Ting Huang

Adapted from the slides of David M. Blei from Princeton University

http://www.cs.princeton.edu/courses/archive/spr08/cos424/slides/clustering-2.pdf

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- Visualizing this tree provides a useful summary of the data

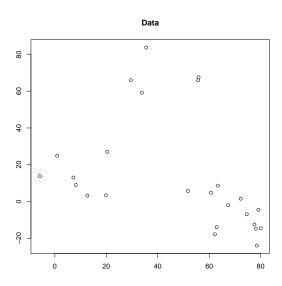
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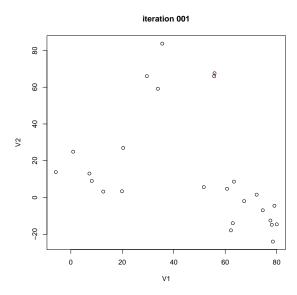
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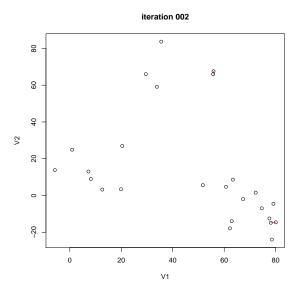
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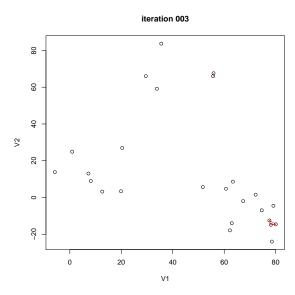
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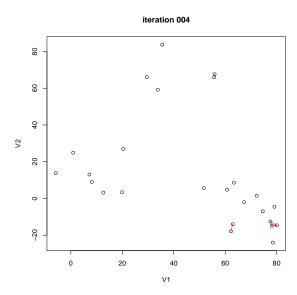
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 - 3 Until: all the data are merged into a single cluster

