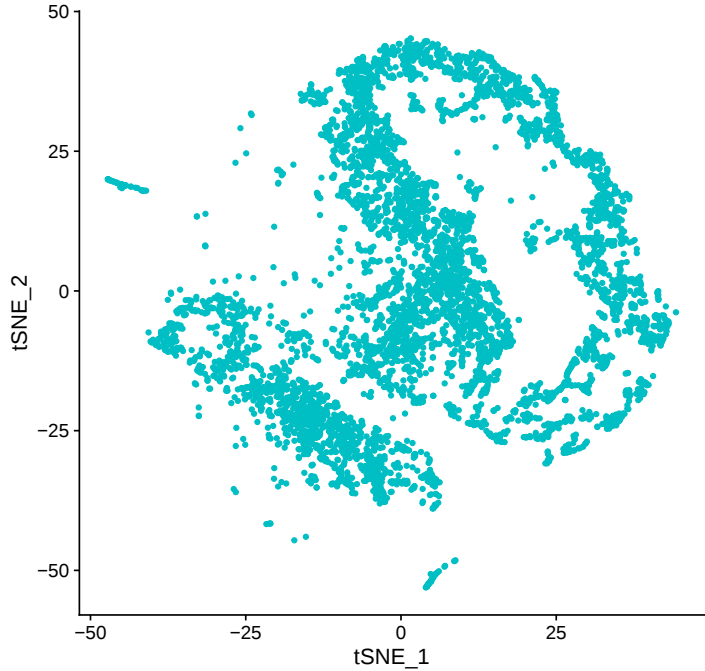
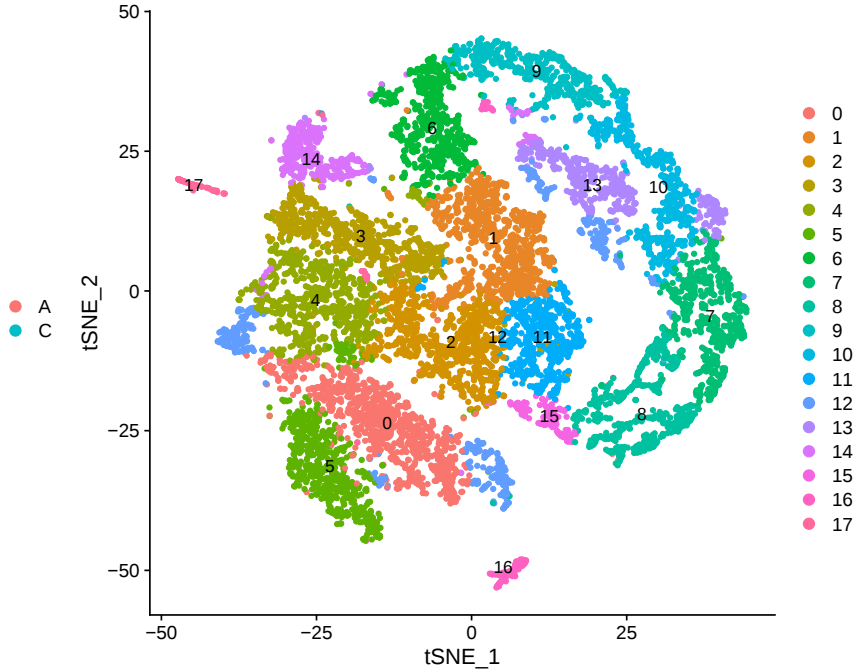
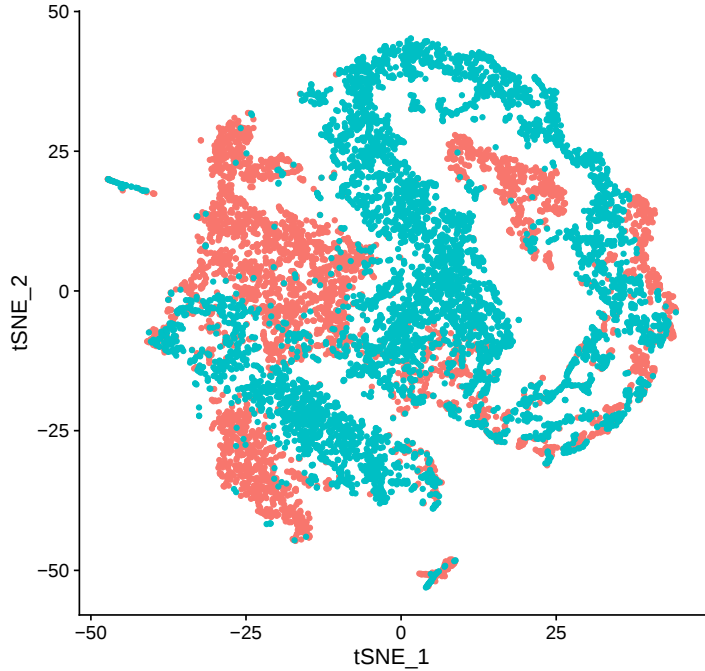








