

ETC5510: Introduction to Data Analysis

Week 9, part A

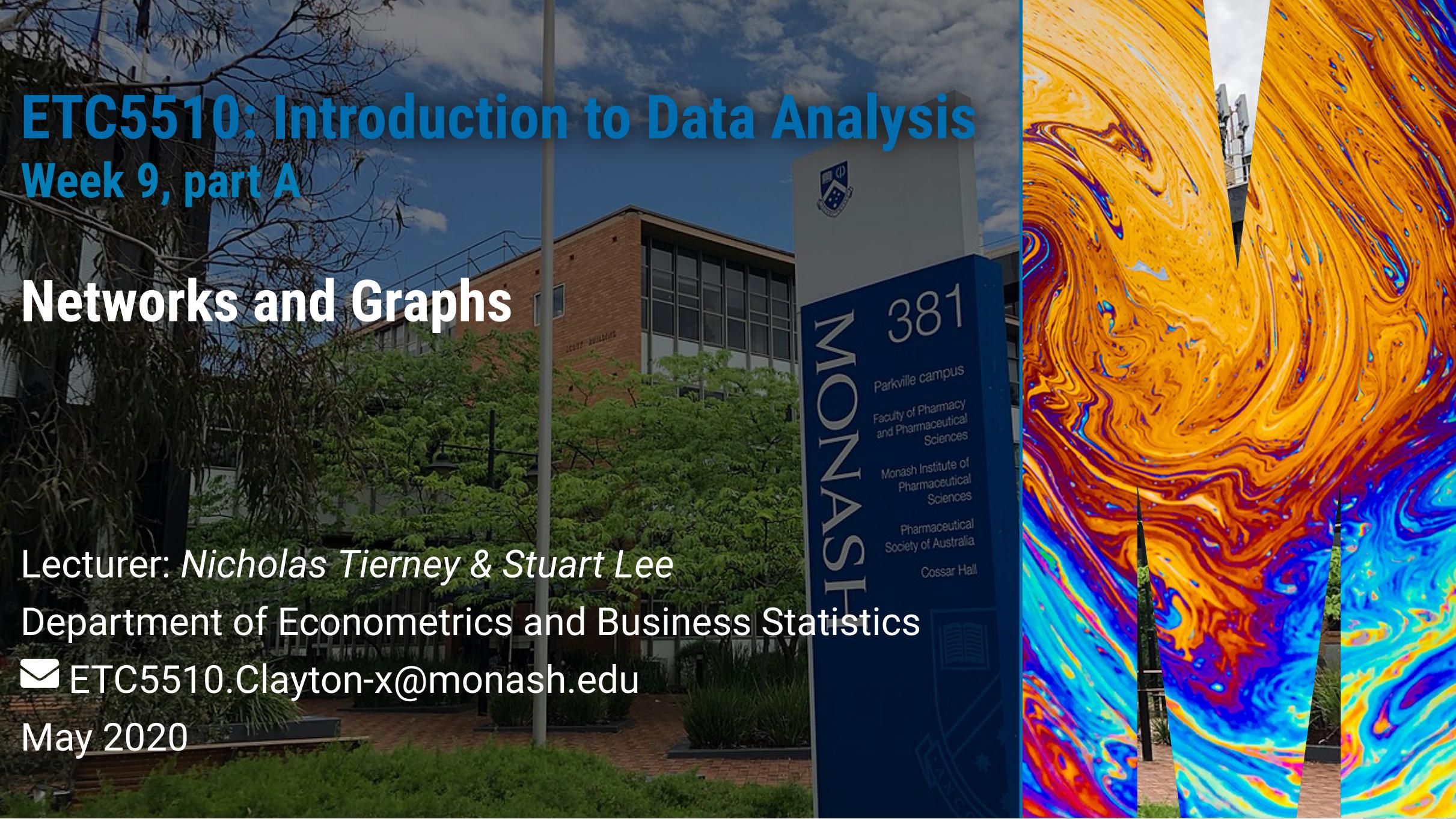
Networks and Graphs

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Announcements

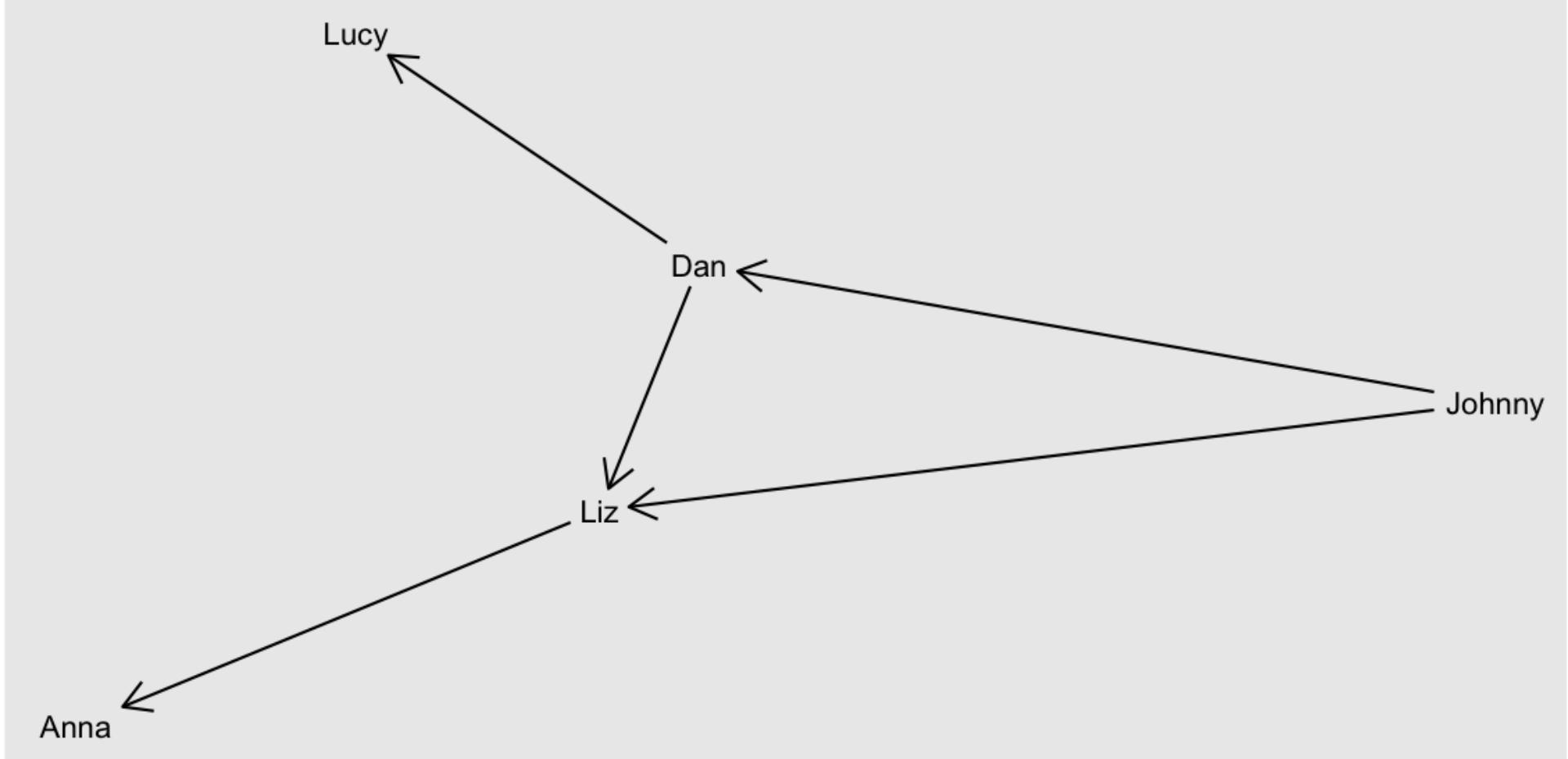
- Assignment 2: due 20th May at 5pm
- Project deadlines:
 - **Milestone 2 (20th May)** : Finalised team/team members, a paragraph about what you're going to do and links to your data sources.
 - **Milestone 3 (27th May)** : Upload data, and Rmd file, and html describing your data, the cleaning you've done and what's left.

Network analysis

A description of phone calls

- Johnny → Liz
- Liz → Anna
- Johnny → Dan
- Dan → Liz
- Dan → Lucy

As a graph



And as an association matrix

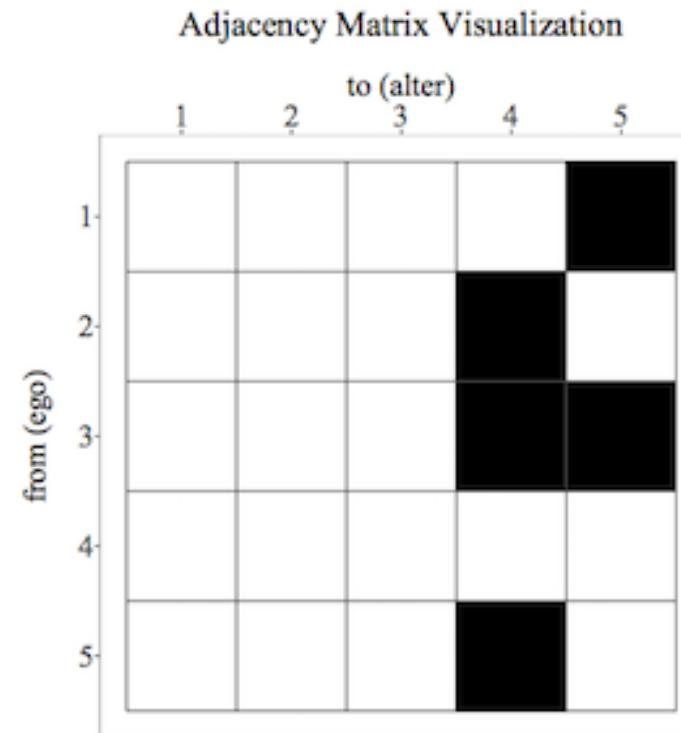
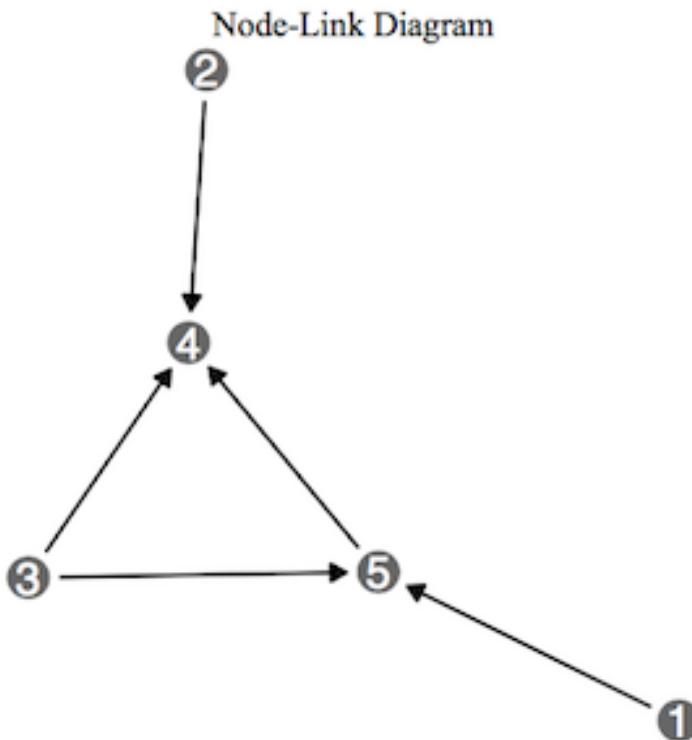
```
##          Johnny Dan Liz Lucy Anna
## Johnny      0   1   1   0   0
## Dan         0   0   1   1   0
## Liz         0   0   0   0   1
## Lucy        0   0   0   0   0
## Anna        0   0   0   0   0
```

