

# PsychoNet

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The goal of this assignment is to create a social network from your free recall data, to plot it and to evaluate it for smallworldness and centrality.

## Overview

This assignment contains of 3 steps.

1. Create a graph from the responses.
2. Plot the graph using `ggraph`.
3. Analyze it using `igraph`.

## Step I - Create social network

1. First, download the data using this [link](#) and store it inside your project. Then load the data using the following command (you probably don't need the `..`). Inspect the data. You will see that the object is composed of a list of vectors, with every vector representing the responses of one person. Which person the responses belong to is coded in the list's names. E.g., `free_recall$"Zana Hightower"` would give you the responses of Zana Hightower.

```
# load data
free_recall = readRDS("../1_data/psychonet_responses.RDS")
free_recall$"Zana Hightower"
```

```
## [1] "Juliana Lemus"      "Phylicia Belcher" "Cassy Martino"
## [4] "Sigrid March"      "Reena Place"      "Lourie Henke"
## [7] "Gerry Dolan"       "Velva Burley"     "Tona Timm"
## [10] "Jarvis Chapin"     "Deandre Talbert"  "Alysha Harwood"
## [13] "Lory Ralston"      "Luciano Aiken"    "Dee Bartholomew"
## [16] "Sharie Gable"      "Maudie Arroyo"    "Maisha Van"
```

2. Extract all of the respondents names using `names(free_recall)` and the unique responses using `unique(unlist(free_recall))`. Store these in objects named `respondents` and `responses` and create a third one containing the unique names across both, using `unique(c(respondents, responses))`, and name it `persons`.

```
# get names & persons
respondents = names(free_recall)
responses = unique(unlist(free_recall))
persons = unique(c(respondents, responses))
```

3. Now create an adjacency matrix with enough rows and columns to store the edges between individuals using `matrix(0, ncol = XX, nrow = XX)` (tipp: what is the `length()` of `persons`?) and name it `social_network`. Assign the matrix' `rownames()` and `colnames()` to be the names contained in `persons` (e.g., `rownames(XX) <- XX`).

```
# create social network
social_network = matrix(0, ncol = length(persons), nrow = length(persons))
colnames(social_network) = persons
rownames(social_network) = persons
```

4. Now comes the somewhat difficult part. Iterate over the free recall list and include an edge, if respondent *i* produced response *j*, i.e., set the cell *i*, *j* to 1. Do this using an outer-loop iterating over the respondents and an inner-loop iterating over the respondents' responses. For every respondent this means that you need to pull the respondent's responses and then iterate over those. Note that objects of type `list` and `matrix` can also be accessed using names. Here, we can make use of this by iterating directly over the names rather than their indices. See below.

```
# fill social network
for(i in respondents){

  # HERE EXTRACT RESPONSES OF RESPONDENT i
  responses_i = free_recall[[XX]]

  # loop over responses
  for(j in responses_i){

    # add edges
    social_network[XX, YY] = social_network[YY, XX] = ZZ

  }
}
```

```
# fill social network
for(i in respondents){

  # HERE EXTRACT RESPONSES OF RESPONDENT i
  responses_i = free_recall[[i]]

  # loop over responses
  for(j in responses_i){

    # add edges
    social_network[i, j] = social_network[j, i] = 1

  }
}
```

