NB Coauthor

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In this file, we implement methods for Author Disambiguation problem in paper (Naive Bayes Model). We will first give an overall instruction about our process of Naive Bayes Method based on Nameset "AGupta", and show the results of all the Namesets at the end.

Step 0: Load the packages, specify directories

```
setwd("C:/Users/yftang/Documents/GitHub/Spr2017-proj4-team-5/lib/")
library(stringr)
```

Step 1: Load and process the data

```
source("DataCleaningTZ.R")
n=1 # Nameset "AGupta"
num.paper<-length(data list[[n]])</pre>
 auther.id<-NULL
 coauther<-NULL
 for(i in 1:num.paper){
    auther.id[i]<-data list[[n]][[i]][[1]]</pre>
   coauther<-unique(c(coauther,data list[[n]][[i]][[3]]))</pre>
 }
 a.i<-factor(auther.id)</pre>
 uni.auther<-length(levels(a.i))</pre>
 levels(a.i)<-seq(uni.auther)</pre>
 auther.id<-a.i
 df<-data.frame(auther.id)</pre>
 num.coauther<-length(coauther)</pre>
 get.coauther<-function(i){</pre>
   coauthers<-rep(0, num.coauther)</pre>
   for(j in 1:length(data_list[[n]][[i]][[3]])){
      coauthers[which(data_list[[n]][[i]][[3]][j]==coauther)]<- 1</pre>
   }
   return(coauthers)
 #### build dataframe for auther coauther information######
 id co<-NULL
 for(i in 1:num.paper){
    id_co<-rbind(id_co,get.coauther(i))</pre>
 colnames(id co)<-coauther</pre>
 df<-cbind(auther.id,id_co)</pre>
 #df contains the auther coauther information in nameset 1: AGupta
 id.num.co<-apply(df[,-1],1,sum)
 df<-as.data.frame(cbind(id.num.co,df))</pre>
 head.matrix(df)
```

##		id.num.co	auther.id E	3 Kvande I	Levinstein	K Maly M	1 Olson R	Mukkamala
##	1	8	10	1	1	1	1	1
##	2	7	10	0	0	1	0	1
##	3	8	10	1	1	1	1	1
##	4	6	10	0	0	0	0	0
##	5	6	10	0	0	1	0	1
##	6	8	10	0	0	1	0	0
## ## ## ## ##	1 2 3 4 5	R Chambers 1 0 1 0 0	1 0 1 0 0 0		d C Vemuru 1 0 0 1 1 0 0 0	H Syed H 0 1 0 0 1	d Abdel-W	ahab M Kholiet 0 6 1 1 0 6 1 6 1 6

Step 2: Naive Bayes Model

We assume that each author's citation data is generated by the naive Bayes model, and use his/her past citations as the training data to estimate the model parameters. Based on the parameter estimates, we use the Bayes rule to calculate the probability that each name entry X_i ($i \in [1, N]$, where N is the total number of candidate name entries in the citation database) would have generated the input citation.

Given an input test citation C with the omission of the query author, the target function is to find a name entry X_i in the citation database with the maximal posterior probability of producing the citation C, i.e.

$$max_i P(X_i|C) \tag{1}$$

Using the Bayes rule, the problem becomes finding:

$$max_i P(C|X_i) P(X_i) / P(C)$$
 (2)

Since P(C) does not depend on X_i Then Function (2) becomes:

$$\max_{i} P(C|X_i) P(X_i) \tag{3}$$

Function (3) is the target of our method.

First, we use coauthor as the attribute to do the Naive Bayes Model.

2.1 Attribute - Coauthor

```
freq.id<-as.data.frame(table(df$auther.id)/length(df$auther.id))
# numbers of paper for each id
colnames(freq.id)<-c("auther.id", "Freq")
num.id<-dim(freq.id)[1]</pre>
```

Split the Nameset into Train set and Test set

```
set.seed(29)
sample_split<-function(i){
    Data<- df[df$auther.id==i,]
    bound <- floor(nrow(Data)/2) #define 50 % of training and test set
    Data <- Data[sample(nrow(Data)),] #sample rows</pre>
```

```
df.train <- Data[1:bound, ]</pre>
                                                  #get training set
      df.test <- Data[(bound+1):nrow(Data), ] #get test set</pre>
      return(list(df.train=df.train,df.test=df.test))
}
train<-NULL
test<-NULL
for(i in seq(num.id)){
   train[i]<-list(sample_split(i)[[1]])</pre>
  test[i]<-list(sample_split(i)[[2]])</pre>
}
TT<-NULL
for(i in 1:num.id){
TT <- rbind(TT,test[[i]])</pre>
}
test_x<-TT[,-c(1,2)] # test set as data.frame
test_y<-TT[,2] # test lable
```