Why care about these relationships?

- **Telephone exchanges:** Nodes are the phone numbers. Edges would indicate a call was made between two numbers.
- **Book or movie plots:** Nodes are the characters. Edges would indicate whether they appear together in a scene, or chapter. If they speak to each other, various ways we might measure the association.
- **Social media:** nodes would be the people who post on facebook, including comments. Edges would measure who comments on who's posts.

Drawing these relationships out:

One way to describe these relationships is to provide association matrix between many objects.



(Image created by Sam Tyner.)

Example: Madmen



Source: [wikicommons](#)

Generate a network view

- Create a layout (in 2D) which places nodes which are most related close,
- Plot the nodes as points, connect the appropriate lines
- Overlaying other aspects, e.g. gender

introducing madmen data

```
glimpse(madmen)
## List of 2
## $ edges   :'data.frame':   39 obs. of  2 variables:
##   ..$ Name1: Factor w/ 9 levels "Betty Draper",...: 1 1 2 2 2 2 2 2 2 ...
##   ..$ Name2: Factor w/ 39 levels "Abe Drexler",...: 15 31 2 4 5 6 8 9 11 21 ...
## $ vertices:'data.frame':   45 obs. of  2 variables:
##   ..$ label : Factor w/ 45 levels "Abe Drexler",...: 5 9 16 23 26 32 33 38 39 17 ...
##   ..$ Gender: Factor w/ 2 levels "female","male": 1 2 2 1 2 1 2 2 2 2 ...
```

Nodes and edges?

Network data can be thought of as two related tables, **nodes** and **edges**:

- **nodes** are connection points
- **edges** are the connections between points

Example: Mad Men. (Nodes = characters from the series)

```
madmen_nodes
## # A tibble: 45 x 2
##   label      gender
##   <chr>     <chr>
## 1 Betty Draper  female
## 2 Don Draper   male
## 3 Harry Crane  male
## 4 Joan Holloway female
## 5 Lane Pryce   male
## 6 Peggy Olson  female
## 7 Pete Campbell male
## 8 Roger Sterling male
## 9 Sal Romano   male
## 10 Henry Francis male
## # ... with 35 more rows
```

Example: Mad Men. (Edges = how they are associated)

```
madmen_edges
## # A tibble: 39 x 2
##   Name1      Name2
##   <chr>     <chr>
## 1 Betty Draper Henry Francis
## 2 Betty Draper Random guy
## 3 Don Draper Allison
## 4 Don Draper Bethany Van Nuys
## 5 Don Draper Betty Draper
## 6 Don Draper Bobbie Barrett
## 7 Don Draper Candace
## 8 Don Draper Doris
## 9 Don Draper Faye Miller
## 10 Don Draper Joy
## # ... with 29 more rows
```

Let's get the madmen data into the right shape

```
madmen_edges %>%  
  rename(from_id = Name1, to_id = Name2)  
## # A tibble: 39 x 2  
##   from_id     to_id  
##   <chr>      <chr>  
## 1 Betty Draper Henry Francis  
## 2 Betty Draper Random guy  
## 3 Don Draper Allison  
## 4 Don Draper Bethany Van Nuys  
## 5 Don Draper Betty Draper  
## 6 Don Draper Bobbie Barrett  
## 7 Don Draper Candace  
## 8 Don Draper Doris  
## 9 Don Draper Faye Miller  
## 10 Don Draper Joy  
## # ... with 29 more rows
```

Let's get the madmen data into the right shape

```
madmen_net <- madmen_edges %>%  
  rename(from_id = Name1, to_id = Name2) %>%  
  full_join(madmen_nodes,  
            by = c("from_id" = "label"))
```

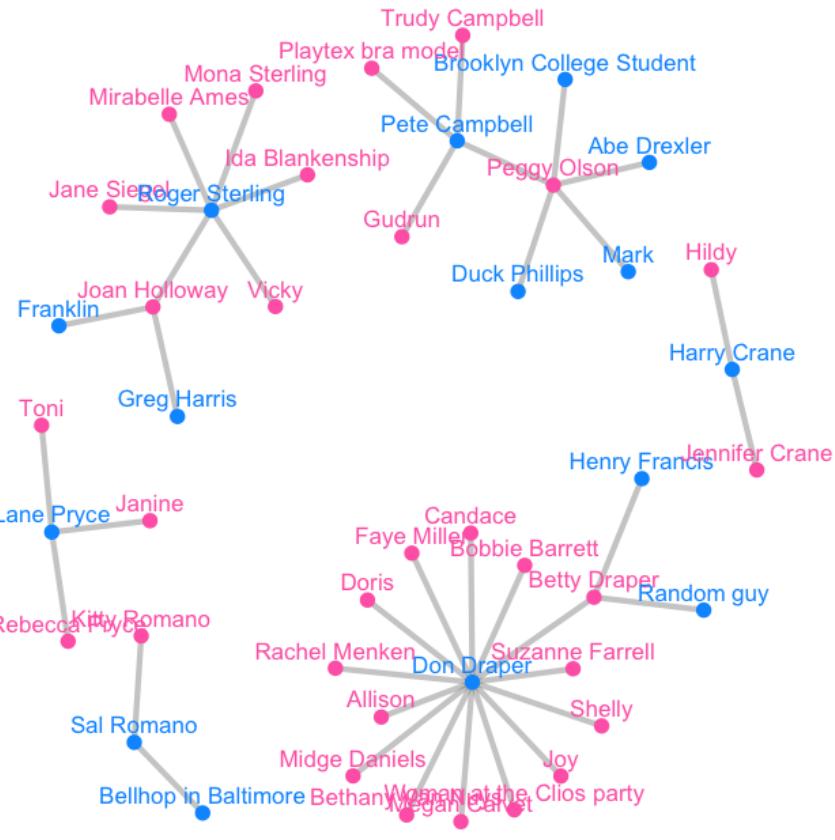
```
madmen_net  
## # A tibble: 75 x 3  
##   from_id      to_id    gender  
##   <chr>        <chr>    <chr>  
## 1 Betty Draper Henry Francis  female  
## 2 Betty Draper Random guy  female  
## 3 Don Draper  Allison    male  
## 4 Don Draper  Bethany Van Nuys male  
## 5 Don Draper  Betty Draper male  
## 6 Don Draper  Bobbie Barrett male  
## 7 Don Draper  Candace    male  
## 8 Don Draper  Doris     male  
## 9 Don Draper  Faye Miller male  
## 10 Don Draper Joy       male  
## # ... with 65 more rows
```

Full join?

