



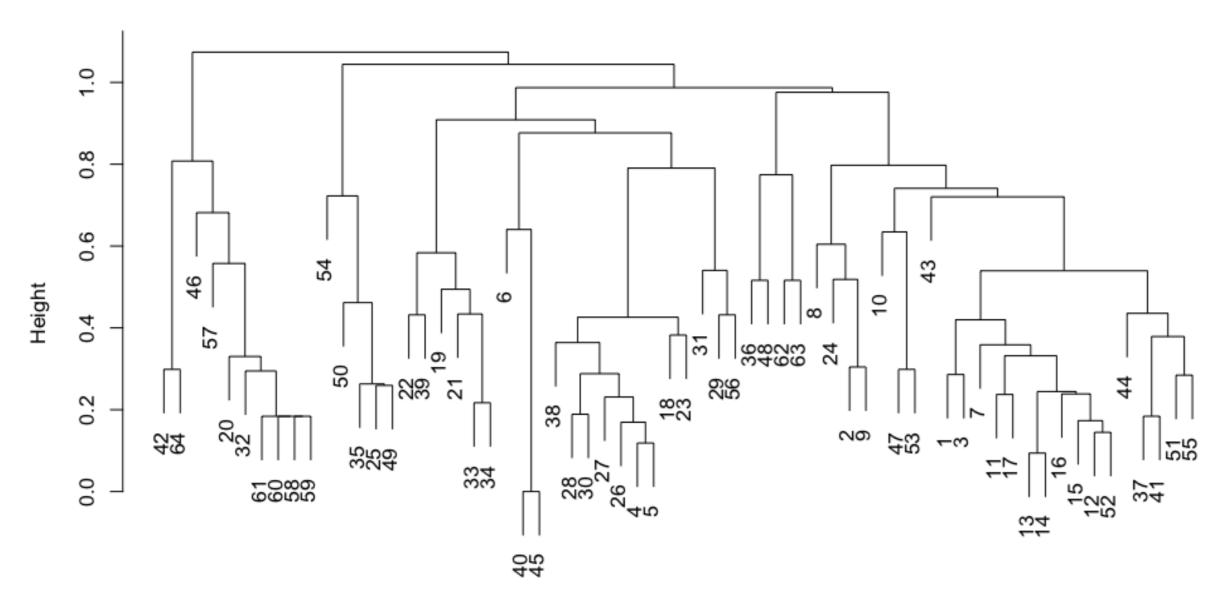
NETWORK ANALYSIS IN THE TIDYVERSE

Hierarchical Clustering

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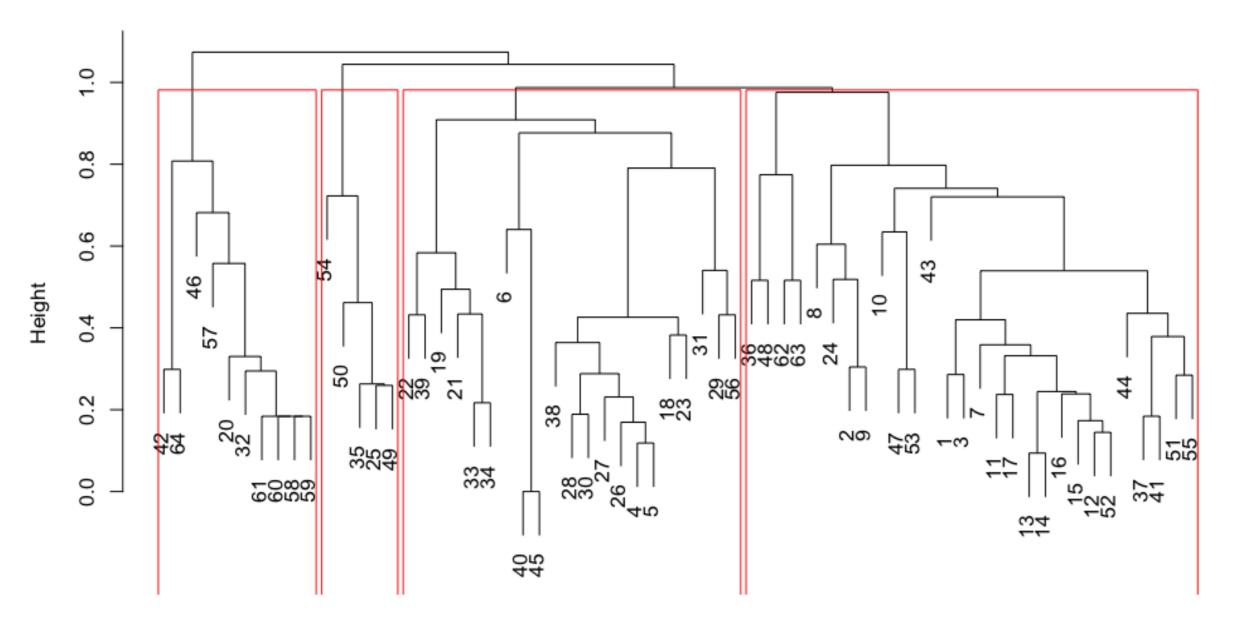
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Cluster Dendrogram



