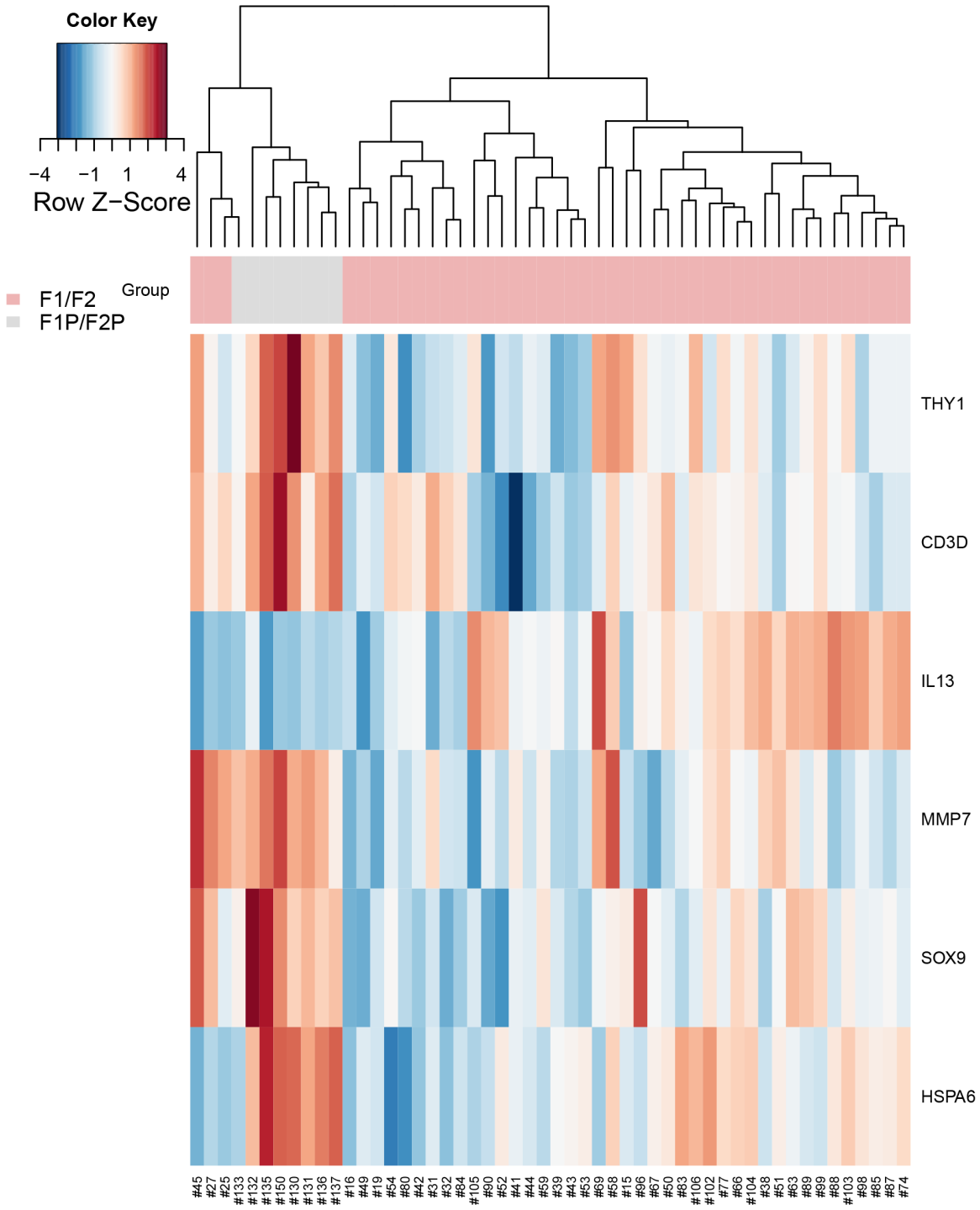


Supplementary Figure S1. Hierarchical clustering of NAFLD patients with F1/F2 including 8 fast progressors (F1P/F2P). The six genes with the highest ROC performances in discriminating fast progressors were used to generate a heatmap using log10-transformed gene expression data. Color key represents row Z-scores.