`full_join(x, y)`

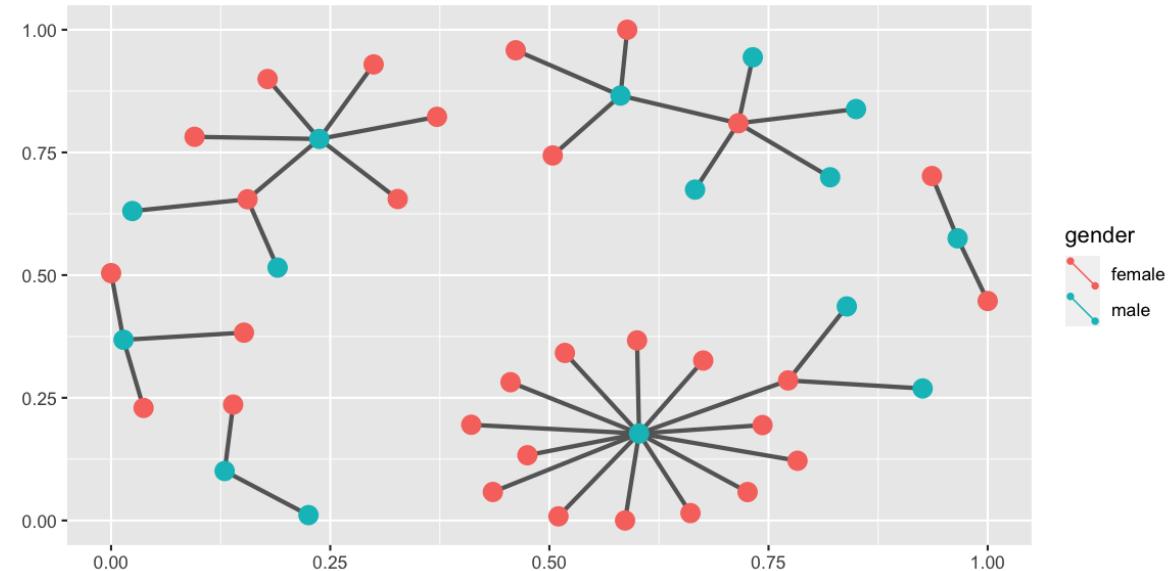
1	x1	1	y1
2	x2	2	y2
3	x3	4	y4

Plotting the data



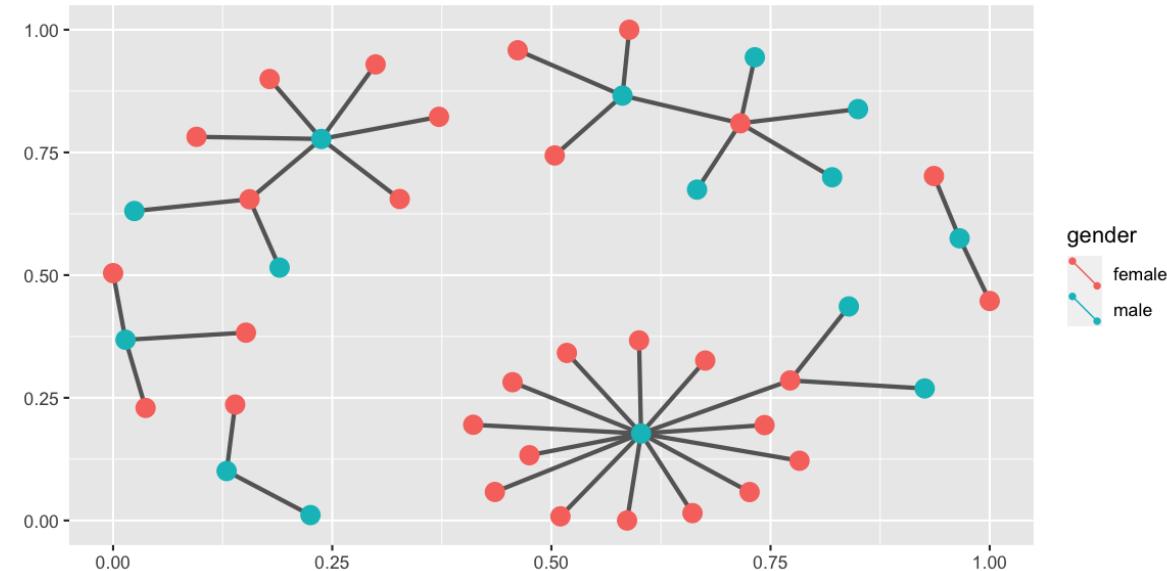
How to plot

```
set.seed(5556677)
ggplot(data = madmen_net,
       aes(from_id = from_id,
           to_id = to_id)) +
  geom_net(aes(colour = gender))
```



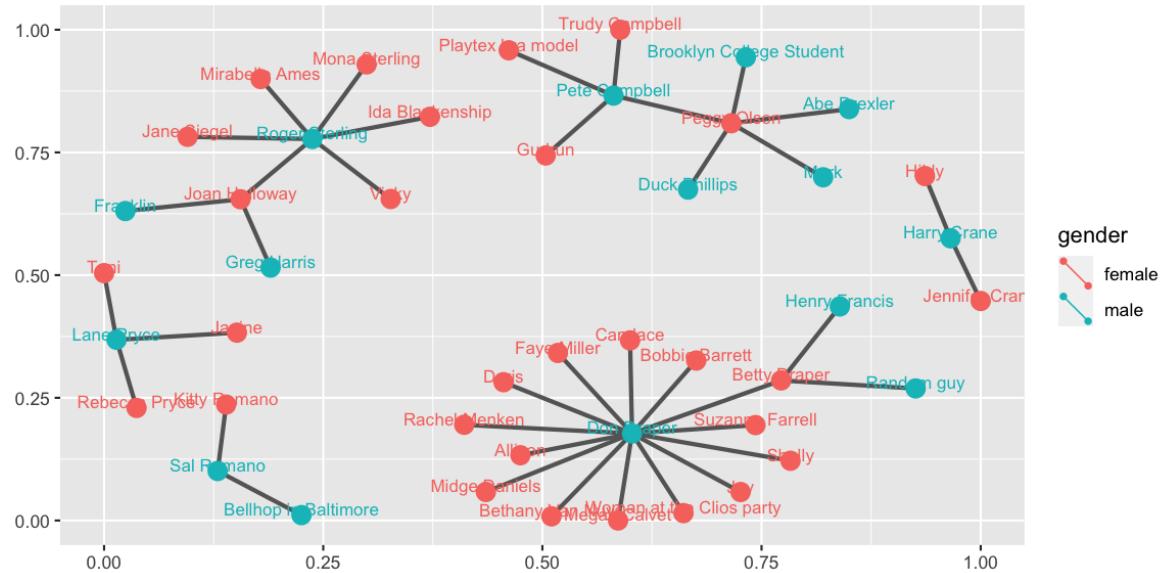
How to plot: specify the layout algorithm

```
set.seed(5556677)
ggplot(data = madmen_net,
       aes(from_id = from_id,
           to_id = to_id)) +
  geom_net(aes(colour = gender),
           layout.alg = "kamada_kawai")
```



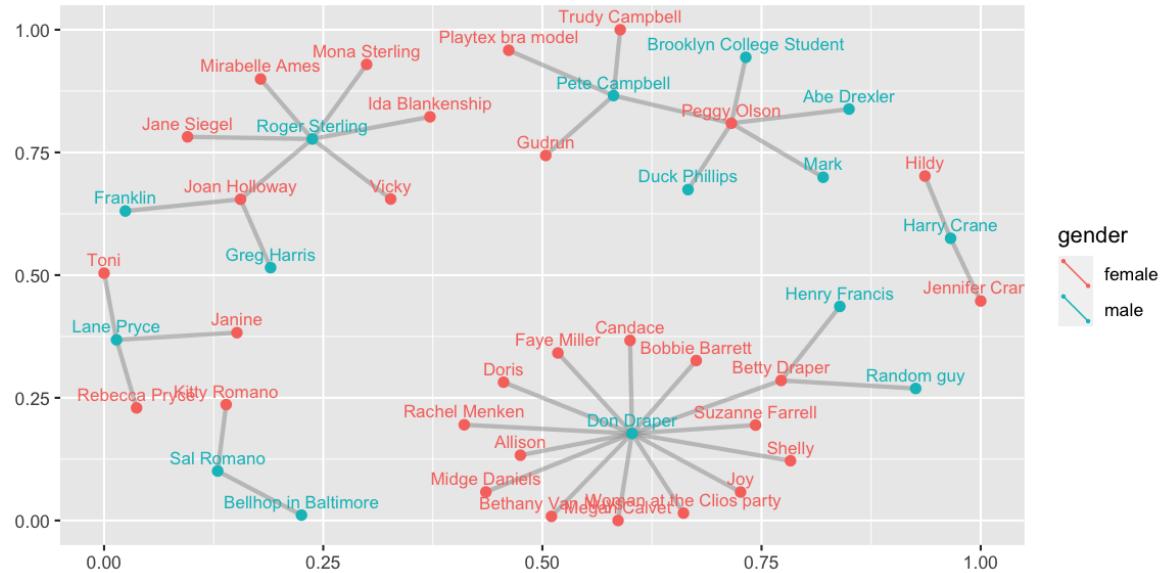
How to plot

```
set.seed(5556677)
ggplot(data = madmen_net,
       aes(from_id = from_id,
           to_id = to_id)) +
  geom_net(aes(colour = gender),
            layout.alg = "kamada",
            directed = FALSE,
            labelon = TRUE,
            fontsize = 3)
```



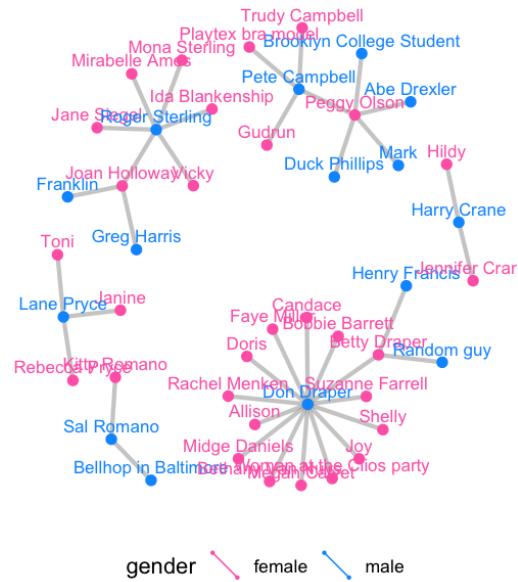
How to plot

```
set.seed(5556677)
ggplot(data = madmen_net,
       aes(from_id = from_id,
           to_id = to_id)) +
  geom_net(aes(colour = gender),
           layout.alg = "kamada",
           directed = FALSE,
           labelon = TRUE,
           fontsize = 3,
           size = 2,
           vjust = -0.6,
           ecolour = "grey60",
           ealpha = 0.5)
```



How to plot

```
set.seed(5556677)
ggplot(data = madmen_net,
       aes(from_id = from_id,
           to_id = to_id)) +
  geom_net(aes(colour = gender),
            layout.alg = "kamada",
            directed = FALSE,
            labelon = TRUE,
            fontsize = 3,
            size = 2,
            vjust = -0.6,
            ecolour = "grey60",
            ealpha = 0.5) +
  scale_colour_manual(values =
    lims(x = c(-0.05, 1.05)) +
  theme_net() +
  theme(legend.position = "bot
```



gender ↗ female ↖ male

Which character was most connected?

```
madmen_edges %>%  
  pivot_longer(cols = c(Name1, Name2),  
               names_to = "List",  
               values_to = "Name")  
## # A tibble: 78 x 2  
##   List   Name  
##   <chr> <chr>  
## 1 Name1 Betty Draper  
## 2 Name2 Henry Francis  
## 3 Name1 Betty Draper  
## 4 Name2 Random guy  
## 5 Name1 Don Draper  
## 6 Name2 Allison  
## 7 Name1 Don Draper  
## 8 Name2 Bethany Van Nuys  
## 9 Name1 Don Draper  
## 10 Name2 Betty Draper  
## # ... with 68 more rows
```

Which character was most connected?

```
madmen_edges %>%  
  pivot_longer(cols = c(Name1, Name2),  
               names_to = "List",  
               values_to = "Name") %>%  
  count(Name, sort = TRUE)  
## # A tibble: 45 x 2  
##   Name           n  
##   <chr>        <int>  
## 1 Don Draper     14  
## 2 Roger Sterling    6  
## 3 Peggy Olson      5  
## 4 Pete Campbell     4  
## 5 Betty Draper      3  
## 6 Joan Holloway     3  
## 7 Lane Pryce        3  
## 8 Harry Crane       2  
## 9 Sal Romano        2  
## 10 Abe Drexler       1  
## # ... with 35 more rows
```

Example: American college football

Early American football outfits were like Australian AFL today!