d hclust (*, "average")

Cluster Dendrogram



d hclust (*, "average")



The similarity measure

- **Single-linkage**: the similarity between two groups is the maximum of the similarities between nodes of different groups.
- Complete-linkage: the similarity between two groups is the minimum of the similarities between nodes of different groups.
- Average-linkage: the similarity between two groups is the average of the similarities between nodes of different groups.

The clustering algorithm

- 1. Evaluate the similarity measures for all node pairs.
- 2. Assign each node to a group of its own.
- 3. Find the pair of groups with the highest similarity and join them together into a single group.
- 4. Calculate the similarity between the new composite group and all others.
- 5. Repeat steps 3 and 4 until all nodes have been joined into a single group.



Hierarchical clustering in R

```
# distance matrix from similarity matrix
D <- 1-S

# distance object from distance matrix
d <- as.dist(D)

# average-linkage clustering method
cc <- hclust(d, method = "average")

# cut dendrogram at 4 clusters
hclust(d, method = "average")

[1] 1 1 1 2 2 2 1 1 1 1 1 1 1 1 1 1 2 2 3 2 2 2 1 4 2 2 2
[29] 2 2 2 3 2 2 4 1 1 2 2 2 1 3 1 1 2 3 1 1 4 4 1 1 1 4 1 2
[57] 3 3 3 3 3 3 1 1 3</pre>
```





Let's cluster our network!

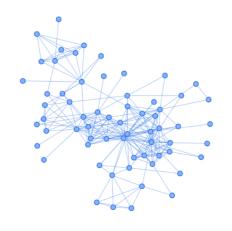


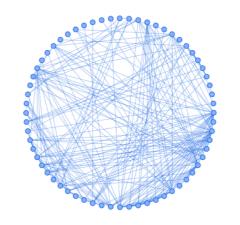


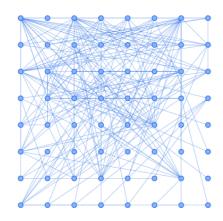
Interactive visualizations with visNetwork