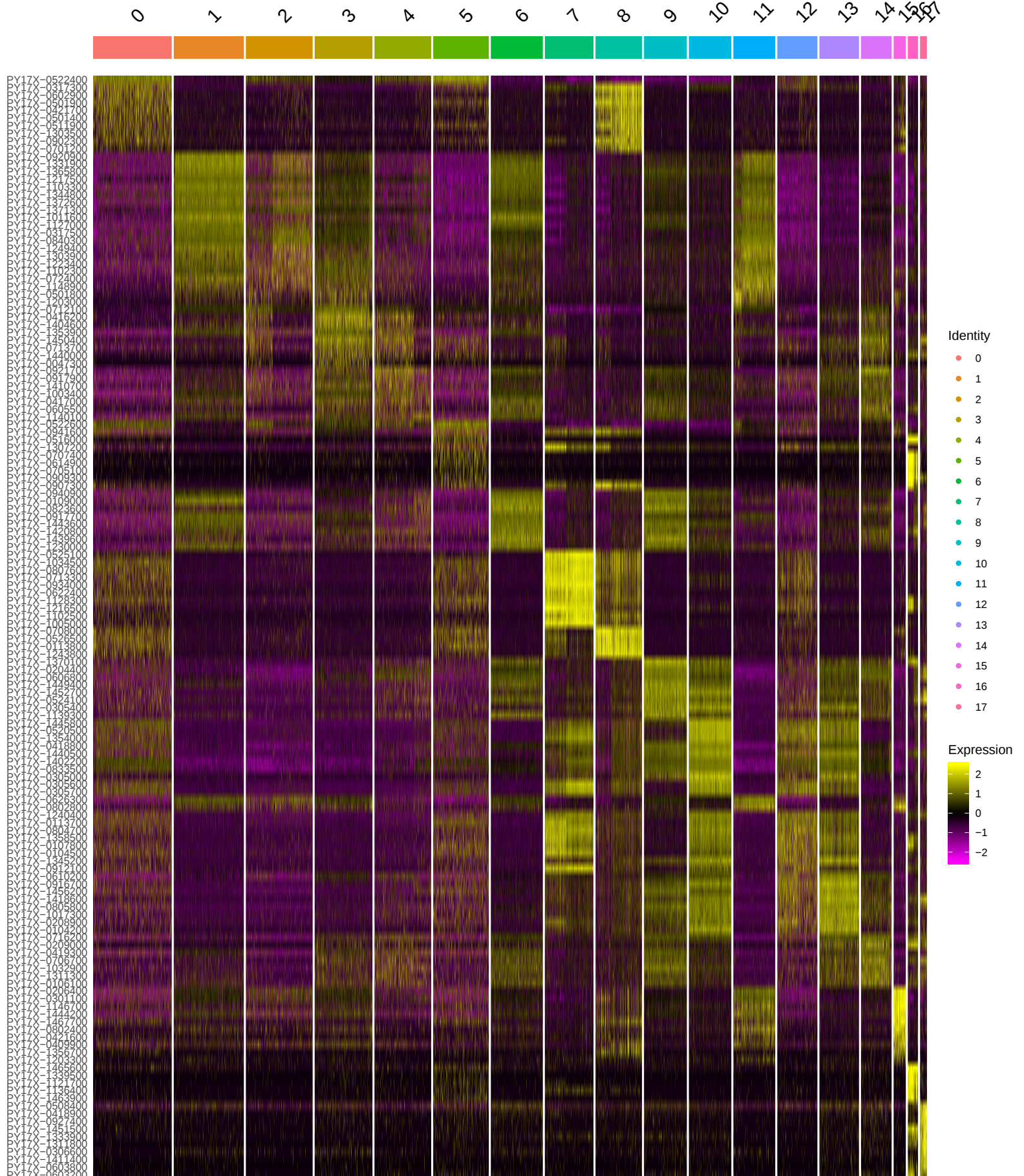


Fig. 1. t-distributed stochastic neighbour embedding (tSNE) plots of single-cell transcriptomes (SCTs) from the ART-treated groups showing 18 clusters.

Fig. 2. t-distributed stochastic neighbour embedding (tSNE) plots of single-cell transcriptomes (SCTs) from the control groups showing 18 clusters.

Fig. 3. t-distributed stochastic neighbour embedding (tSNE) plots of single-cell transcriptomes (SCTs) from the ART-treated and control groups showing 18 clusters.

Fig. 4. The heatmap of marker genes which determine 18 clusters.