Source: [wikicommons](#)

Example: American college football

Fall 2000 Season of Division I college football.

- Nodes are the teams, edges are the matches.
- Teams are broken into "conferences" which are the primary competition, but they can play outside this group.

American college football data

Edges:

```
football$edges
```

```
##          from
## 1    BrighamYoung FloridaState
## 2        Iowa           Kansas
## 3    BrighamYoung           K
## 4        NewMexico
## 5    KansasState
## 6        Iowa
## 7    PennState SouthernCalifornia
## 8 SouthernCalifornia ArizonaState
## 9        ArizonaState
## 10   BrighamYoung           San
## 11   NewMexico           San
## 12   TexasTech
## 13        Baylor
## 14   TexasTech
## 15   KansasState
## 16 NorthernIllinois
```

Nodes:

```
football$vertices
```

```
##          from
## 1    BrighamYoung
## 2    FloridaState
## 3        Iowa
## 4    KansasState
## 5        NewMexico
## 6    TexasTech
## 7    PennState
## 8 SouthernCalifornia
## 9    ArizonaState
## 10   SanDiegoState
## 11   Baylor
## 12   NorthTexas
## 13 NorthernIllinois
## 14   Northwestern
```

American college football: joining the data

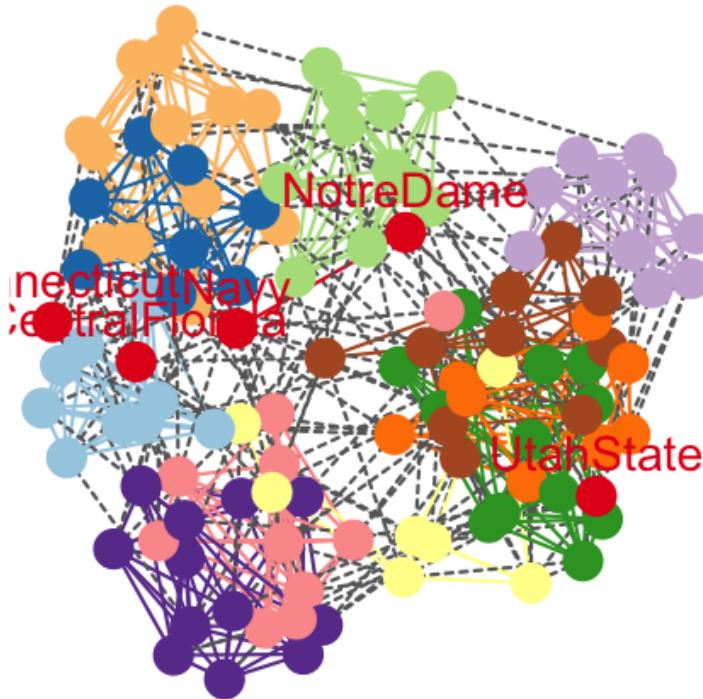
```
# data step: merge vertices and edges
ftnet <- full_join(
  football$edges, football$vertices,
  by = c("from" = "label"))
) %>%
  mutate(schools = if_else(value == "Independents", from, ""))
```

ftnet

##	from	to	same.conf	value
## 1	BrighamYoung	FloridaState	0	Mountain West
## 2	Iowa	KansasState	0	Big Ten
## 3	BrighamYoung	NewMexico	1	Mountain West
## 4	NewMexico	TexasTech	0	Mountain West
## 5	KansasState	TexasTech	1	Big Twelve
## 6	Iowa	PennState	1	Big Ten
## 7	PennState	SouthernCalifornia	0	Big Ten
## 8	SouthernCalifornia	ArizonaState	1	Pacific Ten
## 9	ArizonaState	SanDiegoState	0	Pacific Ten
## 10	BrighamYoung	SanDiegoState	1	Mountain West
## 11	NewMexico	SanDiegoState	1	Mountain West

Example: American college football

```
ggplot(data = ftnet,
       aes(from_id = from, to_id = to)) +
  geom_net(
    aes(colour = value,
        group = value,
        linetype = factor(1-same.conf),
        label = schools),
    linewidth = 0.5,
    size = 5,
    vjust = -0.75,
    alpha = 0.3,
    layout.alg = 'fruchtermanreingold'
  ) +
  theme_net() +
  theme(legend.position = "bottom") +
  scale_colour_brewer("Conference", palette = "Paired")
```



$\text{factor}(1 - \text{same.conf})$



What do we learn?

- Remember layout is done to place nodes that are more similar close together in the display.
- The colours indicate conference the team belongs too. For the most part, conferences are clustered, more similar to each other than other conferences.
- There are some clusters of conference groups, eg Mid-American, Big East, and Atlantic Coast
- The Independents are independent
- Some teams play far afield from their conference.

Example: Harry Potter characters



Source: [wikicommons](#)

Example: Harry Potter characters

There is a connection between two students if one provides emotional support to the other at some point in the book.

- Code to pull the data together is provided by Sam Tyner [here](#).

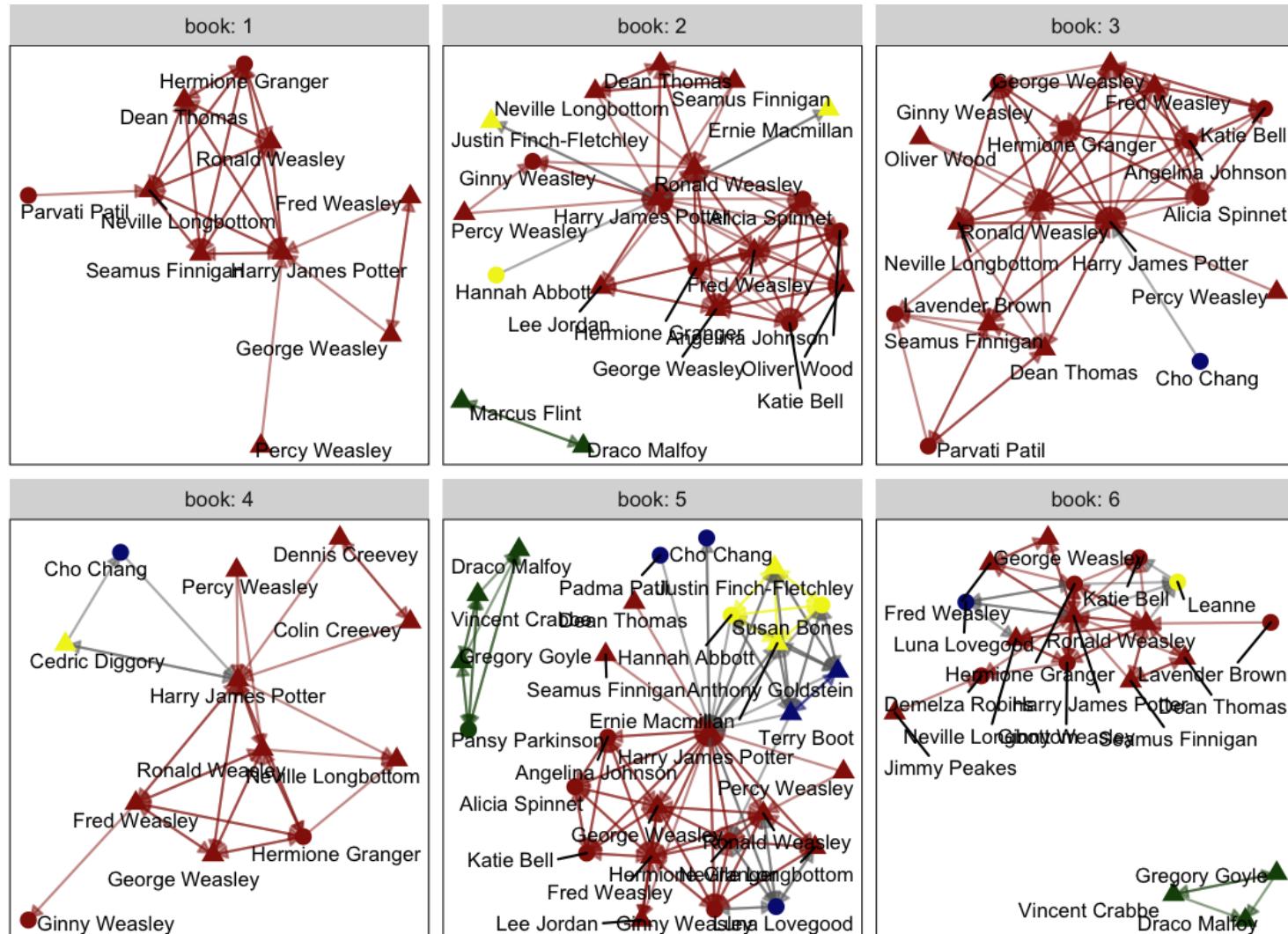
Harry potter data as nodes and edges

```
hp_all
## # A tibble: 720 x 6
##   book  from_id          to_id      schoolyear gender house
##   <chr> <chr>           <chr>       <dbl> <chr> <chr>
## 1 1     Dean Thomas     Harry James Potter  1991 M    Gryffindor
## 2 1     Dean Thomas     Hermione Granger  1991 M    Gryffindor
## 3 1     Dean Thomas     Neville Longbottom 1991 M    Gryffindor
## 4 1     Dean Thomas     Ronald Weasley    1991 M    Gryffindor
## 5 1     Dean Thomas     Seamus Finnigan  1991 M    Gryffindor
## 6 1     Fred Weasley    George Weasley   1989 M    Gryffindor
## 7 1     Fred Weasley    Harry James Potter  1989 M    Gryffindor
## 8 1     George Weasley  Fred Weasley    1989 M    Gryffindor
## 9 1     George Weasley  Harry James Potter  1989 M    Gryffindor
## 10 1    Harry James Potter Dean Thomas    1991 M    Gryffindor
## # ... with 710 more rows
```

