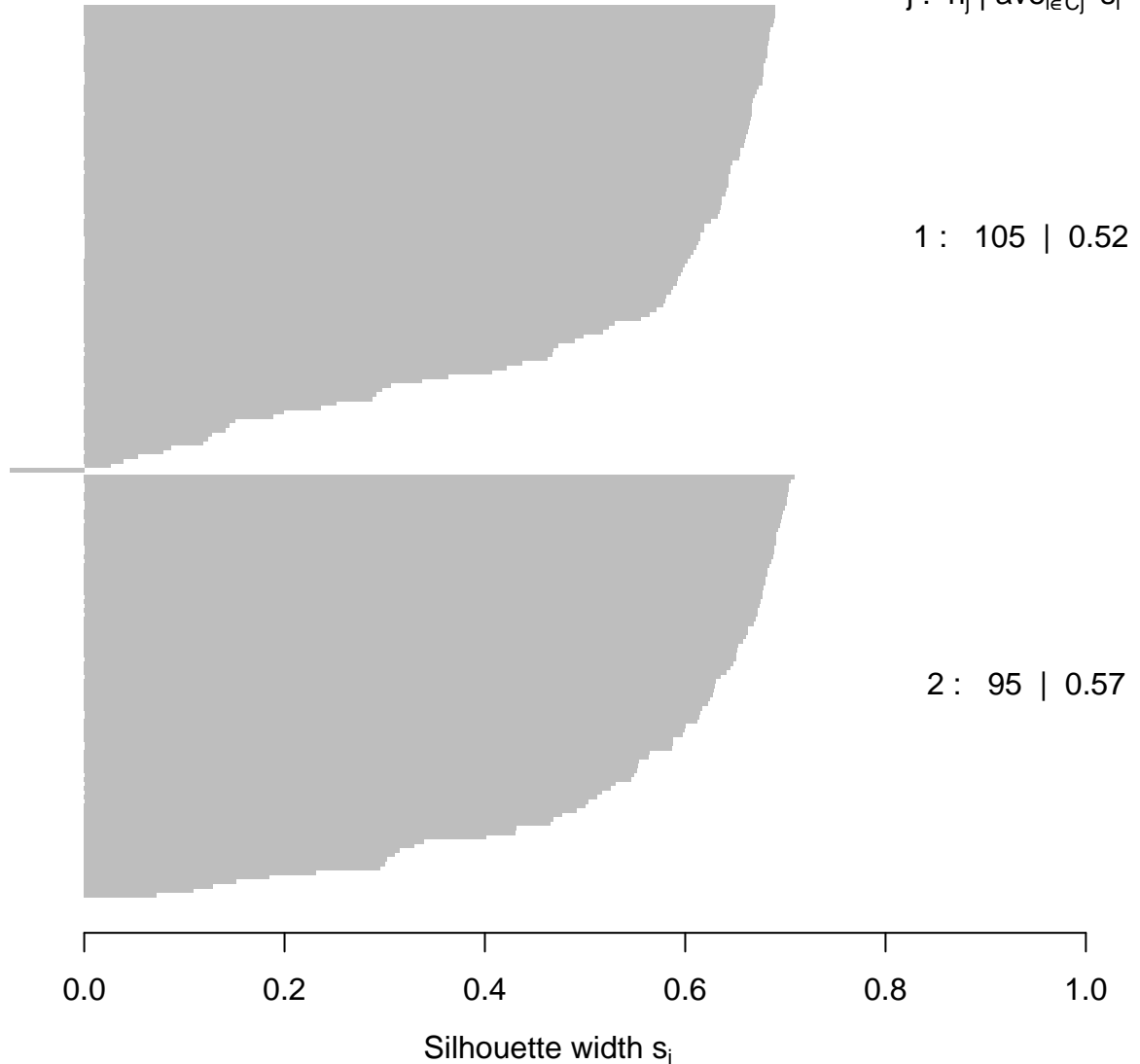


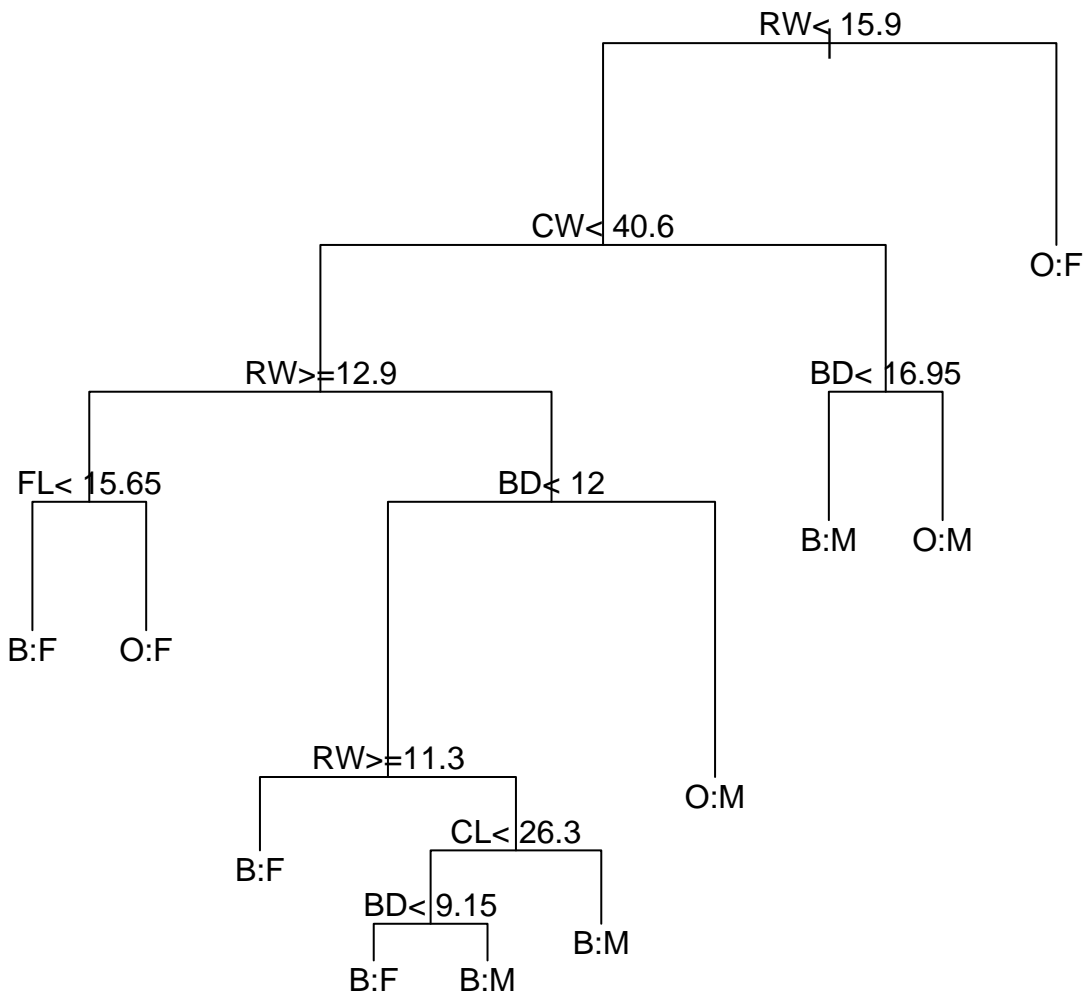
Gene names (e.g., LOC100000000, LOC100000001, etc.) are listed along the bottom of the heatmap, corresponding to the rows of data.

