Machine Learning Workshop - iGraph Example

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Contents

Overview	1
iGraph Package	1
References	16

Overview

This vignette provides a brief introduction to networks using R's igraph package which is explained here. It also shows an example of fitting a stochastic block model using the blockmodels package. For more information on stochastic block models, see this document by Emmanuel Abbe.

iGraph Package

Network analysis or graph theory involves describing a network of individuals and how they interact. Common finance applications include models to assess systemic risk (Billio et al. (2012)) who develop several measures of systemic risk based on networks. Another good example is a series of papers by Gerard Hoberg (USC) and Gordon Phillips (Dartmouth). They developed a new measure of industries based on text mining of corporate reports. Essentially, what they do is look for similarities in the product market space based on SEC reports. They then convert the text mining to similarity scores and use those scores to cluster firms into groups. The clusters represent industries. The data is freely available here. We will be using one of their annual datasets in this vignette. However, this document only very briefly describes network theory; the main intent is to provide an example of how to run some basic network models in R.

The fist step is to import the data. We will search over the current directory for a pattern matching our file. In practice, this is a useful way if yo have batched data (e.g., a series of years of data with structured names - data1997.txt, data1998.txt, etc.). The list.files function is goin gto search over a given directory for files matching a patter - in this example tnicall*txt.

```
rm(list=ls())
setwd("C:/Users/CLARKB2/Documents/Classes/ML Course")
dir()
##
    [1] "10Algorithms-08.pdf"
##
    [2] "13_datafest_cart_talk.pdf"
    [3] "2019_Shaft_MPF_Woods.xlsx"
##
##
    [4] "APM.pdf"
##
    [5] "Best paper award PPBRC 2019 Brian Clark.pdf"
    [6] "bib.bib"
##
##
       "cart.tree.png"
    [8] "Class01_001_Intro_to_R.Rmd"
```

```
[9] "Class01_001_Intro_to_R[rmd2r].R"
  [10] "Class05_001_optimization_Ndim.Rmd"
  [11] "Class05 001 optimization Ndim[rmd2r].R"
## [12] "Class06_002_portfolioOptimization_nl.Rmd"
  [13] "Class06 002 portfolioOptimization nl[rmd2r].R"
  [14] "Comparative Study Id3 Cart And C4.5 Deci.pdf"
## [15] "Compress Mort Data Annual.R"
## [16] "ESLII.pdf"
  [17] "ISLR Seventh Printing.pdf"
## [18] "kMeannsCluster.R"
## [19] "KNN.png"
## [20] "M 2017-12-05.pdf"
  [21] "Master Machine Learning Algorithms.pdf"
## [22] "ML_002_KNN.nb.html"
## [23] "ML_002_KNN.Rmd"
## [24] "ML_Course_Agenda.pdf"
  [25] "ML_Course_Agenda.Rmd"
  [26] "ML Course C45.R"
  [27] "ML_Course_Ensemble.nb.html"
## [28] "ML Course Ensemble.pdf"
## [29] "ML_Course_Ensemble.R"
## [30] "ML Course Ensemble.Rmd"
## [31] "ML_Course_Ensemble[rmd2r].R"
## [32] "ML Course iGraph.pdf"
## [33] "ML Course iGraph.Rmd"
## [34] "ML Course Ind.Rda"
## [35] "ML_Course_Keras.R"
  [36] "ML_Course_Kmeans.nb.html"
## [37] "ML_Course_Kmeans.pdf"
## [38] "ML Course Kmeans.Rmd"
## [39] "ML_Course_Kmeans[rmd2r].R"
  [40] "ML_Course_KNN_Function.nb.html"
  [41] "ML_Course_KNN_Function.pdf"
## [42] "ML_Course_KNN_Function.Rmd"
  [43] "ML Course KNN Function[rmd2r].R"
## [44] "ML Course KNN Script.R"
## [45] "ML Course Mortgage Data.pdf"
## [46] "ML_Course_Mortgage_Data.R"
## [47] "ML_Course_Mortgage_Data.Rmd"
## [48] "ML_Course_Mortgage_Data[rmd2r].R"
## [49] "ML Course OCC 001.nb.html"
## [50] "ML Course OCC 001.pdf"
  [51] "ML Course OCC 001.Rmd"
##
## [52] "ML_Course_Optimization_Basics.pdf"
## [53] "ML_Course_Optimization_constrOptim_nl.pdf"
## [54] "ML_Course_Ridge_LASSO.nb.html"
  [55] "ML_Course_Ridge_LASSO.pdf"
##
  [56] "ML_Course_Ridge_LASSO.Rmd"
  [57] "ML_Course_Ridge_LASSO[rmd2r].R"
  [58] "ML_Course_SVM.nb.html"
## [59] "ML_Course_SVM.pdf"
## [60] "ML_Course_SVM.Rmd"
## [61] "ML_Course_SVM[rmd2r].R"
## [62] "ML Course Trees.nb.html"
```

```
## [63] "ML_Course_Trees.pdf"
##
  [64] "ML_Course_Trees.R"
  [65] "ML Course Trees.Rmd"
  [66] "ML_Course_Trees[rmd2r].R"
##
##
  [67] "ML_Seminar_NonQuant_OCC_001.nb.html"
       "ML Seminar NonQuant OCC 001.pdf"
  [68]
##
       "ML Seminar NonQuant OCC 001.Rmd"
  Г691
  [70] "Mortgage_Annual.Rda"
##
  [71] "Motrgage_Annual.Rda"
##
  [72]
       "New folder"
  [73] "Non-Quant-OCC_FINAL_FINAL-forECON.pptx"
   [74] "Regression_Trees_and_Rule_Based_Models_Install.R"
  [75]
       "rmarkdown-reference.pdf"
## [76] "RMD2R.R"
## [77]
       "Rpart_details.pdf"
  [78]
        "SBM_ML_Course"
##
  [79]
       "The caret Package.pdf"
  [80] "tnicall1997.txt"
  [81] "trees.txt"
  [82] "Value_of_Location_in_Derivative_Markets.pdf"
files <- list.files(pattern = glob2rx("tnicall*txt"),</pre>
                    full.names=TRUE)
print(files)
```