Let's plot the characters

```
ggplot(data = hp_all,  
       aes(from_id = from_id,  
           to_id = to_id)) +  
  geom_net(aes(colour = house, group = house, shape = gender),  
           fiteach=T,  
           directed = T,  
           size = 3,  
           linewidth = .5,  
           ealpha = .5,  
           labelon = T,  
           fontsize = 3,  
           repel = T,  
           labelcolour = "black",  
           arrowsize = .5,  
           singletons = FALSE) +  
  scale_colour_manual(values = c("#941B08", "#F1F31C", "#071A80", "#154C07")) +  
  facet_wrap(~book, labeller = "label_both", ncol=3) +  
  theme_net() +  
  theme(panel.background = element_rect(colour = 'black'),  
        legend.position="bottom")
```

Let's plot the characters



house

Gryffindor

Hufflepuff

Ravenclaw

Slytherin

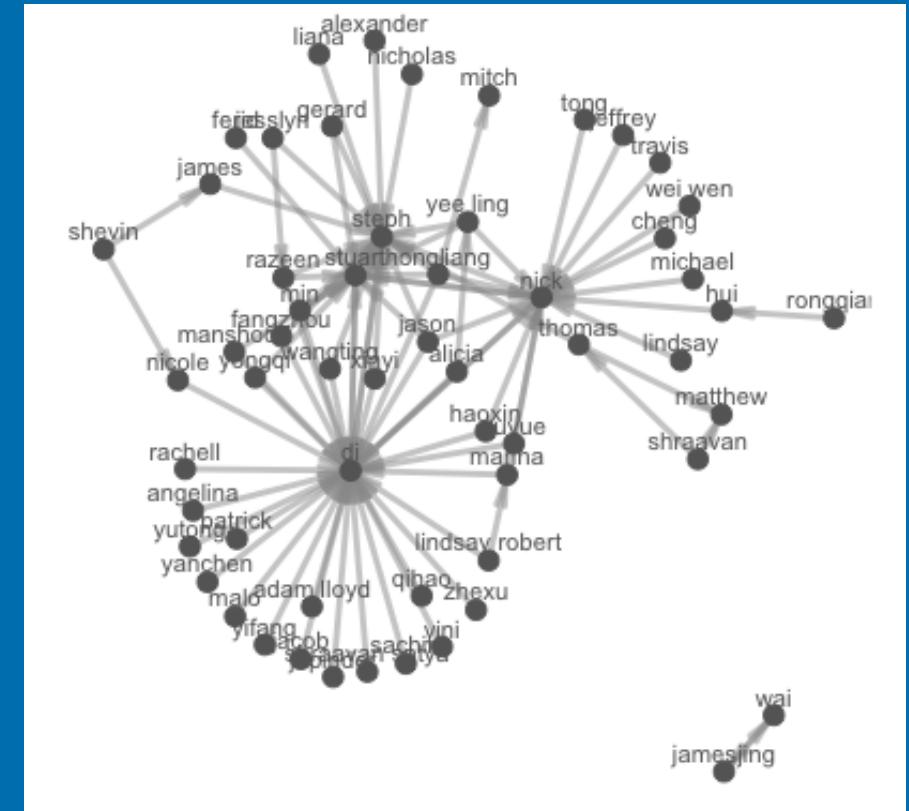
gender

F

M

Your turn: lab exercise

- Read in last semesters class data, which contains s1_name and s2_name are the first names of class members, and tutors, with the latter being the "go-to" person for the former.
- Write the code to produce a class network that looks something like below

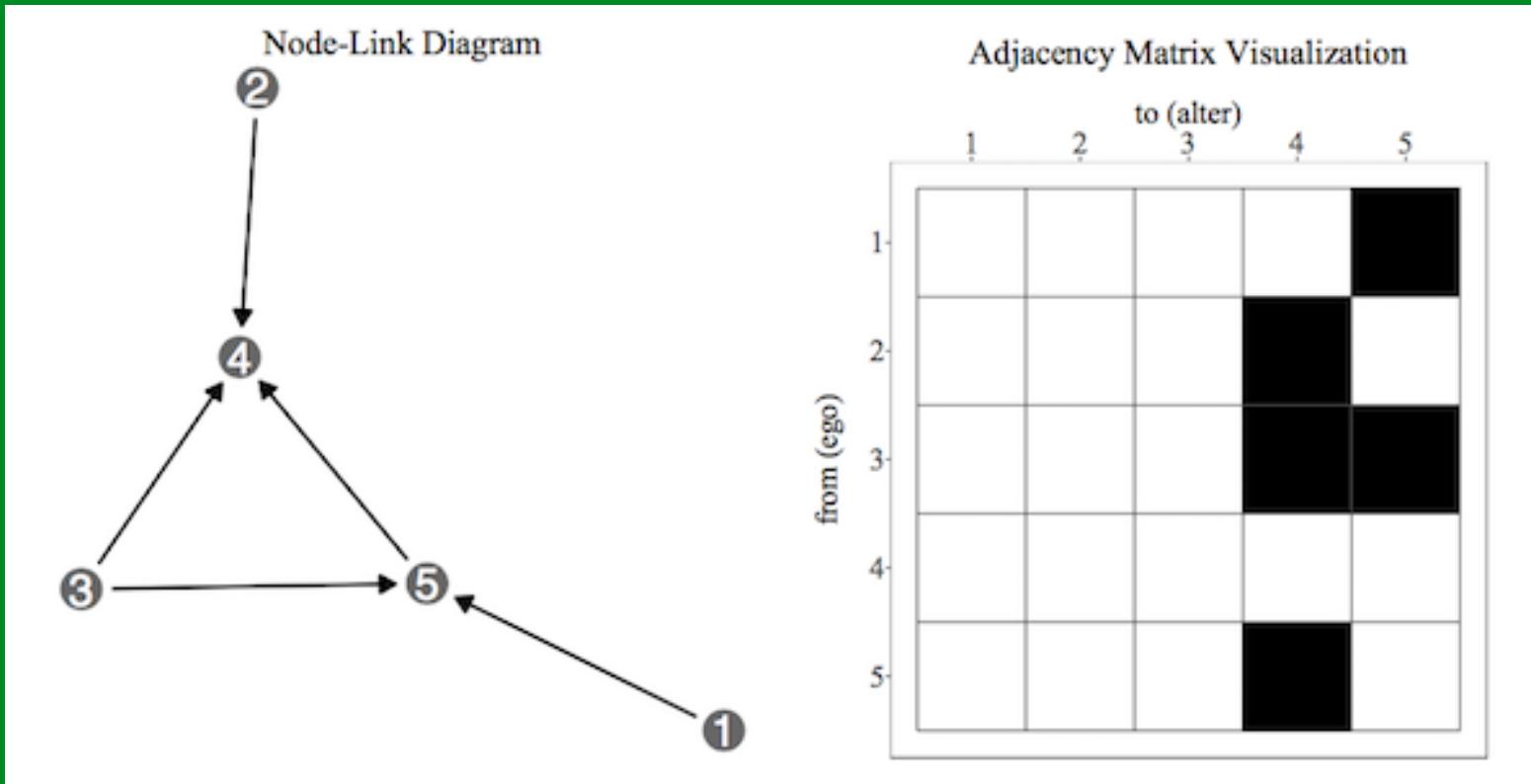


Summary

- To make a network analysis, you need:
 - an association matrix, that describes how nodes (vertices) are connected to each other
 - a layout algorithm to place the nodes optimally so that the fewest edges cross, or that the nodes that are most closely associated are near to each other.

Quantitative association matrices

Previous association matrices were black and white:



Quantitative association matrices

- You could have the association between nodes described as real numbers.
- E.g., these are the number of times that these people called each other in the last week:

	Meg	Tay	Yat	Zili	Jess
Meg	0	5	4	1	1
Tay	5	0	4	2	1
Yat	4	4	0	0	0
Zili	1	2	0	0	6
Jess	1	1	0	6	0

Quantitative association matrices

We would need to turn this into an edge data set:

```
## # A tibble: 25 x 3
##   from   to   count
##   <chr> <chr> <dbl>
## 1 Meg    Meg     0
## 2 Tay    Meg     5
## 3 Yat    Meg     4
## 4 Zili   Meg     1
## 5 Jess   Meg     1
## 6 Meg    Tay     5
## 7 Tay    Tay     0
## 8 Yat    Tay     4
## 9 Zili   Tay     2
## 10 Jess   Tay    1
## # ... with 15 more rows
```

Quantitative association matrices

- We need to decide what corresponds to a "connection".
- Let's say they need to have called each other at least 4 times, to be considered connected.

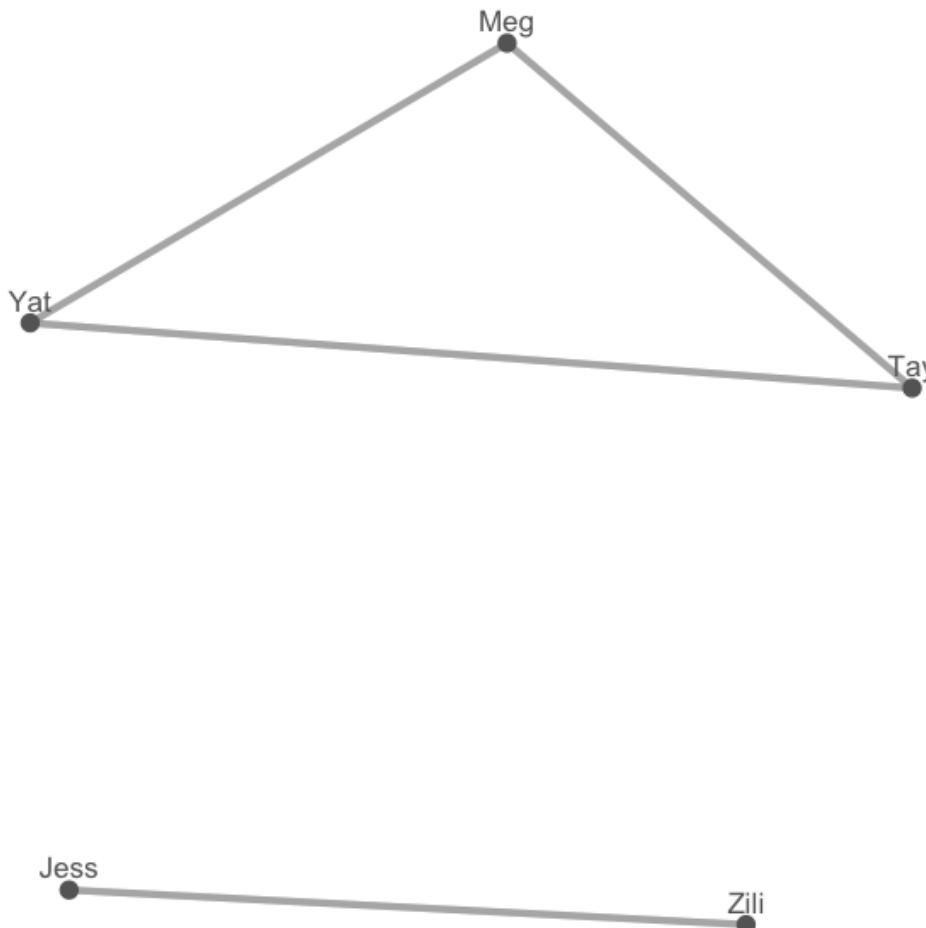
```
d_edges_filter <- d_edges %>% filter(count > 3)

d_edges_filter
## # A tibble: 8 x 3
##   from     to   count
##   <chr>  <chr> <dbl>
## 1 Tay    Meg      5
## 2 Yat    Meg      4
## 3 Meg    Tay      5
## 4 Yat    Tay      4
## 5 Meg    Yat      4
## 6 Tay    Yat      4
## 7 Jess   Zili     6
## 8 Zili   Jess     6
```

