Homework 2 (CSC 495)

Kai Chung Ying Spring 2017

Working with the Beer Advocate data

Bipartite networks and ego networks

```
library(knitr)
setwd("/Users/KevQuant/Desktop/Depaul/csc495/wk2/hwk2")
read_chunk("hwk2_2.R")
knitr::opts_chunk$set(echo = TRUE)
```

Step 1: Load libraries

```
# Load libraries

library("ggplot2")
# Must load other packages first before SAND
library("sand")
library("intergraph")
```

Step 2: Load the data 1 pt

Load a bipartite network built from the "Beer Advocate" review data. For the purposes of this assignment, I selected only very active users (> 1000 reviews), chose the beers with the very top ratings (> 4.5 out of 5) and looked at just "American IPA" style beers. This gives a network of managable size with 278 nodes (131 beers and 147 users).

The data is in edgelist and attribute form, so we have to shape it into a network first. The two files are

- \bullet beer_edges.csv
- beer_attrs.csv

```
# Load edge and attribute data
setwd("/Users/KevQuant/Desktop/Depaul/csc495/wk2/hwk2")
beer_attrs<- read.csv("beer_attrs.csv",stringsAsFactors = TRUE)
beer_edges<-read.csv("beer_edges.csv",stringsAsFactors = FALSE)</pre>
```

Step 3: Use graph.data.frame to convert to graph 1 pt

Step 4: Looking at attributes 1 pt

List the vertex attributes and note that most attributes are for beers. We don't have attributes for users.

Use the head function to list the first 6 (head) beer names (not the numeric IDs)

```
# Examine attributes
list.vertex.attributes(beer_network)

## [1] "name" "brewery" "beer.name" "alcohol" "type"
head(V(beer_network)$beer.name,6)

## [1] "Hoppin' To Heaven IPA"

## [2] "Point Defiance IPA"

## [3] "Founders Centennial IPA"

## [4] "Big Man Ale"

## [5] "Founders Harvest Ale"

## [6] "Sierra Nevada Anniversary Ale (2007-2009)"
```

Step 5: Projection 1 pt

Compute the user-user and beer-beer projections and print the summary information for each.

```
# Projections
beer_network_b2b<-bipartite_projection(beer_network, which="TRUE")
beer_network_u2u<-bipartite_projection(beer_network, which="FALSE")
summary(beer_network_b2b)

## IGRAPH UNW- 131 2373 --
## + attr: name (v/c), brewery (v/c), beer.name (v/c), alcohol (v/n),
## | weight (e/n)

summary(beer_network_u2u)

## IGRAPH UNW- 147 5070 --
## + attr: name (v/c), brewery (v/c), beer.name (v/c), alcohol (v/n),
## | weight (e/n)</pre>
```

Step 6: Fixing the attributes 1 pt

Remove the irrelevant attributes from the user-user network. Print the summary at the end. It should show only "name" and "weight" attributes. Use delete_vertex_attr.

```
# Remove non-user attributes
beer_network_u2u<-delete_vertex_attr(beer_network_u2u, "brewery")
beer_network_u2u<-delete_vertex_attr(beer_network_u2u, "beer.name")
beer_network_u2u<-delete_vertex_attr(beer_network_u2u, "alcohol")
summary(beer_network_u2u)

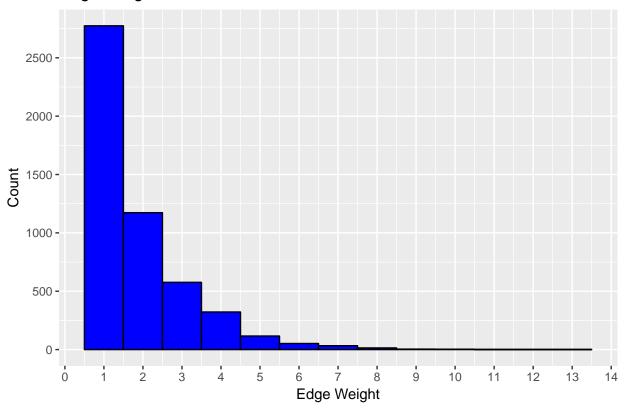
## IGRAPH UNW- 147 5070 --
## + attr: name (v/c), weight (e/n)</pre>
```

Step 7: Plot the edge weight distribution 2 pt

You can use either base plotting or ggplot. Label your plots appropriately (x and y axes, main title). Make sure that the whole distribution is included – set x axis limits correctly.

```
# Edge weight histogram
library(ggplot2)
par(mar=c(1,1,1,1))
g<-ggplot(data.frame(weight=E(beer_network_u2u)$weight),aes(x=weight))
g<-g+geom_histogram(binwidth = 1,col="black",fill="blue")
g<-g+xlab("Edge Weight")
g<-g+ylab("Count")
g<-g+ylab("Count")
g<-g+ggtitle("Edge Weight Distribution")
g<-g+scale_x_continuous(breaks=seq(0,14,1))
g<-g+scale_y_continuous(breaks=seq(0,3000,500))
print(g)</pre>
```

Edge Weight Distribution



```
#Alternative plotting option
#barplot(table(E(beer_network_u2u)$weight))
```