[1] "./tnicall1997.txt"

Once we have the list of files, we can read it in. Note that this step takes a bit of time since the raw file is large. We will reduce the file size before running any network analyses.

```
# Read the first file in (for testing use 1997):
df <- read.table(files[1],sep="\t",header=T)

# Take a look at the first 10 observations
head(df,10)</pre>
```

```
##
       score gvkey1 gvkey2 ball year
## 1
          NA
                1004
                       1004
                               NA 1997
## 2
      0.0000
                1004
                       1013
                                1 1997
## 3
      0.0363
                1004
                       1021
                                1 1997
## 4
      0.0000
                1004
                       1034
                                1 1997
## 5
     0.0000
                1004
                       1038
                                1 1997
     0.0000
## 6
                1004
                       1043
                                1 1997
## 7
      0.0567
                1004
                       1045
                                1 1997
## 8 0.0000
                1004
                       1050
                                1 1997
## 9 0.0699
                1004
                       1056
                                1 1997
## 10 0.0000
                1004
                       1072
                                1 1997
# drop ball and year vars:
```

 $df \leftarrow df[,c(-4,-5)]$

The next thing that we want to know is how many unique firms we have. Our graph (network) is going to be defined by what's called an adjacency matrix which is a square matrix of dimension n, where n is the number of nodes (in our case firms). The links between the nodes are called edges and those are given by the pairwise connections between our nodes. In the Gordon and Phillips data, the edges are the pairwise similarity scores.

The next step is then to get the number of unique firms so we know how large our graph is going to be.

```
# Get the unique firms:
unq.1 <- unique(df$gvkey1)
unq.1.n <- length(unq.1)
print(unq.1.n)

## [1] 7518
unq.1.n*unq.1.n

## [1] 56520324
unq.2 <- unique(df$gvkey2)
unq.2.n <- length(unq.2)
print(unq.2.n)

## [1] 7518
unq.2.n*unq.2.n</pre>
## [1] 56520324

## [1] 56520324

## [1] 56520324

## [1] 56520324

## [1] 56520324

## [1] 56520324

## [1] 56520324

## [1] 56520324
```

Note that we have 7,518 firms s our adjacency matrix wil have over 56 million edges! If we don't reduce the

size of the problem, the rest of the code will not run (on my machine anyway).