Association matrices: Make the network diagram.

```
library(geomnet)
set.seed(2019-10-09)
ggplot(data = d_edges_filter,
       aes(
         from_id = from,
         to_id = to)) +
  geom_net(
    layout.alg = "kamadakawai",
    size = 2,
    labelon = TRUE,
    vjust = -0.6,
    ecolour = "grey60",
    directed = FALSE,
    fontsize = 3,
    ealpha = 0.5
  ) +
  theme_net()
```

Association matrices: Make the network diagram.



Data: Last 2018, 4 months of currency USD cross-rates

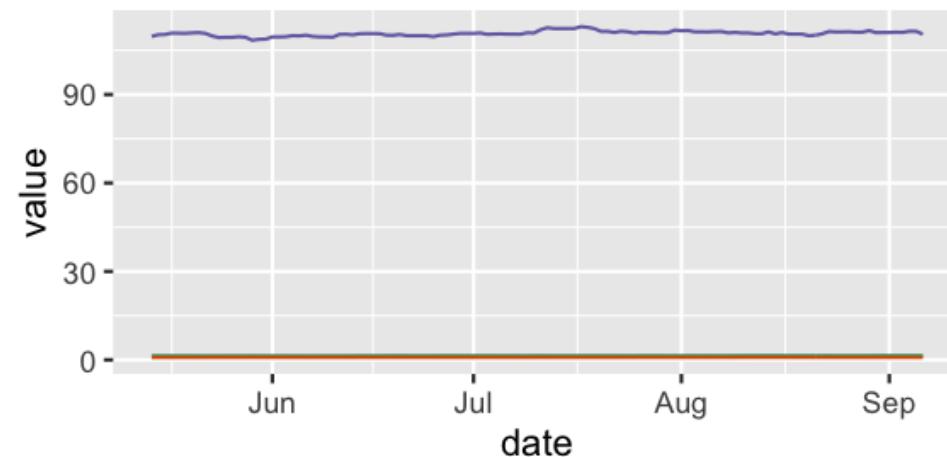
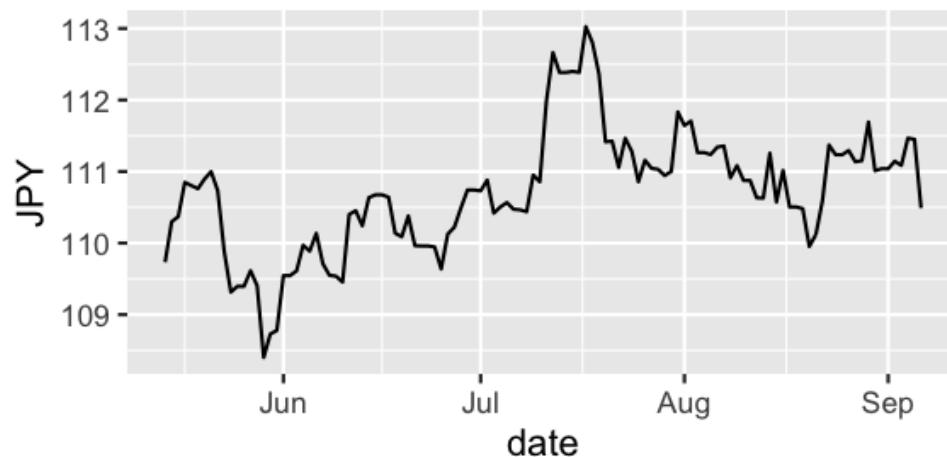
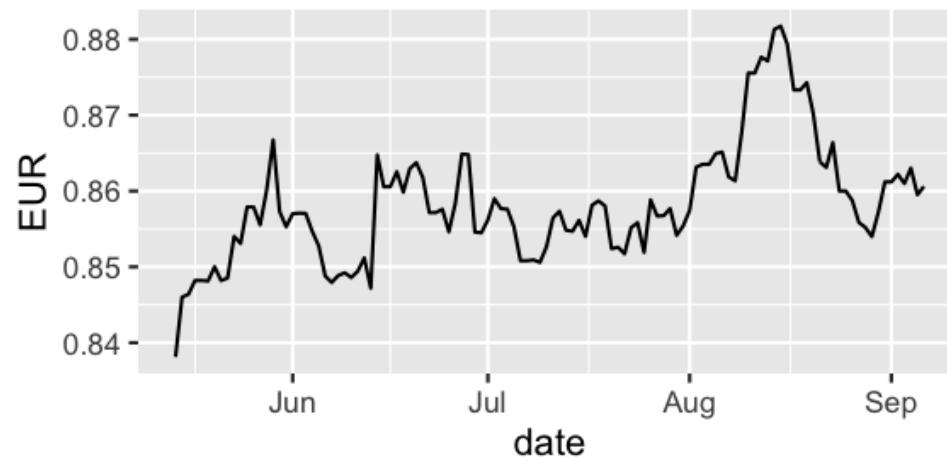
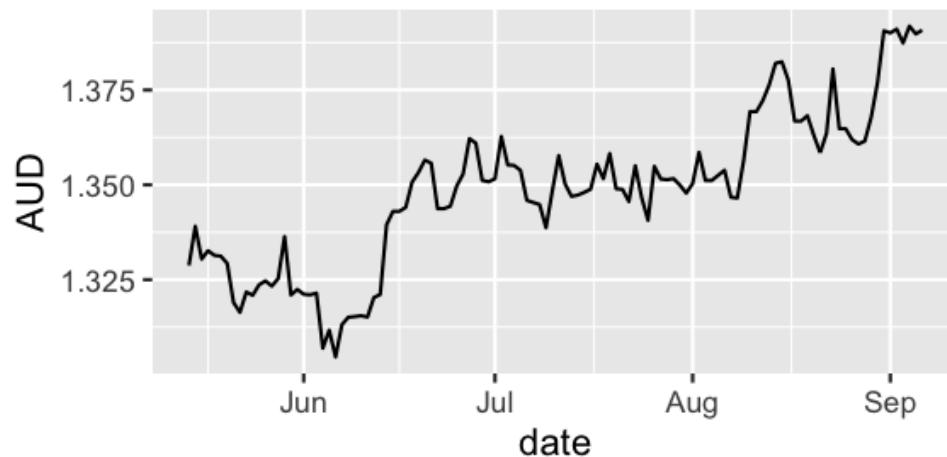
SO let's try this with cross-currency rates across the globe!

- Data extracted from <http://openexchangerates.org/api/historical>
- R packages jsonlite, processed with tidyverse, lubridate

Data: Last 4 months of currency USD cross-rates

```
## # A tibble: 6 x 171
##   date      AED    AFN    ALL    AMD    ANG    AOA    ARS    AUD    AWG    AZN    BAM
##   <date>     <dbl>  <dbl>
## 1 2018-05-14  3.67  71.2  106.  485.  1.79  230.  25.0  1.33  1.79  1.70  1.63
## 2 2018-05-15  3.67  71.2  107.  485.  1.80  230.  24.1  1.34  1.79  1.70  1.64
## 3 2018-05-16  3.67  71.0  108.  484.  1.80  232.  24.3  1.33  1.79  1.70  1.66
## 4 2018-05-17  3.67  71.0  108.  483.  1.80  233.  24.3  1.33  1.79  1.70  1.66
## 5 2018-05-18  3.67  71.0  108.  483.  1.80  233.  24.4  1.33  1.79  1.70  1.66
## 6 2018-05-19  3.67  70.9  108.  482.  1.79  233.  24.4  1.33  1.79  1.70  1.66
## # ... with 159 more variables: BBD <dbl>, BDT <dbl>, BGN <dbl>, BHD <dbl>,
## #   BIF <dbl>, BMD <dbl>, BND <dbl>, BOB <dbl>, BRL <dbl>, BSD <dbl>,
## #   BTC <dbl>, BTN <dbl>, BWP <dbl>, BYN <dbl>, BZD <dbl>, CAD <dbl>,
## #   CDF <dbl>, CHF <dbl>, CLF <dbl>, CLP <dbl>, CNH <dbl>, CNY <dbl>,
## #   COP <dbl>, CRC <dbl>, CUC <dbl>, CUP <dbl>, CVE <dbl>, CZK <dbl>,
## #   DJF <dbl>, DKK <dbl>, DOP <dbl>, DZD <dbl>, EGP <dbl>, ERN <dbl>,
## #   ETB <dbl>, EUR <dbl>, FJD <dbl>, FKP <dbl>, GBP <dbl>, GEL <dbl>,
## #   GGP <dbl>, GHS <dbl>, GIP <dbl>, GMD <dbl>, GNF <dbl>, GTQ <dbl>,
## #   GYD <dbl>, HKD <dbl>, HNL <dbl>, HRK <dbl>, HTG <dbl>, HUF <dbl>,
## #   IDR <dbl>, ILS <dbl>, IMP <dbl>, INR <dbl>, IQD <dbl>, IRR <dbl>,
## #   ISK <dbl>, JEP <dbl>, JMD <dbl>, JOD <dbl>, JPY <dbl>, KES <dbl>,
```