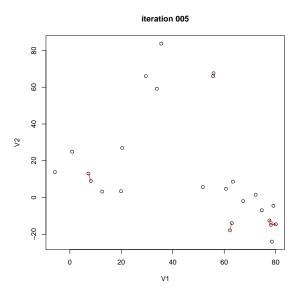


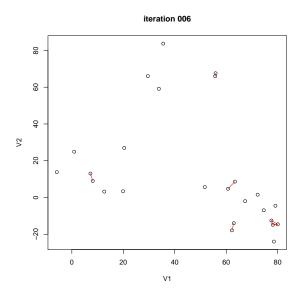


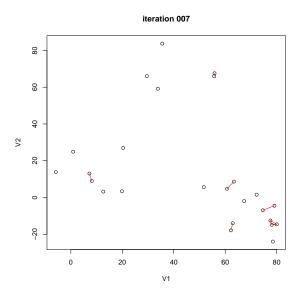


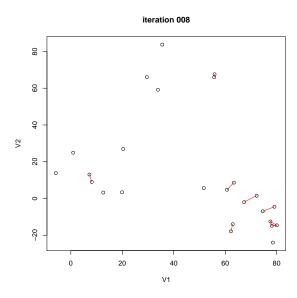


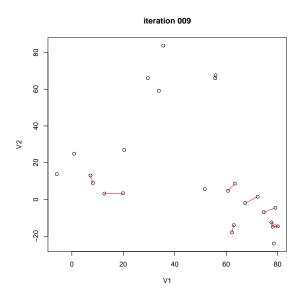


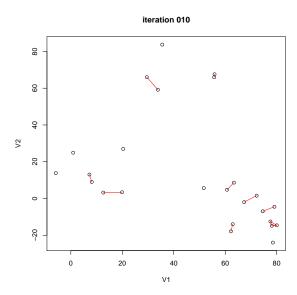


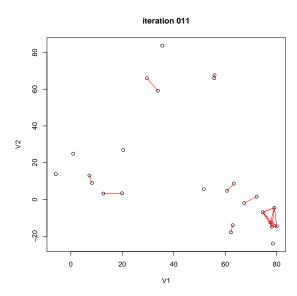


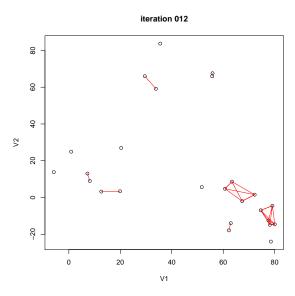


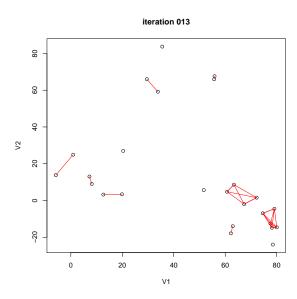


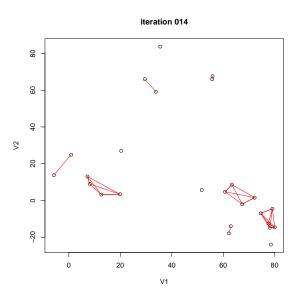


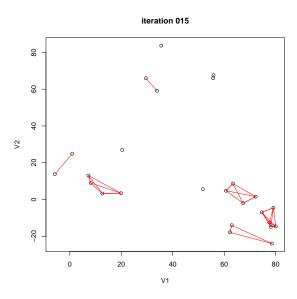


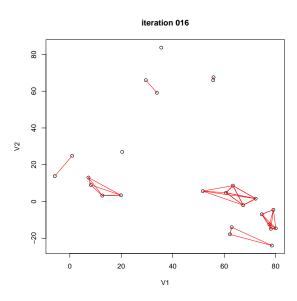


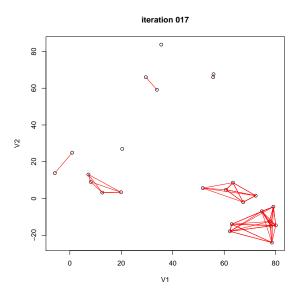


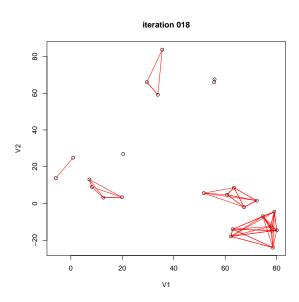


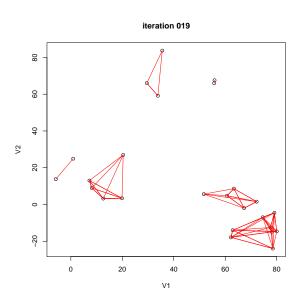


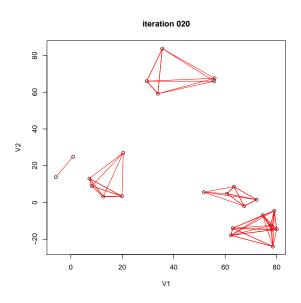


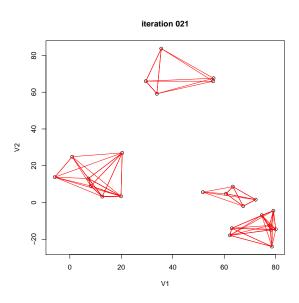


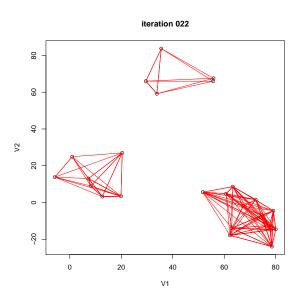


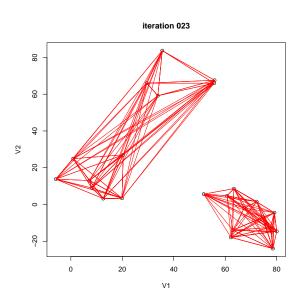


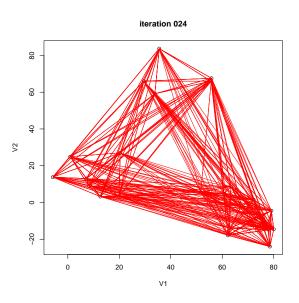




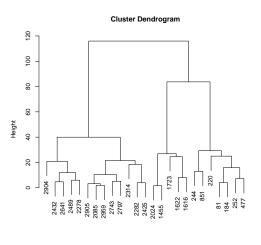








Dendrogram of example data



dist(x) hclust (*, "complete")