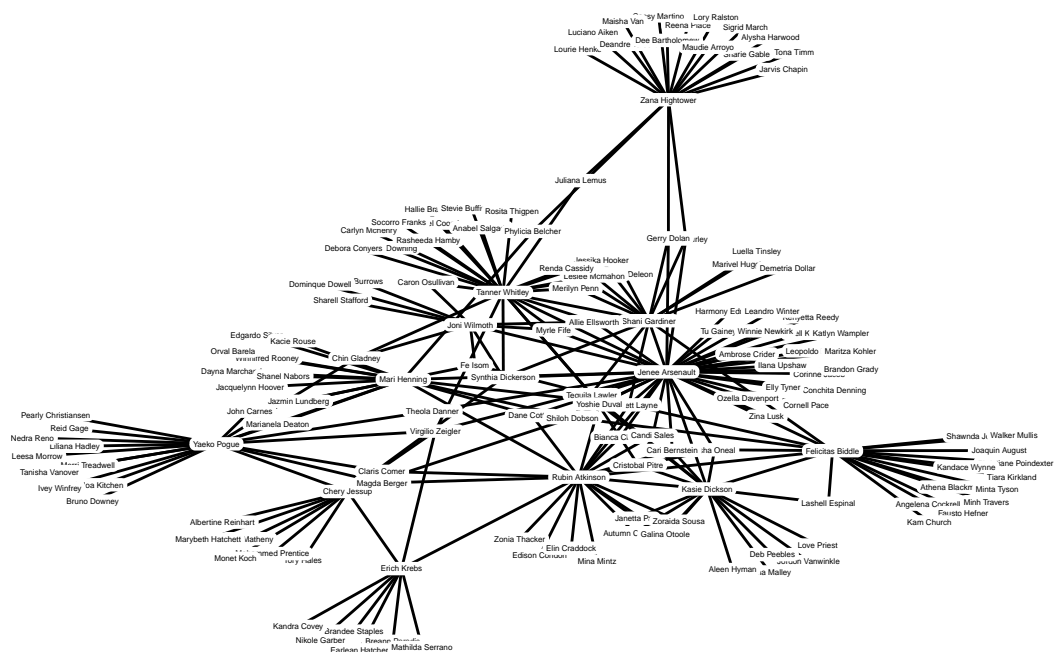
## Step II - Plot social network

1. The next step is to plot the network. A lot of plotting can be done using `igraph`, but `ggraph` clearly creates nicer plots (and has nicer syntax). Install the package using `install.packages('ggraph')` and check out the **Intro**. Then plot the network using the code below. If the graph and the labels appear too small, increase the numbers in `geom_node_label()`.

```
require(igraph)
require(ggraph)
```

```
# create social graph
social_graph <- graph_from_adjacency_matrix(social_network, mode = 'undirected')

# plot
ggraph(social_graph) +
  geom_edge_link() +
  geom_node_label(aes(label = V(social_graph)$name),
    repel = FALSE,
    label.size = unit(0.15, "lines"),
    label.padding = unit(.1, "lines"),
    size = 1.2) +
  theme_graph(base_family = "Roboto Condensed")
```



### Step III - Analyze social network

1. Is your social network a *small world*? To evaluate this, calculate its clustering coefficient (using `transitivity(XX, type = 'localaverage')`) and average shortest path length (using `average.path.length(XX)`). What do you think, is it small world?

```
# calculate statistics
transitivity(social_graph, type = 'localaverage')
```

```
## [1] 0.4980782
```

```
average.path.length(social_graph)
```

```
## [1] 3.323176
```

2. Identify central individuals. Use the functions `centr_degree()`, `centr_clo()`, `centr_betw()`, and `page_rank()` to calculate the centrality with regard to the respective definition for every person (i.e.,

node). Then identify the, respectively, most central person using `persons[which.max(XX)]`. Note, each of the four functions returns a list. This means that you first need to extract the vector containing the centrality values (usually called `res` or `vector`).

```
# get centralities
deg = centr_degree(social_graph)$res
clo = centr_clo(social_graph)$res
bet = centr_betw(social_graph)$res
pgr = page_rank(social_graph)$vector

# identify most central person
persons[which.max(deg)]
```

```
## [1] "Jenee Arsenault"
```

```
persons[which.max(clo)]
```

```
## [1] "Jenee Arsenault"
```

```
persons[which.max(bet)]
```

```
## [1] "Jenee Arsenault"
```

```
persons[which.max(pgr)]
```

```
## [1] "Jenee Arsenault"
```

3. BONUS: If you are interested in finding communities you can play around with `cluster_louvain()` and `cluster_optimal()` from the `igraph` packages.