But first, let's get some basic information on the scores. We want to count the number of scores that are approximately equal to zero. This gives us an idea of how sparse our graph is going to be.

```
# Count no. of scores = 0:
score.0 <- (df$score[df$score <= 0.00001])
# print(score.0)
length(score.0)

## [1] 29601126
score.0plus <- (df$score[df$score > 0.00001])
length(score.0plus)

## [1] 26517790
cutoff <- 0.2132
n.score.below <- length((df$score[df$score <= cutoff]))
n.score.above <- length((df$score[df$score > cutoff]))
n.score.above/m
```

[1] 0.004549307

[1] 56111398

The next step is to sample the data by selecting 50 random firms. We will have an adjacency matrix of dimesion 50×50 .

```
# Make the adjency matrix:
library("igraph")

##
## Attaching package: 'igraph'
## The following objects are masked from 'package:stats':
```

```
##
##
       decompose, spectrum
## The following object is masked from 'package:base':
##
##
       union
library("blockmodels")
## Loading required package: Rcpp
## Loading required package: parallel
## Loading required package: digest
# make the network:
# remove missing scores (when gukey1=gukey2):
df <- df[-which(is.na(df$score)),]</pre>
m <- dim(df)[1]
# Reorder the columns:
df \leftarrow df[,c(2,3,1)]
# make a sample network by retaining a x% of the firms:
keep <- sample(unq.1,50)
df.50 <- df[df$gvkey1 %in% keep,]</pre>
df.50 <- df.50[df.50$gvkey2 %in% keep,]
```

At this point, it makes sense to save the data so next time we run this we don't need to wait so long.

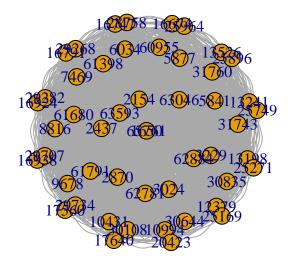
```
save(df.50,file="ML_Course_Ind.Rda")
```

Finally, we can make the adjacency matrix and we have our graph. Since the similarity scores are symmetric, we set directed=FALSE. A directed graph is one in which the edges are not symmetric in the adjacency matrix.

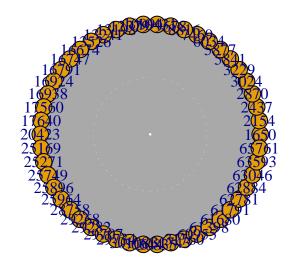
```
# make the network and adjency matrix:
net <- graph_from_data_frame(d=df.50,directed=FALSE)
adj <- as.matrix(as_adjacency_matrix(net,attr="score"))</pre>
```

Next, let's plot the graph. Network plots can be represented in many ways. Here are a few examples (check the help file - component_wise, layout_as_bipartite, layout_as_star, layout_as_tree, layout_in_circle, layout_nicely, layout_on_grid, layout_randomly, layout_with_dh, layout_with_fr, layout_with_gem, layout_with_graphopt, layout_with_kk, layout_with_lgl, layout_with_mds, layout_with_sugiyama, layout_, merge_coords, norm_coords, normalize).

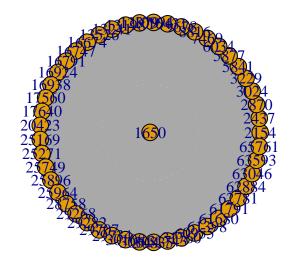
```
1 <- layout_on_sphere(net)
plot(net, layout=1)</pre>
```



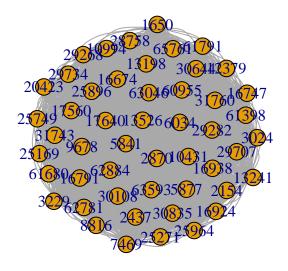
12 <- layout_in_circle(net)
plot(net, layout=12)</pre>



13 <- layout_as_star(net)
plot(net, layout=13)</pre>



14 <- layout_nicely(net)
plot(net,layout=14)</pre>



Finally, we can fit a stochastic block modeling using the **blockmodels** package. The main inputs are the type of block model and the distribution. Since our edges are close to normal, we will use a Gaussion block model. Many times you have a network where the connections are 0/1 - in that case you would use a Bernoulli block model. Note that the block models in this packages can also handle covariates.