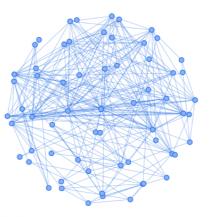


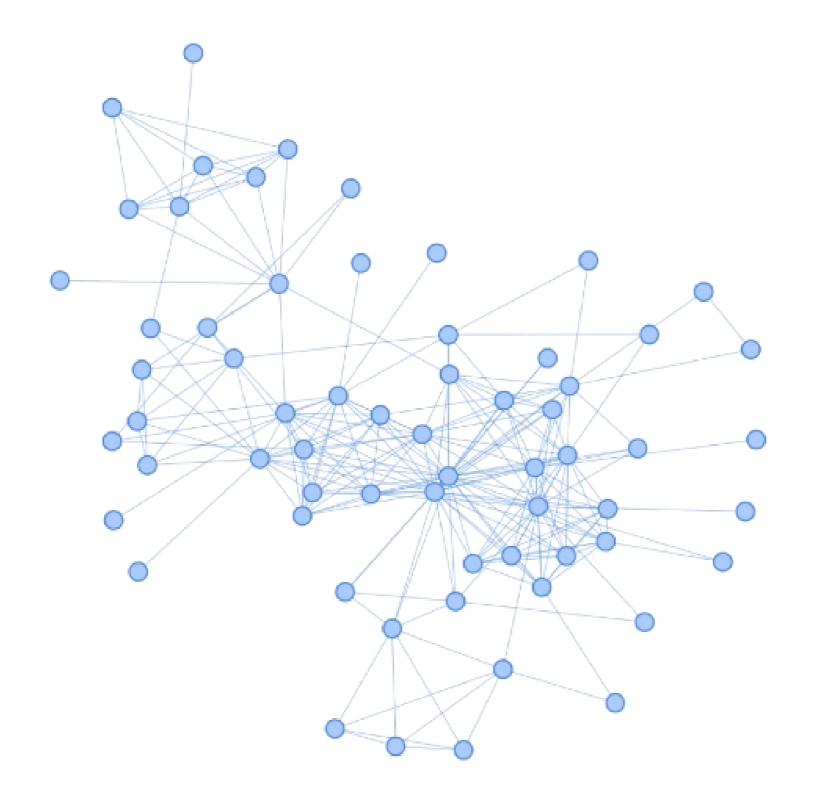
Different layouts



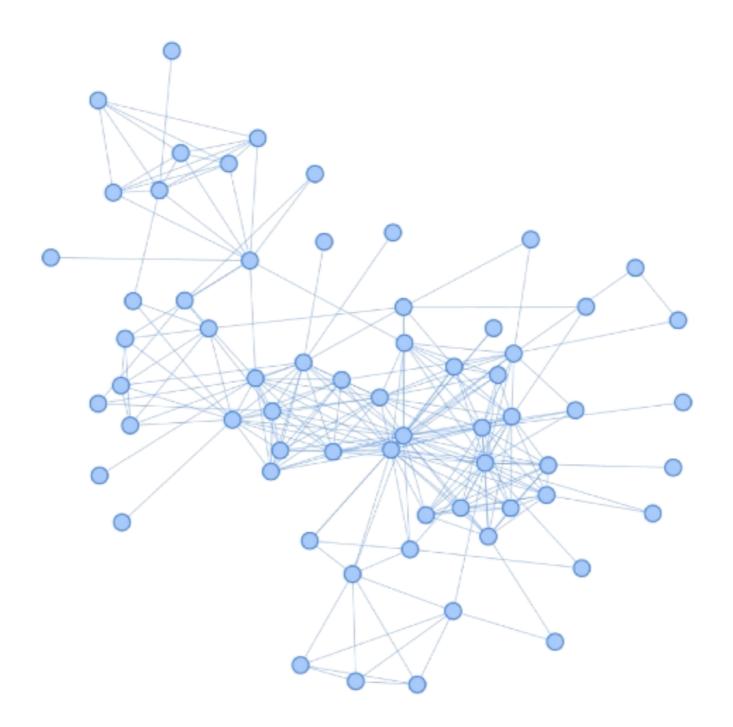






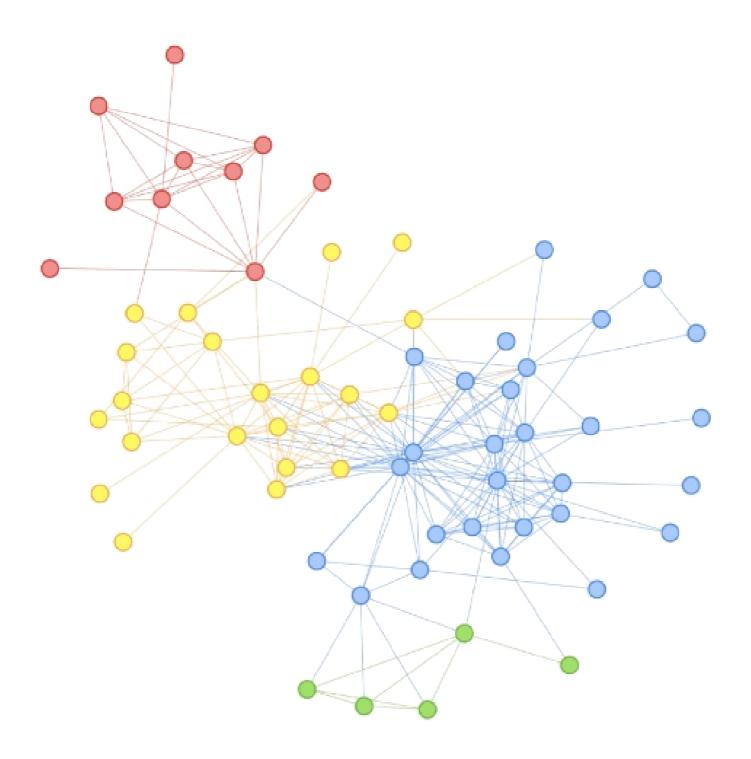


Select by id \$





Select by group \$







Let's interact!





Congratulations!



Deeper inside network science

You now know how to:

- Analyze any network with basic centrality and similarity measures
- Produce beautiful network visualizations, including interactive ones

For more information:

Univeristy of Udine Network Science Course





Continue the journey!