A

Color Key

sample correlation matrix
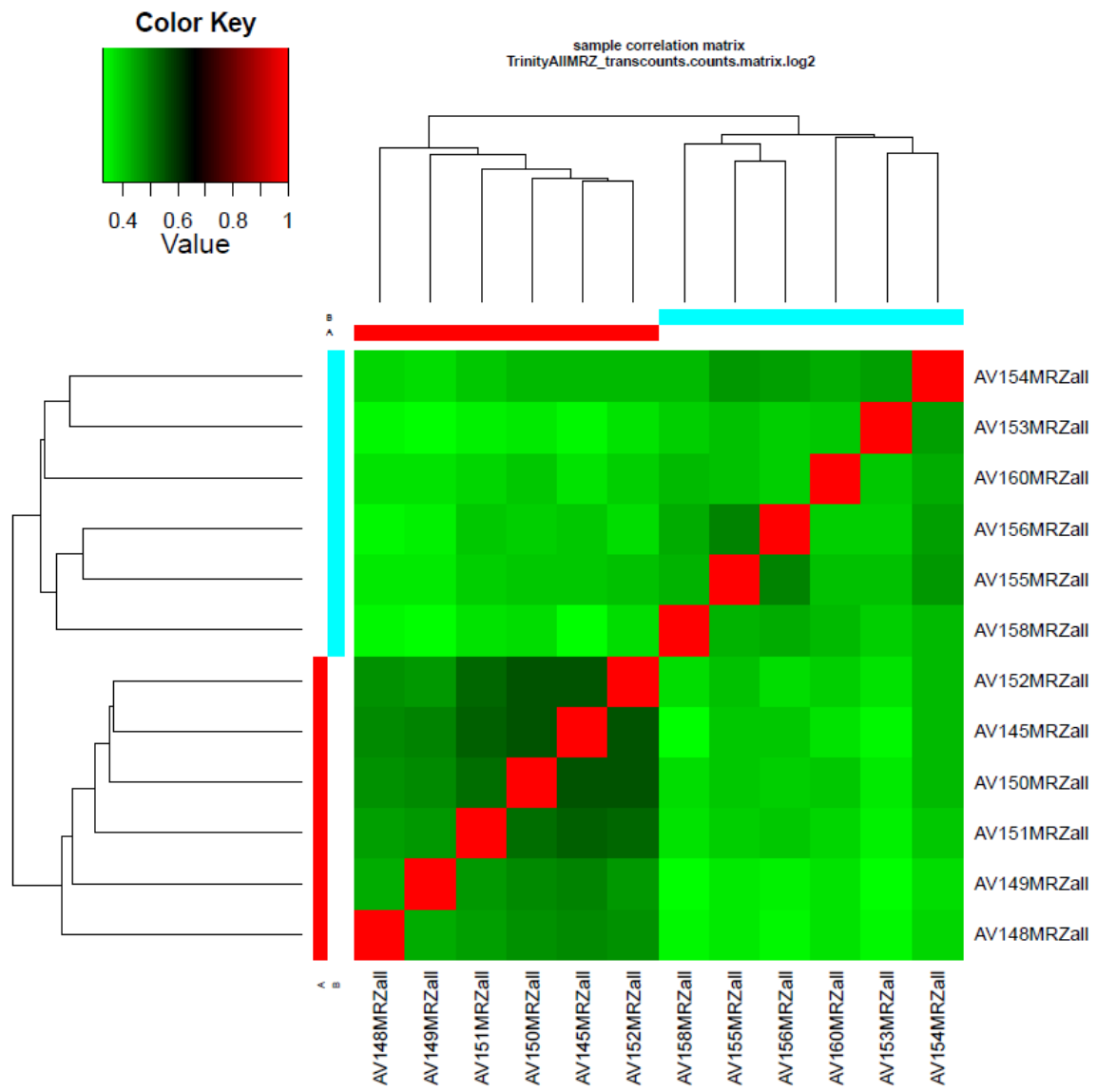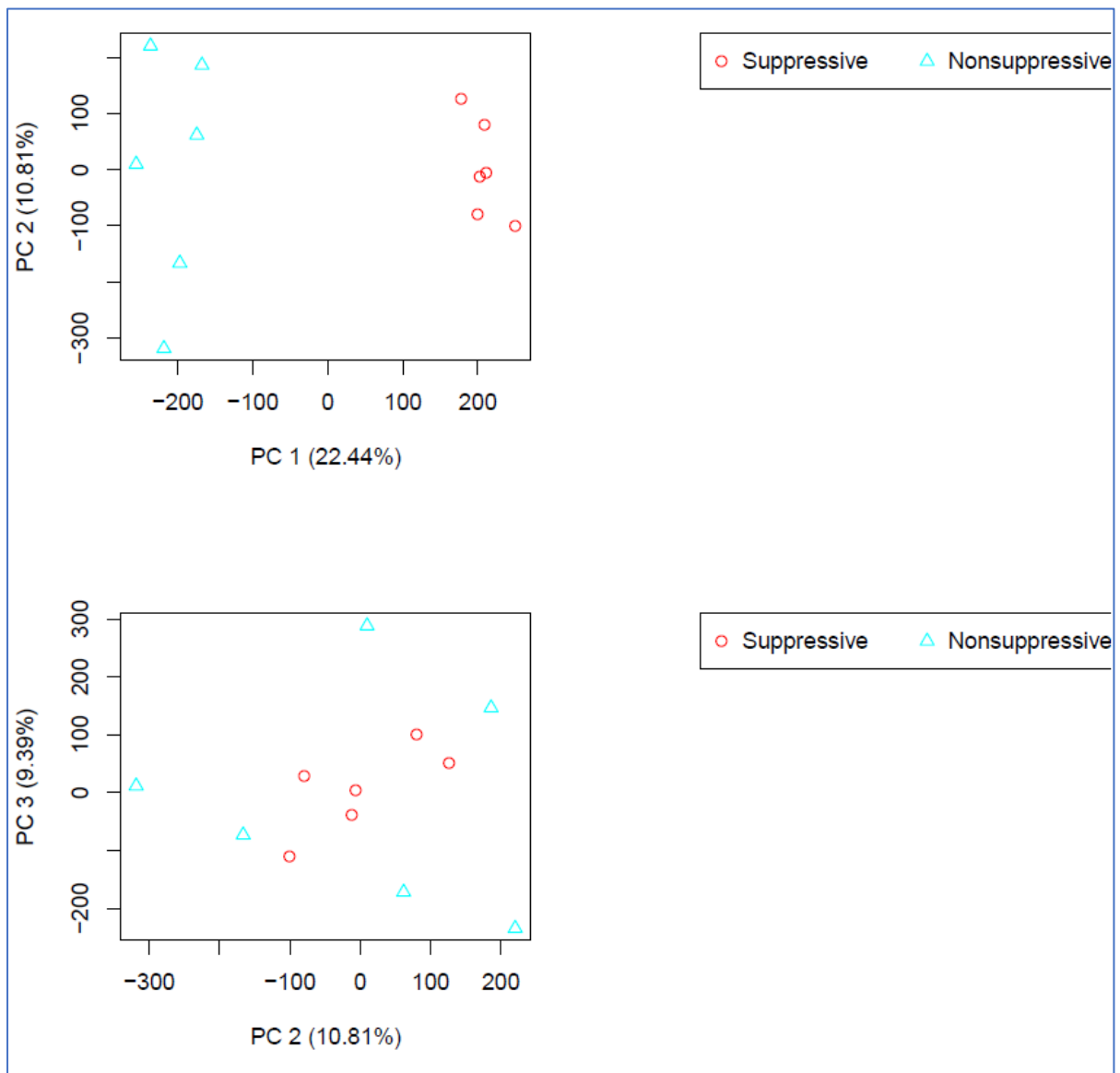TrinityAllMRZ_transcounts.counts.matrix.log2

B

FIGURE S4. The similarity of biological replicate metatranscriptomes from the suppressive (AV145MRZall-AV152MRZall) and non-suppressive (AV153MRZall-AV160MRZall) field was examined by **A)** a Pearson correlation heatmap based on transcript expression values. Replicates are highly similarly when they are closer to a value of 1; and **B)** a principal component analysis (PCA) plot.