Then, we use the train set to train the required probabilities

```
NB_para<-function(data){</pre>
   # Input: df.train
   # Output: 6 probability used for NB model
      paper.num<-dim(data)[1]</pre>
      p_0<-sum(data$id.num.co==0)/paper.num</pre>
      #### P(N|X)
      p_1<-1-p_0
      #### P(Co|x)
      colsum<-apply(data,2,sum)</pre>
      data<-rbind(data,colsum)</pre>
      p_s_cx<-sum(colsum>=2)/sum(colsum>=1)
      #### P(Seen|Co,X)
      p_u cx<-1-p_s cx
      #### P(Unseen Co,X)
      total.num.co<-colsum[1]
      p a scx<-colsum/total.num.co</pre>
      p_a_scx<-p_a_scx[-c(1,2)]
      #### P(A|Seen,Co,X)
      p_a_ucx<-1/(num.coauther -sum(colsum>=1))
      #### P(A|Unseen,Co,X)
      outcome<-list(P.N.X = p_0,
                     P.Co.X = p_1
                     P.S.Co.X = p_s_cx
                     P.U.Co.X = p_u_cx
                     P.Ak.S.Co.X = p_a_scx
                     P.Ak.U.Co.X = p_a_ucx
      return(outcome)
}
    P.N.X<-NULL
    P.Co.X <-NULL
    P.S.Co.X<-NULL
    P.U.Co.X<-NULL
    P.Ak.S.Co.X<-NULL
    P.Ak.U.Co.X<-NULL
```

```
for(i in seq(num.id)){
    P.N.X[i]<-NB_para(train[[i]])$P.N.X
    P.Co.X[i]<-NB_para(train[[i]])$P.Co.X
    P.S.Co.X[i]<-NB_para(train[[i]])$P.S.Co.X
    P.U.Co.X[i]<-NB_para(train[[i]])$P.U.Co.X
    P.Ak.U.Co.X[i]<-NB_para(train[[i]])$P.Ak.U.Co.X
    P.Ak.S.Co.X[i]<-list(NB_para(train[[i]])$P.Ak.S.Co.X)
}
freq.x<-NULL
for(i in seq(num.id)){
    freq.x[i]<-dim(train[[i]])[1]
}
px<-freq.x/sum(freq.x) # px is the prior for the nameset</pre>
```

Build the Naive Bayes Model and use test set to do the prediction and calculate the accuracy rate.

```
NB model<-function(data){</pre>
      # input: Test datafram
      # data.df=test x
      # out<-NULL
      # for(m in seq(dim(data.df)[1])){
      # data = data.df[m,]
      if(sum(data)==0){
        out<-which.max(P.N.X)</pre>
      }
      else{
        condition<-function(id){</pre>
           i=id
           \# i=id, k = k-th coauther
           P_Ak_X<-NULL
           significant<-length(P.Ak.S.Co.X[[i]][data>0])
           for(k in seq(significant)){
             P_Ak_X[k]<-P.Co.X[i] * ( P.S.Co.X[i] *
                                          P.Ak.S.Co.X[[i]][data>0][k] +
                                          P.Ak.U.Co.X[i] * P.U.Co.X[i])
           }
           return(prod(P_Ak_X))
        }
        target<-NULL
        for(i in seq(num.id)){
           target[i]<-condition(i)*px[i]</pre>
        }
        out <-which.max(target)</pre>
      return(out)
    }
    est y<-apply(test x,1,NB model) ## test estimate
    accuracy<-sum(est y==test y)/length(test y)</pre>
    accuracy
## [1] 0.902027
```

Step 3 Outcomes for all nameset

```
source("NB_Co.R")
```

```
summary.all
```

```
##
         Nameset mean.accuracy StdDev.accuracy
## 1
         A Gupta
                     0.9077703
                                    0.01788024
         A Kumar
## 2
                     0.7919355
                                    0.02242651
## 3
         C Chen
                     0.8081928
                                    0.02125709
       D Johnson
## 4
                     0.8053476
                                    0.04941182
## 5
           J Lee
                     0.7986486
                                    0.01655061
        J Martin
## 6
                     0.7844828
                                    0.06514818
## 7 J Robinson
                     0.8534091
                                    0.03278212
        J Smith
## 8
                     0.7951064
                                    0.01085130
## 9
        K Tanaka
                     0.9062937
                                    0.01899445
## 10
      M Brown
                     0.8337500
                                    0.03283481
## 11
        M Jones
                     0.7526316
                                    0.02309303
## 12
        M Miller
                     0.9290476
                                    0.01317556
## 13 S Lee
## 14 Y Chen
                     0.7934840
                                    0.01377755
                     0.8664625
                                    0.01350176
```