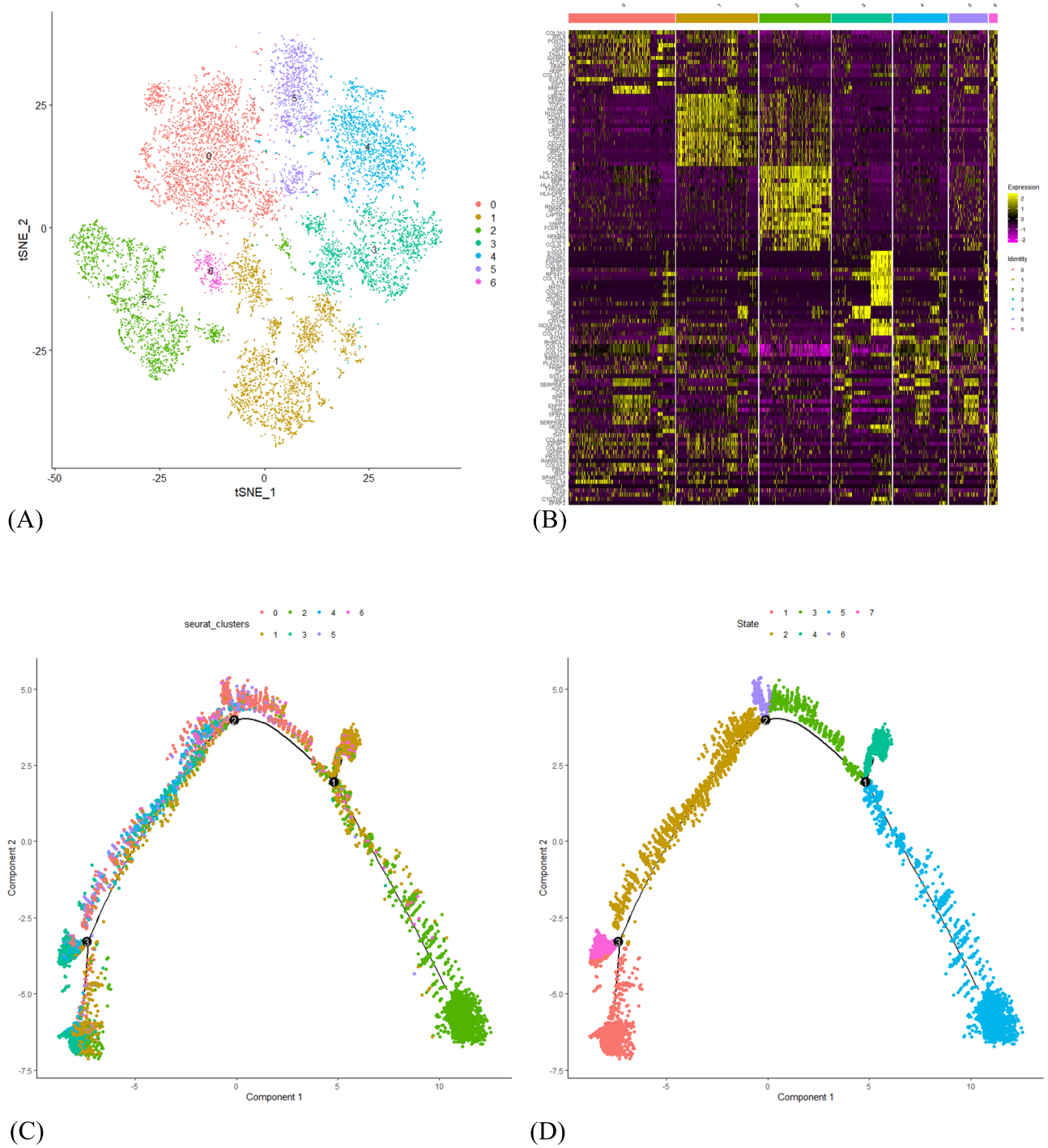

Supplement Figure S4: Single-cell transcriptomic analyses of all cells and the subclusters of osteosarcoma cells. (A) The top 5 marker genes of each subtype of all cells are displayed in the heatmap. (B) The t-distributed stochastic neighbor embedding (t-SNE) plot of the subtypes of osteosarcoma cells. (C) Via cell trajectory analysis, the top 5 marker genes of each state of osteosarcoma cells are displayed in the heatmap. (D) Pseudotime and trajectory analysis.