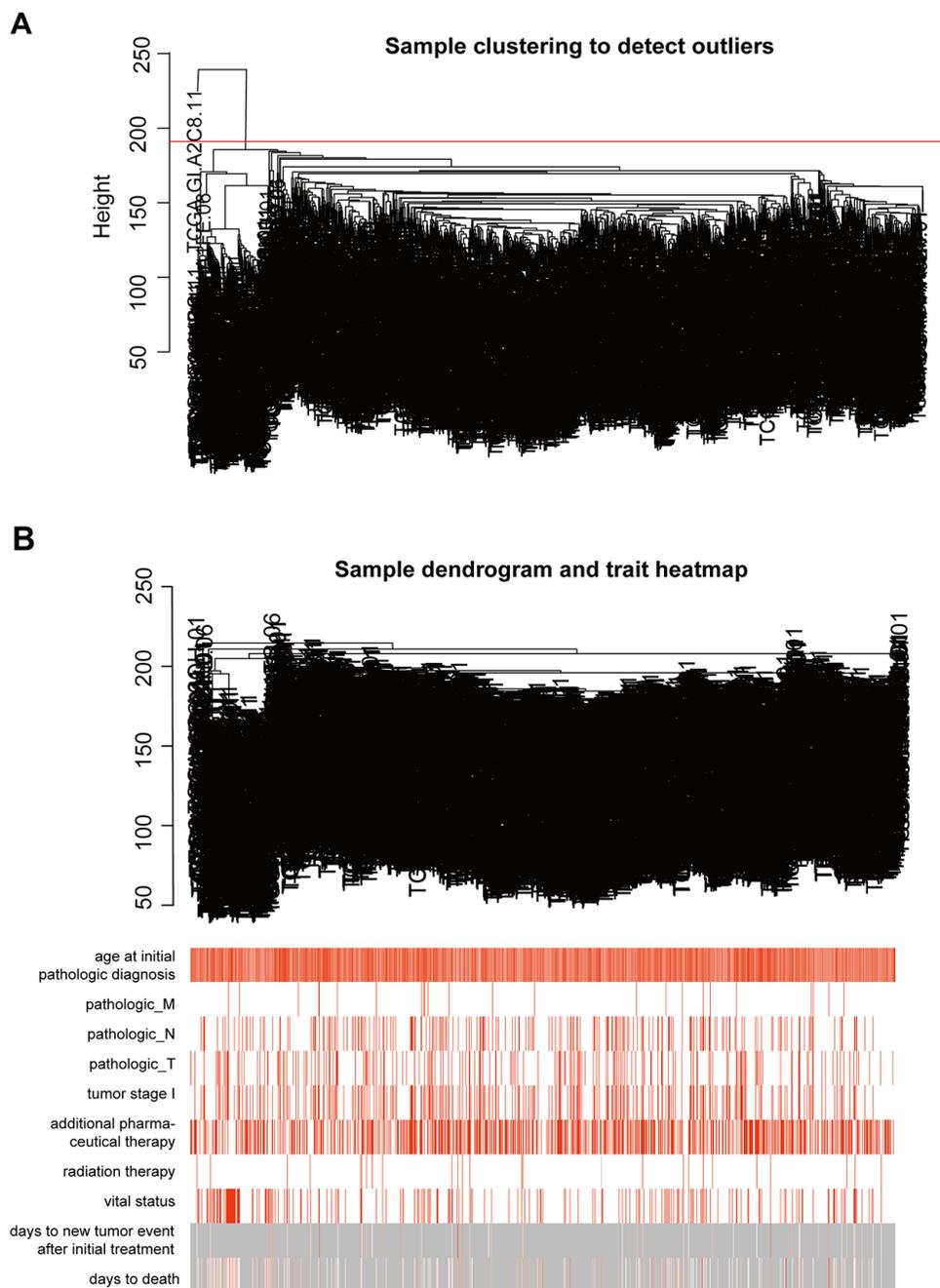


SUPPLEMENTARY FIGURES



Supplementary Figure 1. Sample cluster analysis. (A) Sample clustering to detect outliers based on RNA data (the red line (cut height = 200 cm) was used as the filter to remove outlier samples from the sample tree). (B) Sample dendrogram and clinical trait heatmap based on gene expression and clinical data. TNM = tumor, node, metastasis (classification).

