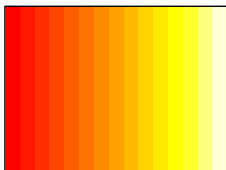
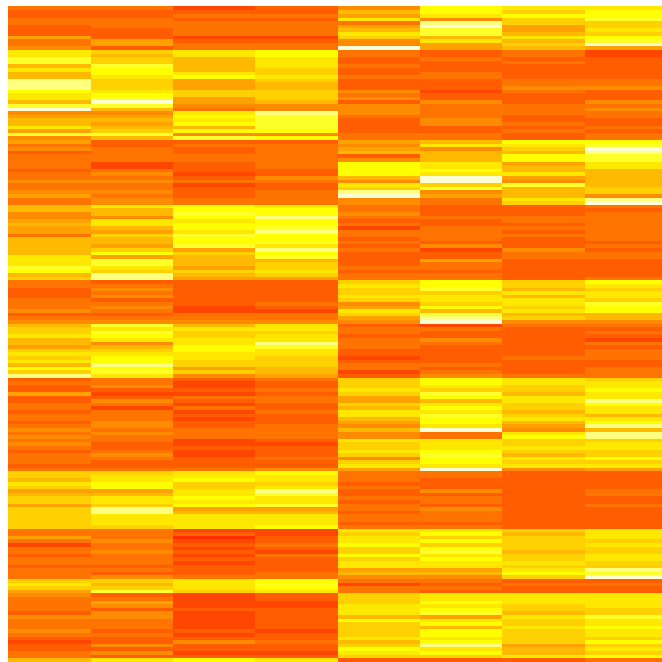
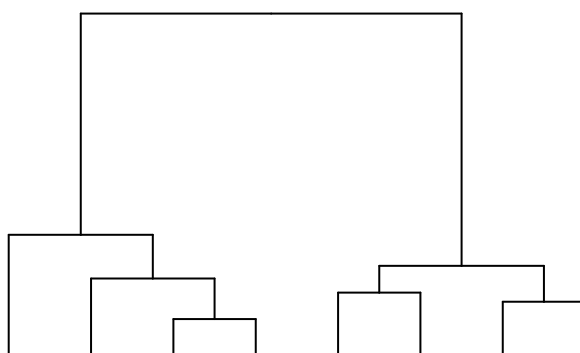


Color Key



-2 0 1 2
Row Z-Score



Seud1-2

Seud1-1

Seud1-3

Seud1-4

NHR40-4

NHR40-1

NHR40-3

NHR40-2

Heatmap visualization of gene expression data across eight samples (Seud1-2, Seud1-1, Seud1-3, Seud1-4, NHR40-4, NHR40-1, NHR40-3, NHR40-2). The color scale represents the Row Z-Score, ranging from -2 (red) to 2 (yellow). The dendrogram on the left indicates hierarchical clustering of the samples.