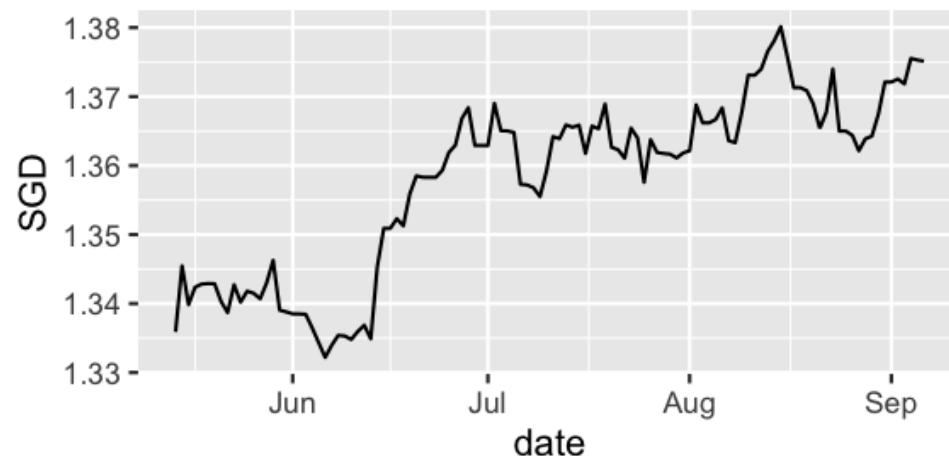
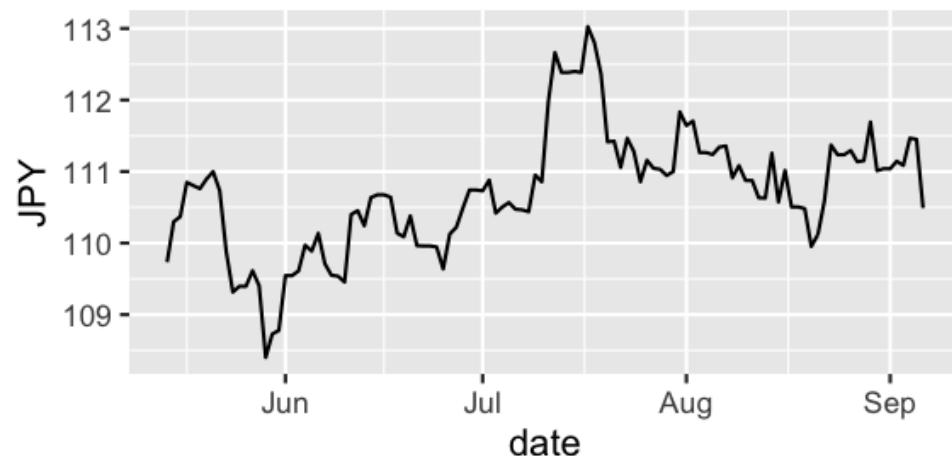
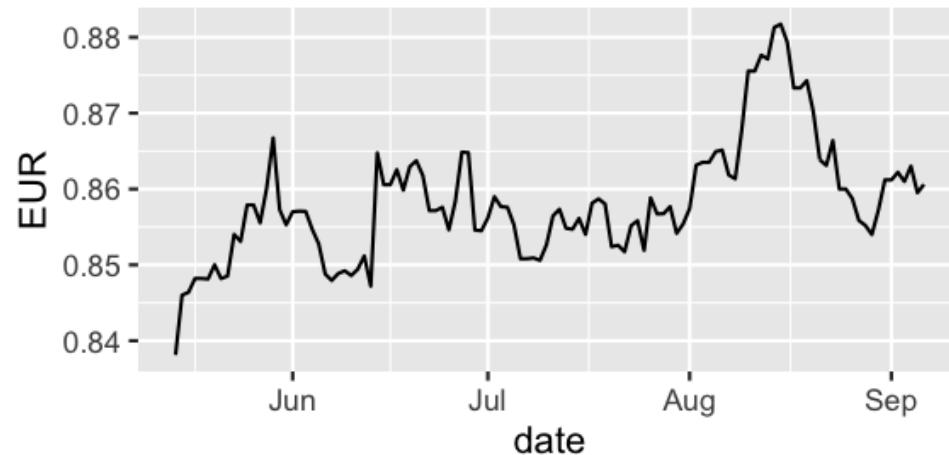
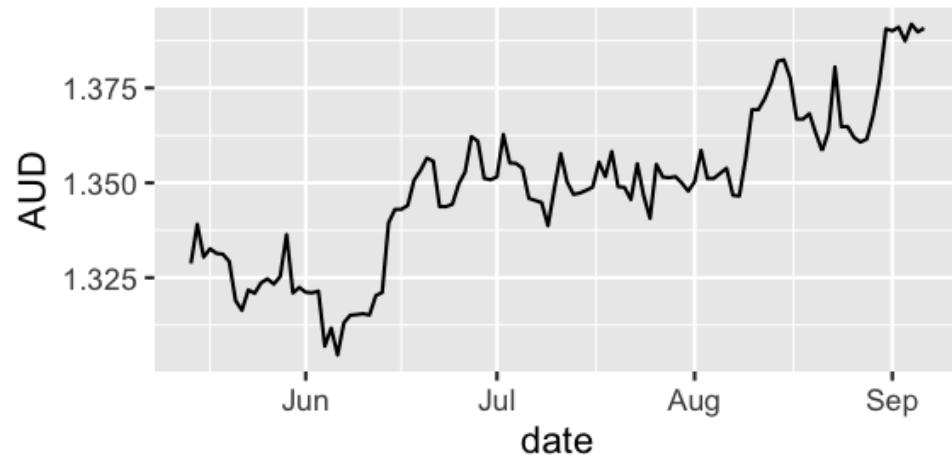
Data: Last 4 months of currency USD cross-rates



Your turn: Rstudio

Make some plots (or google) to answer these questions

- Is the NZD more similar to AUD, EUR, or JPY? (What currency is NZD?)
- Is SGD more similar to AUD, EUR, or JPY? (What currency is SGD?)
- How many currencies are there in the British Isles?



Pre-processing: Keep currencies that change

- Some currencies don't change very much.
- These should be filtered from the analysis, because in a study of currency movement, if it doesn't move then there is nothing more to be said.

Pre-processing: Keep currencies that change

- To filter out these currencies we use a statistic called coefficient of variation:

$$CoefVariation = \frac{\sigma}{\mu}$$

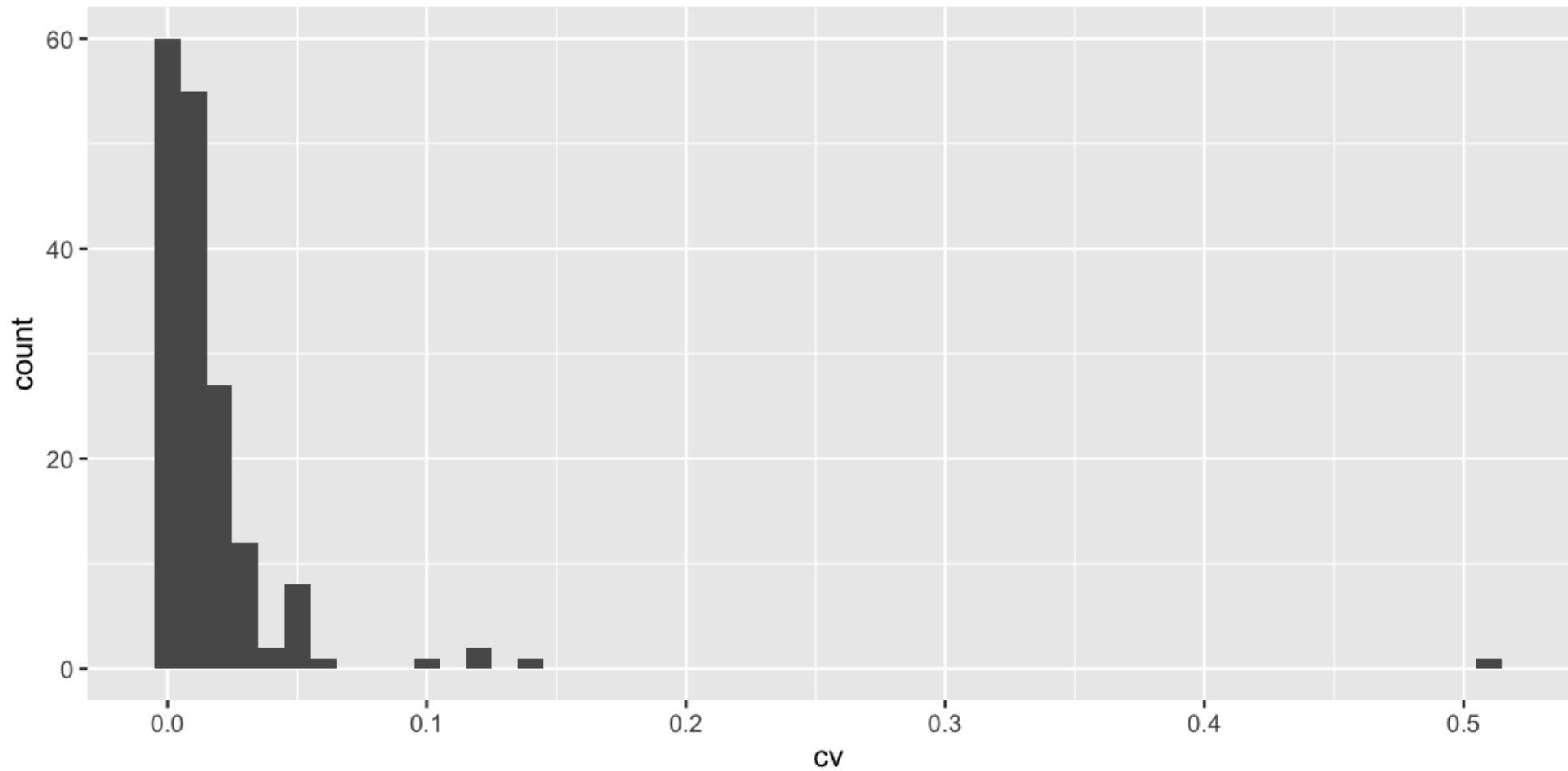
- Measures standard deviation of currency relative to the mean.
- For high means, we expect a currency to change more.
- That is, relatively the standard deviation would be larger to consider it to be changing.