Step 8: Filter out edges of weight < 6 2 pts

As is typical with projections of bipartite networks, we'll filter out the low weight edges of which there are very many.

```
# Filter edges of weight less than 6
beer_network_u2u_filt<-delete.edges(beer_network_u2u,E(beer_network_u2u)[E(beer_network_u2u)$weight<6])</pre>
```

```
#Showing the weight of the edges after filtering
table(E(beer_network_u2u_filt)$weight)
```

```
## ## 6 7 8 9 10 11 12 13
## 52 33 14 4 3 1 1 1
```

Step 9: Remove singletons (nodes of degree == 0) 1 pt

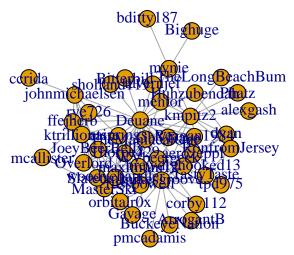
Removing edges leaves some nodes disconnected from the network so we remove them also. Another way to do this would be calculate the largest (giant) component.

Note: Use the filtered version of the network for the rest of the assignment.

Step 10: Plot the network 1 pt

The graph is now simplified enough that it can be visualized. Use the Kamada-Kawai layout: layout=layout_with_kk.

```
# Plot
plot(beer_network_u2u_filt2,layout=layout_with_kk)
```



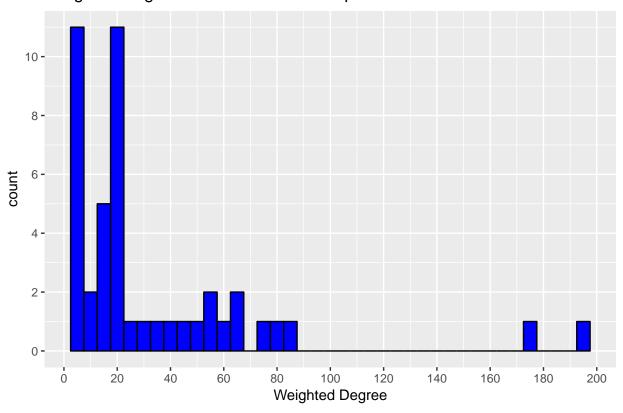
Step 11: Weighted degree 2~pt

Compute a histogram of the weighted degree. (graph.strength function)

```
# Weighted degree histogram
beer_network_u2u_filt2_weight_deg<-graph.strength(beer_network_u2u_filt2)
beer_network_u2u_filt2_weight_deg_df<-data.frame(weight_degree=beer_network_u2u_filt2_weight_deg)
g2<-ggplot(beer_network_u2u_filt2_weight_deg_df,aes(x=weight_degree))
g2<-g2+geom_histogram(binwidth = 5,col="black",fill="blue")
g2<-g2+ggtitle("Weighted Degree of User-to-User Graph")
g2<-g2+xlab("Weighted Degree")</pre>
```

```
g2<-g2+scale_x_continuous(breaks=seq(0,200,20))
g2<-g2+scale_y_continuous(breaks=seq(0,14,2))
print(g2)</pre>
```

Weighted Degree of User-to-User Graph



Step 12: Find outliers 2 pt

Find the names of the two individuals with highest weighted degree (2 outliers)

Step 13: Compute the ego networks 1 pt

Compute the ego networks for these two users.

```
# Create ego networks
which(V(beer_network_u2u_filt2)$name=="Deuane")
```

[1] 12

```
which(V(beer_network_u2u_filt2)$name=="mikesgroove")

## [1] 26

#Compute the ego network with order =1
egos<-make_ego_graph(beer_network_u2u_filt2,1,nodes = c(12,26))</pre>
```

Step 14: Plot the ego networks side by side 1 pt

Use par(mfrow=c(1,2)) to get the side-by-side layout. Remember to switch back to the normal layout c(1,1) after.

```
# Plot ego networks
#Extract the ego graph from the egos
deuane<-egos[[1]]
mikesgroove<-egos[[2]]

#Find their order on the graph
which(V(deuane)$name=="Deuane")

## [1] 5
which(V(mikesgroove)$name=="mikesgroove")

## [1] 15
#The order numbers were found 5 and 15 respectively,
#And then use these information to make the layout
deuane.lo<-layout_as_star(deuane,V(deuane)[5])
mikesgroove.lo<-layout_as_star(mikesgroove,V(mikesgroove)[15])

#plot row=1, col=2 format</pre>
```



Step 15: Question 3 pts

par(mfrow=c(1,2))

plot(deuane,layout=deuane.lo)

plot(mikesgroove,layout=mikesgroove.lo)

In reducing the size of the network through edge and vertex filtering (steps 8 and 9) so that it is easier to visualize, what information about the original user-user network has been lost? What consequences does this have for our interpretation of the filtered version of the network?

```
### From step 8, we removed the edges with weight less than 6.
### From step 9, we removed the node with degree equals 0.

### For the above action, we might skip some network connection for further analysis.
### For the removal of weight less than 6 (Step 8), we might have skip some users with
### less beer preferences, but actually they might be favorite to different kind of beer.
### In other words, this group of users might be our potential users in the future.
### And discovering this area of users might lead to other area of beer industry.

### For those node with degree equals 0 (Step 9.) which takes up 102 out of 147 vertices is
### about two-third of the overall vertices. This isolated group is actually a huge resources for us.
### We could definitly perform another analysis to discover why they don't have any edge connection
### with other vertices OR they might actually have NO any preference from the beer list. This might
### be able to help the prove the beer category/ marketing stragegy / district investment ratio.
```