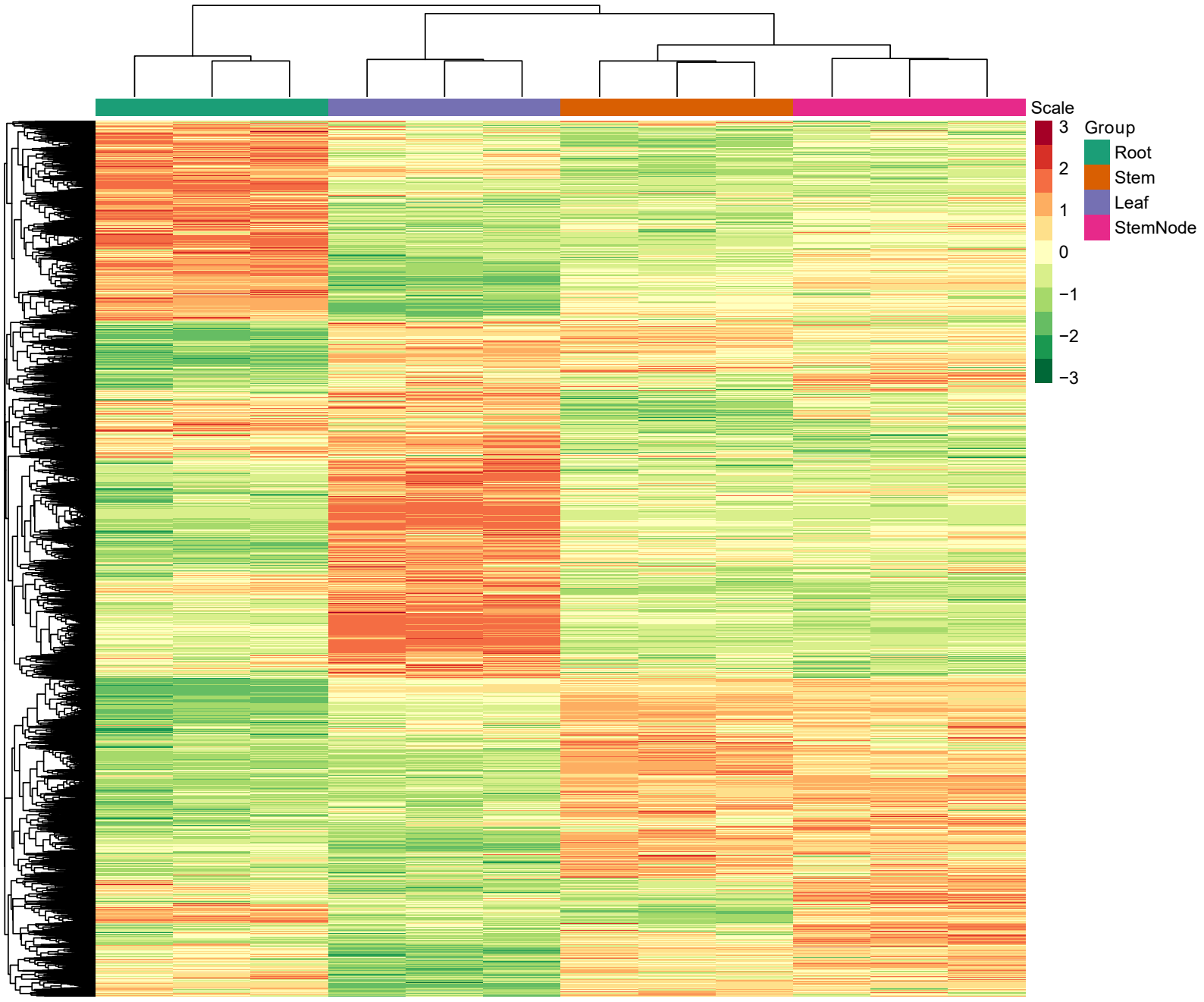


Figure S1: Cluster heatmap of each differential group. The diagram presents the result of a two-way hierarchical clustering of all the DEGs and comparison combinations. The X-axis represents the sample name and hierarchical clustering results, and the Y-axis represents the differentially expressed genes and hierarchical clustering results. Red represents high expression, and green represents low expression