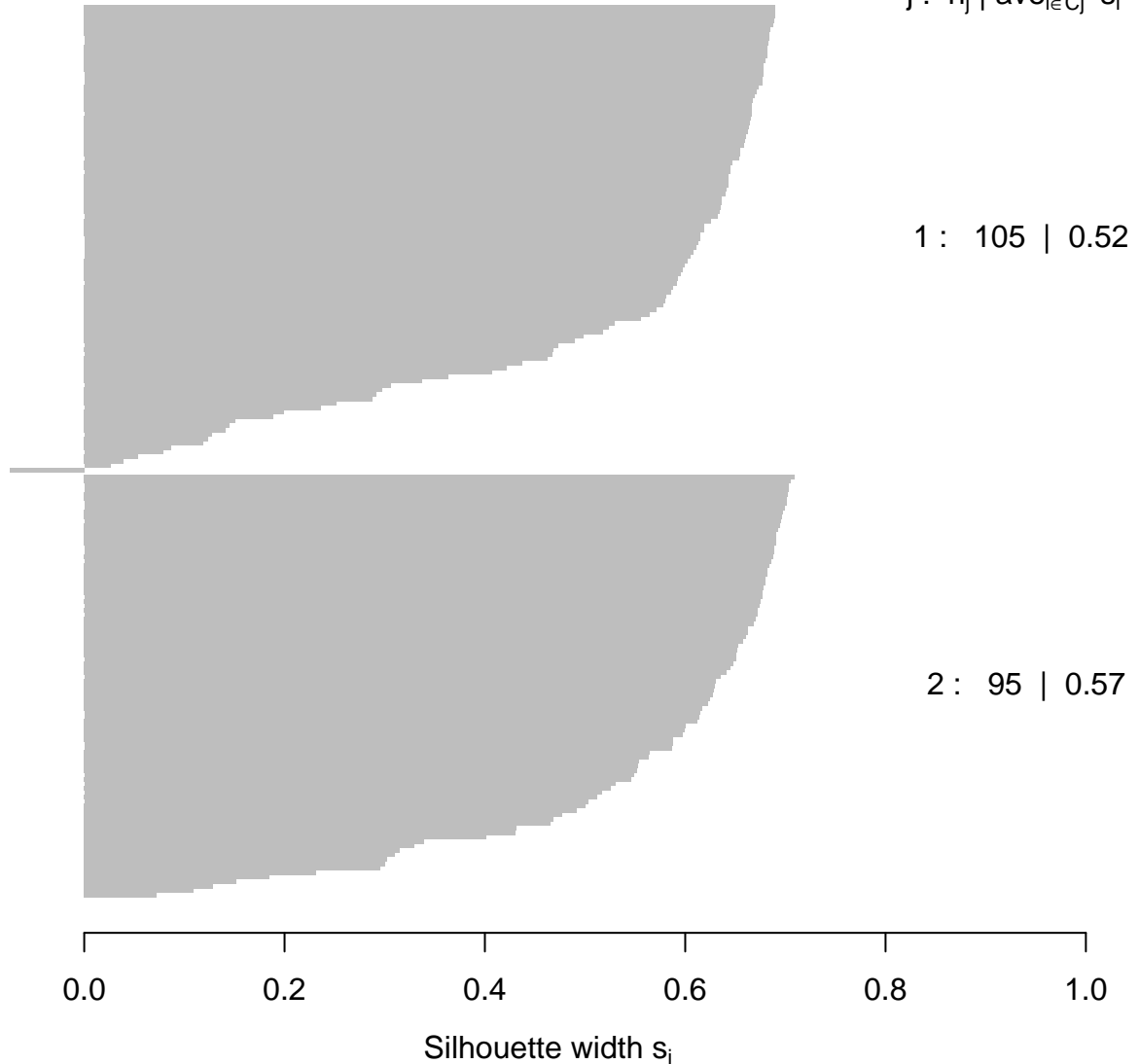


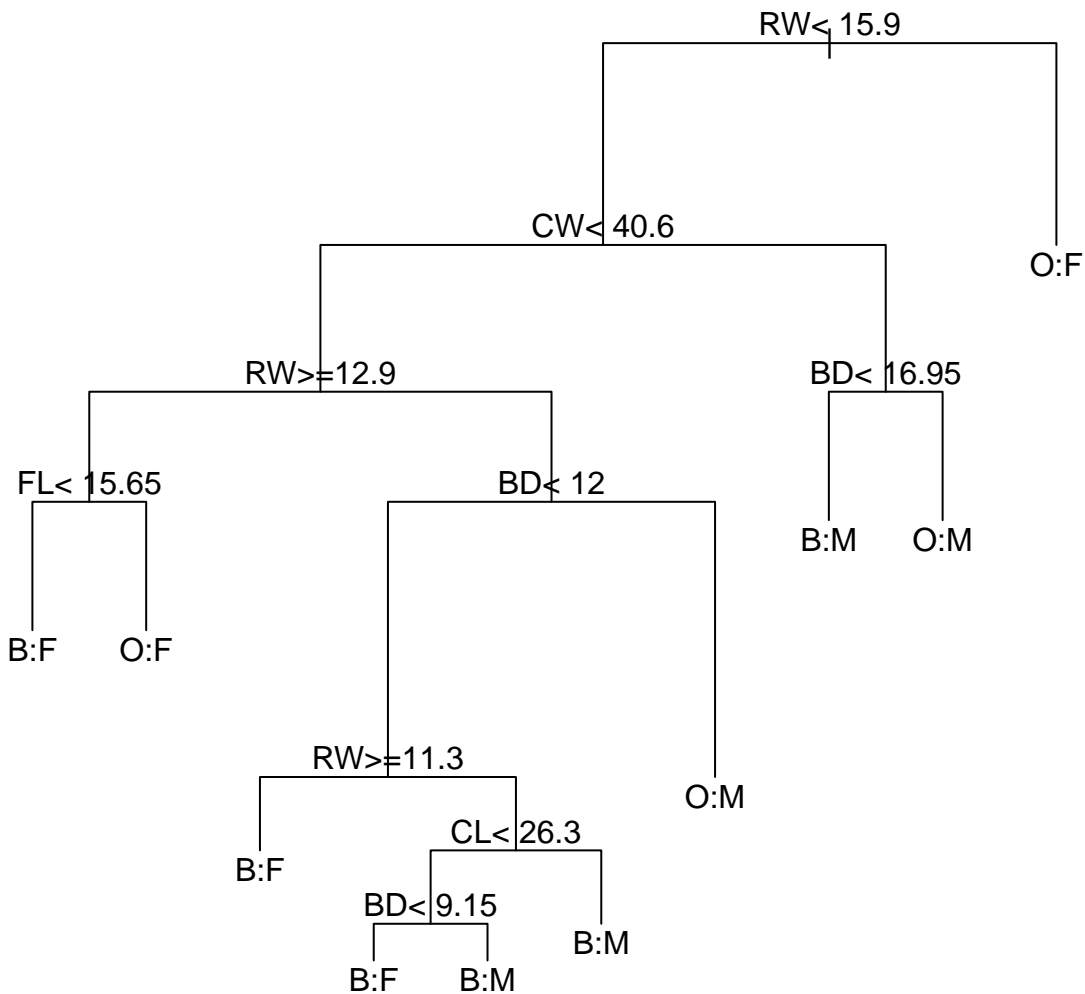
Gene names (e.g., LOC100000000, LOC100000001, etc.) are listed along the bottom of the heatmap, corresponding to the rows of the data matrix.