Computing CV

Stratgey pivot to long form then group and summarize currency values

```
# Compute coefficient of variation. We will only analyse  
# currencies that have changes substantially over this time.  
cv <- function(x){  
  sd(x)/mean(x)  
}  
  
rates_cv <- rates %>%  
  pivot_longer(cols = -date, names_to = "currency") %>%  
  group_by(currency) %>%  
  summarise(cv = cv(value))
```

Distrubtion of CV values



Filter out currencies if they have CVs below the first quantile

```
rates_stable <- rates_cv %>%  
  filter(cv < quantile(cv, 0.25))
```

Filter out low cv currencies using pivot and an anti join

```
rates_sub <- rates %>%
  pivot_longer(cols = -date, names_to = "currency") %>%
  anti_join(rates_stable)
rates_sub
## # A tibble: 14,732 x 3
##   date      currency value
##   <date>    <chr>     <dbl>
## 1 2018-05-14 AFN       71.2
## 2 2018-05-14 ALL      106.
## 3 2018-05-14 ANG       1.79
## 4 2018-05-14 AOA      230.
## 5 2018-05-14 ARS       25.0
## 6 2018-05-14 AUD       1.33
## 7 2018-05-14 BAM       1.63
## 8 2018-05-14 BDT       84.7
## 9 2018-05-14 BGN       1.64
## 10 2018-05-14 BIF      1767.
## # ... with 14,722 more rows
```

Remove currencies that are not currencies

Some of the currencies ... aren't really currencies. Google these ones:
XAG, XDR, XPT - what are they?

Remove currencies that are not currencies

```
# Remove non-currencies  
rates_dropped <- rates_sub %>%  
  filter(!currency %in% c("ALL", "XAG", "XDR", "XPT"))
```

XAG is Gold XPT is Platinum XDR is special drawing rights

Standardize the currencies

To examine overall trend regardless of actual USD cross rate, standardise the values to have mean 0 and standard deviation 1.

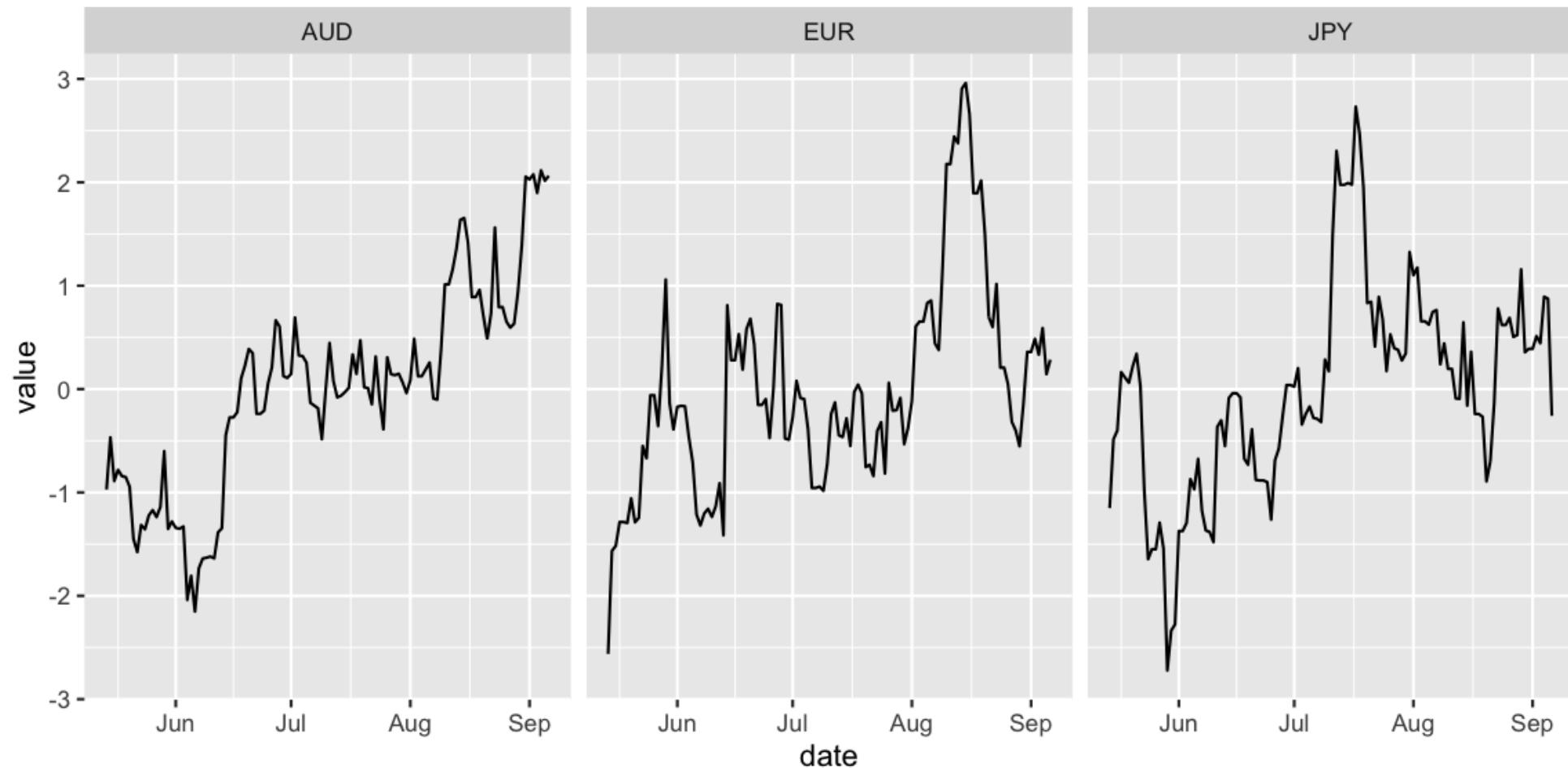
```
scale01 <- function(x) (x - mean(x)) / sd(x)
```

Rescale all values to have standardised values

Use `group_by()` plus `mutate()`!

```
rates_scaled <- rates_dropped %>%  
  group_by(currency) %>%  
  mutate(value = scale01(value))
```

Standardize the currencies



Compute distances between all pairs of currencies

Euclidean distance is used to compute similarity between all pairs of currencies.

$$d_{ij} = \sqrt{\sum_{i=1}^t (C_{1i} - C_{2i})^2}$$

Compute distances between all pairs of currencies

We need to put our data back in wide form! And then turn it into a matrix.

```
rates_wide <- rates_scaled %>%
  pivot_wider(id_cols = "date", names_from = "currency") %>%
  select(-date)

# compute distance between currencies, rows <--> columns
rates_wide_t <- t(rates_wide)
```

Use built in function to compute distance

```
currency_dist <- as.matrix(dist(rates_wide_t,
                                diag = TRUE,
                                upper = TRUE))

currency_dist[1:5, 1:5]
##          AFN       ANG       AOA       ARS       AUD
## AFN 0.000000 8.044527 7.315939 8.014165 7.970993
## ANG 8.044527 0.000000 5.628321 9.601101 7.277124
## AOA 7.315939 5.628321 0.000000 5.760894 5.299254
## ARS 8.014165 9.601101 5.760894 0.000000 5.983452
## AUD 7.970993 7.277124 5.299254 5.983452 0.000000
```

A note on distance matrices:

- A distance matrix is the inverse of an association matrix.
- A distance matrix close to 0 means the pair are most similar.
- For an association matrix far from zero means the pair are close.
- Either can be used to generate a network.

Create network: Pivot data into long form, filter based on similarity

Here only the pairs of currencies who are closer than "4" to each other are kept.

```
distance_tbl <- currency_dist %>%
  as.data.frame() %>%
  rownames_to_column(var = "from_currency") %>%
  pivot_longer(-from_currency,
               names_to = "to_currency",
               values_to = "distance") %>%
  filter(distance < 4) %>%
  filter(from_currency != to_currency)
```

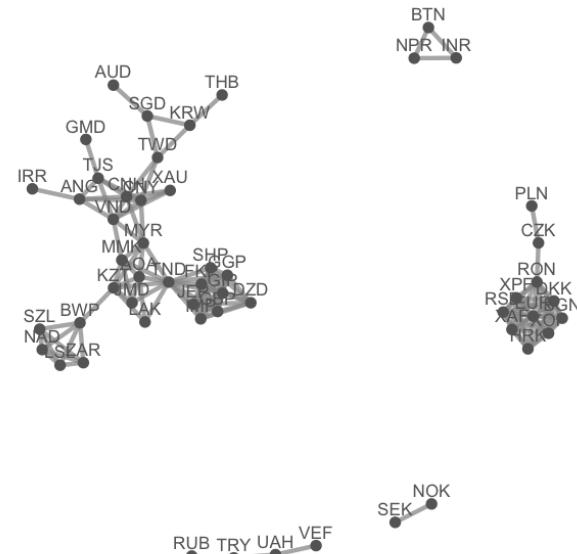
Create network: Gather data into long form, filter based on similarity

Here only the pairs of currencies who are closer than "4" to each other are kept.

```
distance_tbl  
## # A tibble: 266 x 3  
##   from_currency to_currency distance  
##   <chr>        <chr>        <dbl>  
## 1 ANG          CNH          2.98  
## 2 ANG          CNY          3.24  
## 3 ANG          IRR          3.73  
## 4 ANG          TJS          3.60  
## 5 ANG          VND          3.42  
## 6 AOA          JMD          3.66  
## 7 AOA          KZT          2.11  
## 8 AOA          LAK          3.55  
## 9 AOA          MMK          2.19  
## 10 AOA         MYR          2.17  
## # ... with 256 more rows
```

Network laid out

```
# Make network
library(geomnet)
set.seed(10052016)
ggplot(data = distance_tbl,
       aes(
         from_id = from_currency
         to_id = to_currency
       )) +
  geom_net(
    layout.alg = "kamadakawai",
    size = 2,
    labelon = TRUE,
    vjust = -0.6,
    ecolour = "grey60",
    directed = FALSE,
    fontsize = 3,
    ealpha = 0.5
  ) +
  theme_net() +
  theme(
```





That's it!

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