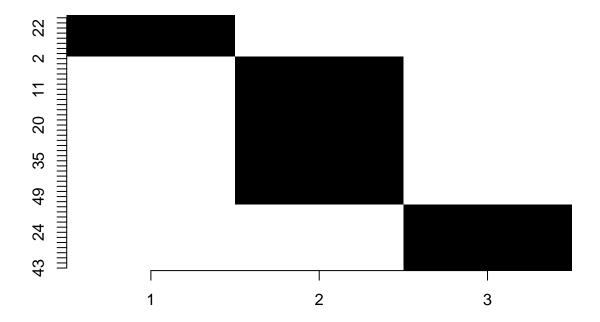
-> Pass 2

```
-> Pass 3 ##
```

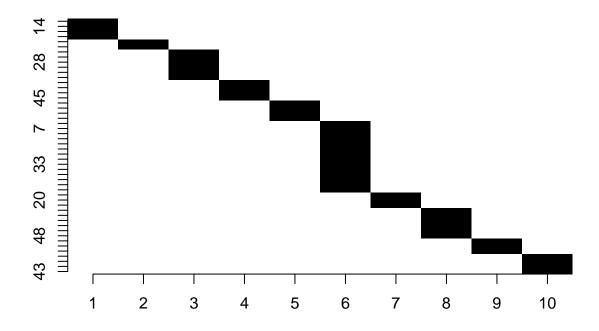
```
-> Pass 4
Z <- model.gaussian$memberships[[10]]$Z
Z.table <- apply(Z,2,table)
for (i in 1:length(Z.table)){
   message(sprintf('-----%d------',i))
   print(Z.table[[i]])
}</pre>
```

- ## [1] 46
- ## -----2-----
- ## [1] 4
- ## -----3-----
- ## [1] 48
- ## -----4
- ## [1] 2
- ## -----5----
- ## [1] 44
- ## -----6-----
- ## [1] 6
- ## -----7
- ## [1] 46
- ## -----8-----
- ## [1] 4
- ## -----9-----
- ## [1] 46
- ## -----10-----
- ## [1] 4
- ## -----11-----
- ## [1] 36
- ## -----12-----
- ## [1] 14
- ## -----13-----
- ## [1] 47
- ## -----14-----
- ## [1] 3
- ## -----15-----
- ## [1] 44
- ## -----16-----
- ## [1] 6
- ## -----17-----
- ## [1] 47
- ## -----18-----
- ## [1] 3
- ## -----19-----

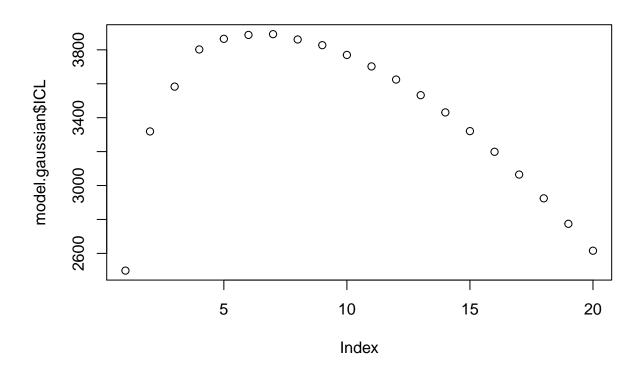
```
## [1] 46
## ----20-----
## [1] 4
model.gaussian$memberships[[3]]$plot()
```



model.gaussian\$memberships[[10]]\$plot()



How to decide the number of communities? The SBM models output a statistic, ICL which is essentially the tuning parameter.



```
print(which.max(model.gaussian$ICL))
```

We can plot the ICL values or just find the maximum which in this case was 7 meaning seven communities gives the best fit.

References

Billio, Monica, Mila Getmansky, Andrew W Lo, and Loriana Pelizzon. 2012. "Econometric Measures of Connectedness and Systemic Risk in the Finance and Insurance Sectors." *Journal of Financial Economics* 104 (3). Elsevier: 535–59.