# Silhouette plot of (x = tr, dist = dist(X))

n = 200

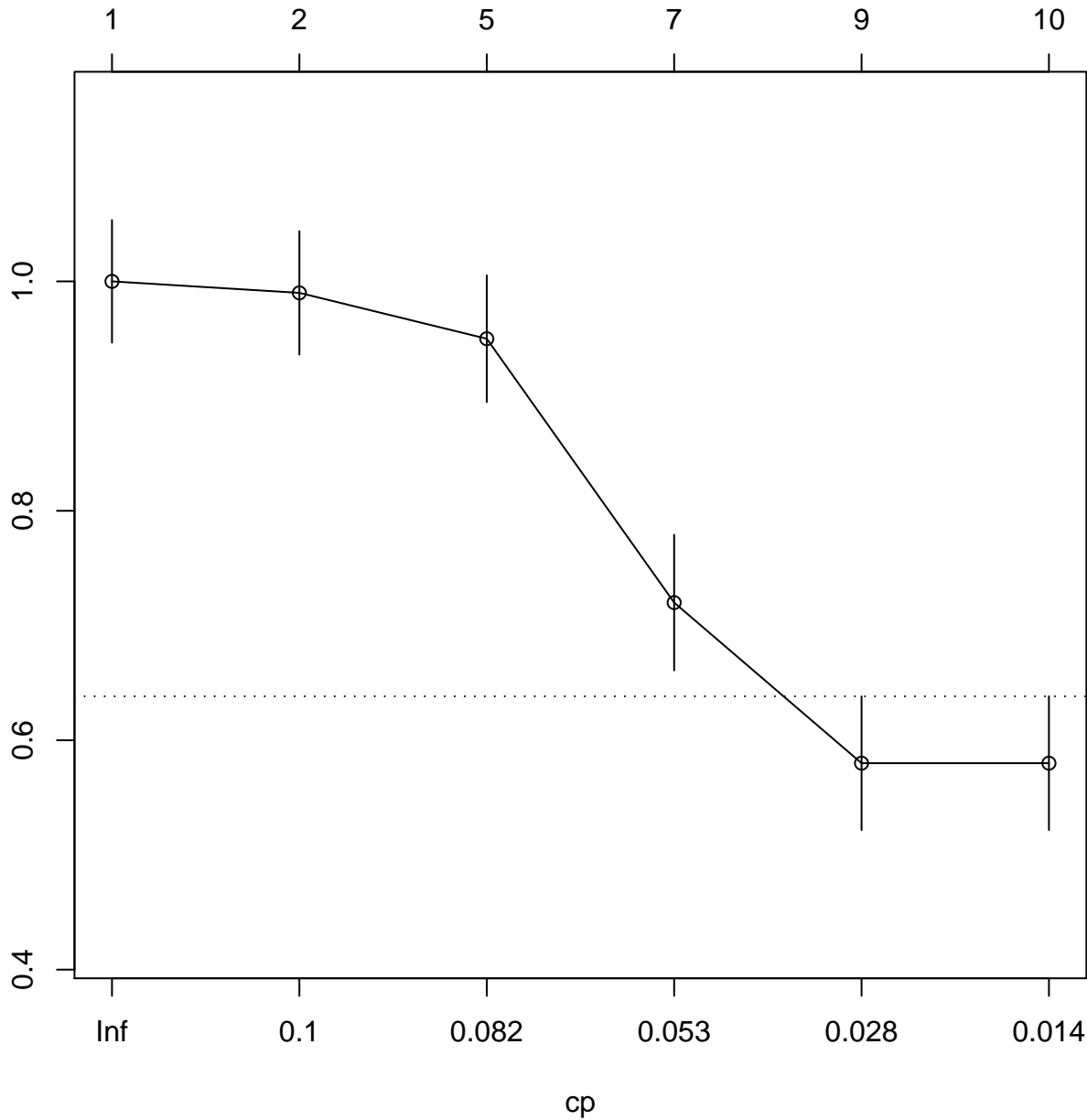
2 clusters  $C_j$   
 $j : n_j \mid \text{ave}_{i \in C_j} s_i$





size of tree

X-val Relative Error





PCg

