

# Network Analysis: Homework

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## 1 Nigeria Data Processing

- a) Process the data: turn this event dataset into a matrix.
- b) Specifically, summarize the interactions across all time periods into an adjacency matrix where:
  1. "1" indicates that  $i$  and  $j$  had a conflictual interaction sometime during the temporal span of the original dataset and zero otherwise.
  2. Make sure all actors that existed at any point during the temporal span are included in the adjacency matrix.

```
1 rm(list=ls())
2 # set working directory to Git location
3 setwd('/Users/jeffziegler/Documents/Git/')
4 # load data
5 load("network2018_hw1/nigeria.rda")
6
7 #####
8 # (1) Nigeria Data Processing
9 #####
10
11 # create variable for row and column length of adjacency matrix to fill
12 rowLength <- length(unique(nigeria$sender))
13 colLength <- length(unique(nigeria$receiver))
14 # create adjacency matrix of sender and receiver, filled w/ zeroes
15 nigeriaAdjMat <- matrix(0, nrow=rowLength, ncol=colLength)
16 # adjust row and column names for sender and receiver
17 rownames(nigeriaAdjMat) <- unique(nigeria$sender)
18 colnames(nigeriaAdjMat) <- unique(nigeria$receiver)
19 # fill in adjacency matrix per year (i)
20 # start by sorting all unique years to iterate over
21 nigeriaAdjMatYearlyList <- lapply(sort(unique(nigeria$year)), function(i){
22   # find just those pairings for (1) a given year
23   currentYear <- nigeria[nigeria$year == i,]
24   # and (2) had a conflict (conflict == 1)
```

```

25 yearlyConflicts <- currentYear[currentYear$conflict == 1,]
26 # now that we know which pairings had conflicts
27 # fill in a "1" based on sender and receiver
28 for(i in 1:nrow(yearlyConflicts)){
29   nigeriaAdjMat[as.character(yearlyConflicts[i,]$sender),
30                 as.character(yearlyConflicts[i,]$receiver)] <- 1
31 }
32 # return the adjacency matrix, which will be placed in a list
33 return(nigeriaAdjMat)
34 })
35 # collapse all the matrices in list into one matrix
36 # since instructions are to "summarize the interactions
37 # across all time periods into a single matrix"
38 nigeriaAdjMatTotalMatrix <- Reduce('+', nigeriaAdjMatYearlyList)

```

## 2 Measurements & Community Detection

- a) Which actor is the most “influential” in the network? Justify your response and the measure you choose to estimate “influence.”
- b) Employ the blockmodel function from the sna package to explore potential group level structure in the data (see slides 61-63 from day 2 for details):
  - Run blockmodel with varying levels of k.
  - Save the node classifications from each run.
  - Now how do we choose k?
    - \* You will do so through an out-of-sample cross-validation exercise (at least 10 folds).
    - \* Report the AUC (ROC) and AUC (PR) statistics from each model.
- c) After having determined the k that gives the best out of sample performance, visualize your results as shown in slide 67 from the day 2 lecture

```

1 #####
2 # (2) Measurements and Community Detection
3 #####
4
5 # (a) We want to discern who the most "influential" actor in the network is
6 # We'll create two measures of "influence":
7 # (1) degree (number of connections)
8 # (2) eigenvector centrality (connections to high-scoring nodes
9 # contribute more to the score of the node)
10
11 # create graph object from igraph package
12 library(igraph)
13 igeriaGraph <- graph_from_adjacency_matrix(nigeriaAdjMatTotalMatrix,
14                                             mode='directed', weighted=TRUE, diag=FALSE)