Groups that merge at high values relative to the merger values of their subgroups are candidates for natural clusters. (Tibshirani et al., 2001)

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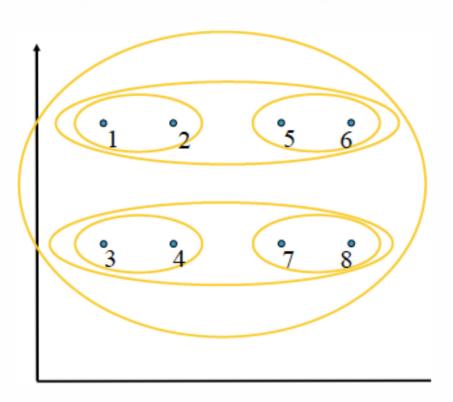
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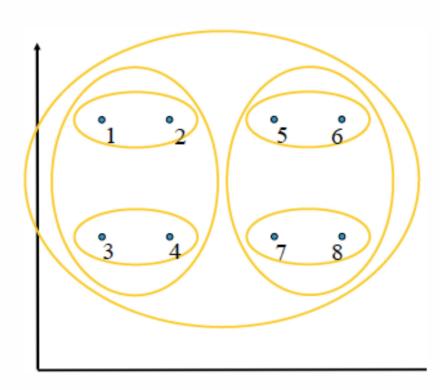
• Group average: the average similarity between groups

$$d_{GA} = \frac{1}{N_G N_H} \sum_{i \in G} \sum_{j \in H} d_{i,j}$$

Closest pair (single-link clustering)

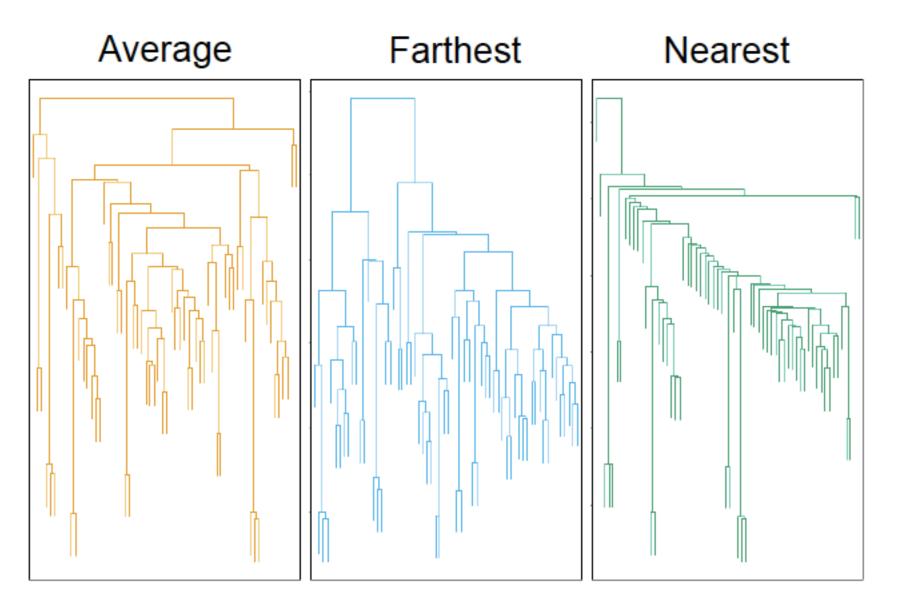


Farthest pair (complete-link clustering)



[Pictures from Thorsten Joachims]

Slide from Lu Wang's machine learning course



Slide from Lu Wang's machine learning course Mouse tumor data from [Hastie *et al.*]

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- Complete linkage has the opposite problem. It might not merge close groups because of outlier members that are far apart.
- Group average represents a natural compromise, but depends on the scale of the similarities. Applying a monotone transformation to the similarities can change the results.