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

n = 200

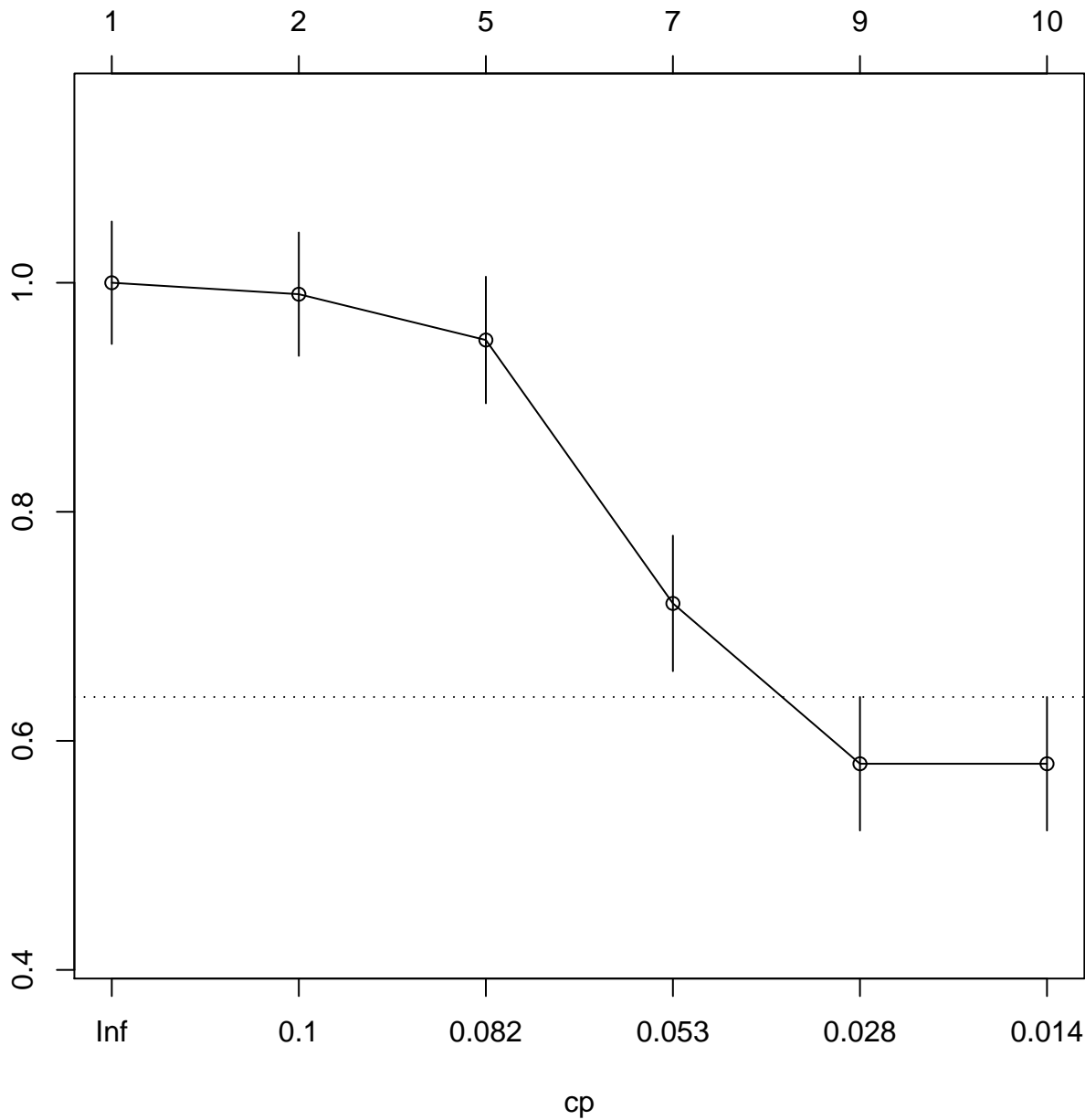
2 clusters C_j
 $j : n_j \mid \text{ave}_{i \in C_j} s_i$





size of tree

X-val Relative Error



PCg