```

```

15
16 # (1) degree
17 head(sort(degree(nigeriaGraph), decreasing=T))
18
19 # interestingly, the Police (Nigeria) and the Military (Nigeria)
20 # are two of the top 3 most engaged actors (Fulani Militia is #2)
21
22 # (2) eigenvector centrality
23 head(sort(eigen_centrality(nigeriaGraph, directed = TRUE)$vector,
24           decreasing=T))
25
26 # again, the police and military are not only more involved in conflicts
27 # but they engage w/ other highly conflicted actors
28
29 # (b) Instruction: Run blockmodel with varying levels of k
30 # Tasks/traits for blockmodel function (each run needs to):
31 # [1] Save the node classifications
32 # [2] Conduct out-of-sample CV (10 folds)
33 # [3] Report the AUC (ROC) and AUC (PR) statistics
34
35 # first, recreate matrices so that they are network objects
36 library(network)
37 nigeriaAdjMatNetworkList <- lapply(sort(unique(nigeria$year)), function(i){
38   # find just those pairings for (1) a given year
39   currentYear <- nigeria[nigeria$year == i,]
40   # and (2) had a conflict (conflict == 1)
41   yearlyConflicts <- currentYear[currentYear$conflict == 1,]
42   # now that we know which pairings had conflicts
43   # fill in a "1" based on sender and receiver
44   for(i in 1:nrow(yearlyConflicts)){
45     nigeriaAdjMat[as.character(yearlyConflicts[i,]$sender),
46                  as.character(yearlyConflicts[i,]$receiver)] <- 1
47   }
48   # return the adjacency matrix, which will be placed in a list
49   return(as.network.matrix(nigeriaAdjMat))
50 })
51
52 # create function that will do tasks 1-3
53 # then we can run CV function for varying levels of k
54 # Arguments:
55 # (remember function takes in igraph object)
56 # nFolds = number of folds (default = 10)
57 # nClusters = number of cluster (default = 2)
58 library(sna); library(caret); library(networkDynamic)
59 library(devtools)
60 install_github("leifeld/btergm", dependencies=TRUE)
61 library(btergm)
62 crossValidateFunc <- function(networkData, nFolds=10, nClusters=2) {
63   # set seed for reproducibility
64   set.seed(5)
65   # createFolds function from caret package

```

```

66 # argument gives a list of the indices in each fold
67 # from the groups that comprise all possible conflicts
68 # return training data
69 cvFolds <- createFolds(y = unique(nigeria$sender),
70                       k=nFolds, returnTrain = T)
71 # create empty vectors to fill w/ goodness-of-fit stats
72 # from TERGMs (AUC (ROC) and AUC (PR))
73 # ROC and PR curves can be used to compare different model specifications,
74 # also for within-sample goodness-of-fit
75 AUC_ROC <- NULL; AUC_PR <- NULL
76 # iterate over folds
77 for (i in 1:nFolds) {
78   # transform input list into network list
79   networkList <- networkDynamic(network.list=networkData)
80   # remove the necessary observations that are exempt from each fold
81   delete.vertices(networkList, (1:dim(nigeriaAdjMat)[1])[-cvFolds[[i]]])
82   # create clusters from structural equivalence
83   equivNetClusters <- equiv.clust(networkList)
84   # perform blockmodel
85   blockModel <- blockmodel(networkList, equivNetClusters, k=nClusters)
86   # take info that pertains to which block actors are placed in
87   groupMembership <- blockModel$block.membership[blockModel$order.vec]
88   # assign the block group values from the model back in the networkList
89   networkList%v%"member" <- groupMembership
90   # now run the out-of-sample prediction with TERGMs
91   outSampleTERGM <- btergm(as.network.networkDynamic(networkList) ~ edges +
92                          gwesp(.5, fixed = TRUE) + nodecov("member"))
93   # now, simulate 100 networks from the model w/ rocpr
94   # to condense the performance into a single measure, the area under
95   # the curve (AUC) can be reported for both curves.
96   goodFitStats <- gof(outSampleTERGM, statistics = rocpr, nsim = 100)
97   # for each iteration/fold, remove and store statistics to existing list
98   AUC_ROC <- c(AUC_ROC, goodFitStats$`Tie prediction`$auc.roc)
99   AUC_PR <- c(AUC_PR, goodFitStats$`Tie prediction`$auc.pr)
100 }
101 # return the mean of each statistic pooled over the folds
102 return(list(avgAUC_ROC=mean(AUC_ROC), avgAUC_PR=mean(AUC_PR)))
103 }

```

### 3 ERGMs

- a) Run a cross-sectional ERGM on the Nigerian conflict network, develop at least one or two network level hypotheses.
- b) Briefly discuss the results.
- c) Make sure to show that you checked for convergence.

### 4 Find your own data

- a) Locate data that relates to your field of interest.

- b) Transform the data, or a subset of it into a matrix, and plot (similar to step 1 in Section 1).
- c) Include descriptive features in your network graph (similar to step 2, but choose your own measurements).
- d) Run a model, it can be any network model from the course but justify your choices!
- e) Discuss the results in a brief write up. Present for 3-5 minutes to the class.