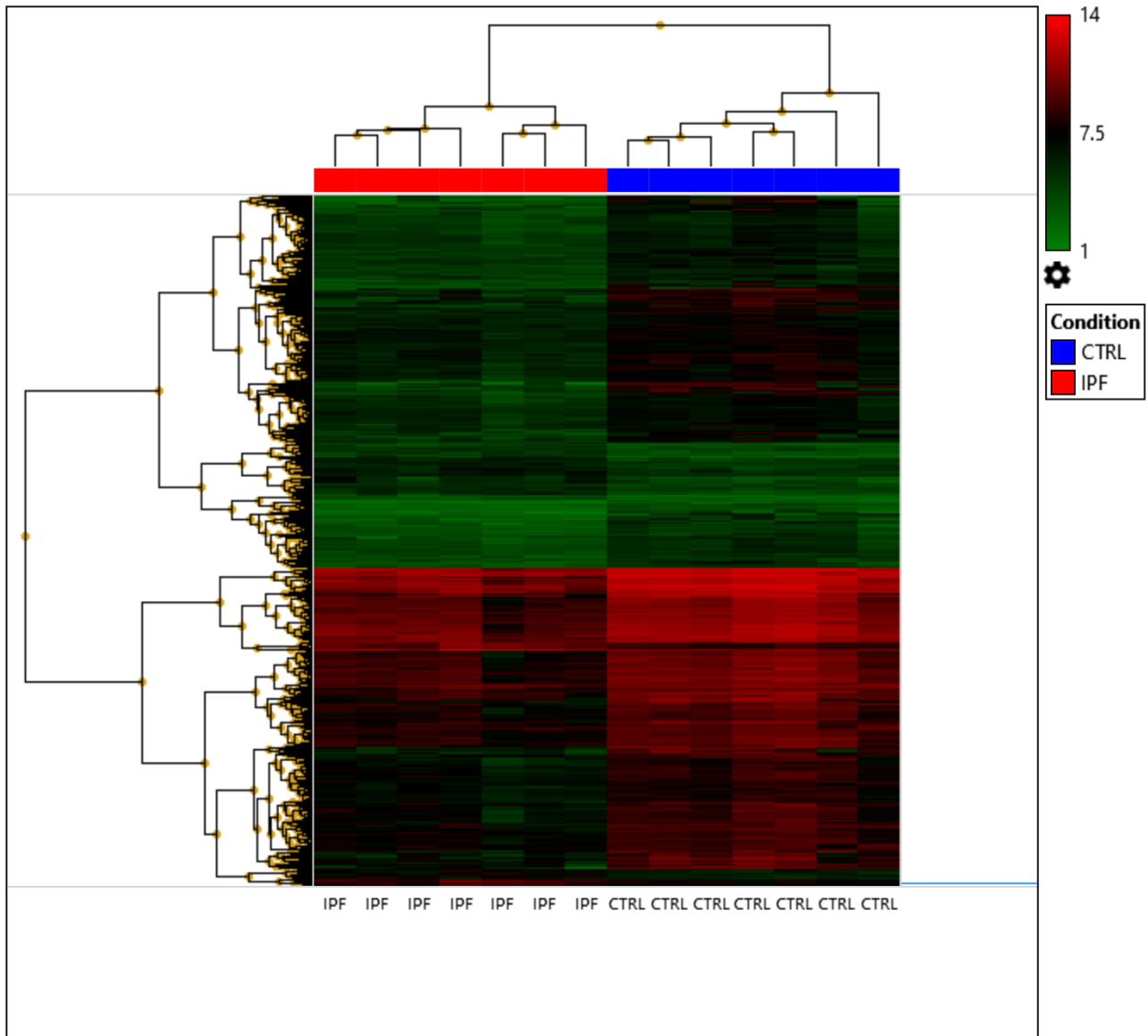


SUPPLEMENTARY DATA



Supplementary figure 1: Hierarchical clustering of DE transcripts between IPF patients versus Controls.

Each row represents the relative expression level of a single transcript and each column represents a single analyzed sample. Fold change >2.0 or <-2 and FDR-corrected p -value <0.05 .