## ProviderScoring

Rohan

```
library(plyr)
nyopth <- read.csv("nyopth.csv")</pre>
agi <- read.csv("agi.csv")</pre>
nyopth.zip = nyopth[,c(2,12)]
func <- function(x){</pre>
  substr(x,1,5)
}
nyopth.zip$nppes_provider_zip <- lapply(nyopth.zip$nppes_provider_zip,FUN = func)</pre>
nyopth.zip$nppes_provider_zip <- as.numeric(nyopth.zip$nppes_provider_zip)</pre>
nyopth.zip = nyopth.zip[!duplicated(nyopth.zip),]
agi \leftarrow agi[,c(1,5)]
nyopth <- nyopth[,c(2,8,12,16,17,18,20,21,22,23,24,25,26,27,28,29)]
tophcpcs <- names((sort(table(nyopth$hcpcs_code),decreasing = TRUE))[1:10])</pre>
spread_scores<-rep(x = 0,times = length(unique(nyopth$npi)))</pre>
names(spread_scores) <- unique(nyopth$npi)</pre>
bene_scores<-rep(x = 0,times = length(unique(nyopth$npi)))</pre>
names(bene_scores) <- unique(nyopth$npi)</pre>
scoreFunction <- function(rowdiff, q, col){</pre>
  score = 0
  rowscore = rowdiff[col]
  if (rowscore>=q[1] && rowscore<q[2]){</pre>
    score = 0
  }
  else if (rowscore>=q[2] && rowscore<q[3]){
    score = 1
  else if (rowscore>=q[3] && rowscore<q[4]){</pre>
    score = 2
  else{
    score = 3
  score
}
```

We loop over all the top hcpcs codes and add to the scores based on the spread of the submitted - payment amount and based on the beneficiaries counts.

```
for (i in tophcpcs){
    tmp = nyopth[nyopth$hcpcs_code==i,]
    tmp.avgpayment <- ddply(tmp,~npi,summarise,mean=mean(average_medicare_payment_amt))</pre>
    tmp.avgsubmitted <- ddply(tmp,~npi,summarise,mean=mean(average_submitted_chrg_amt))</pre>
    tmp.spread <- merge(tmp.avgpayment,tmp.avgsubmitted,by.x = "npi",by.y = "npi")</pre>
    #TODO:Need to reassess agi index?
    tmp.spread$diff = tmp.spread$mean.y - tmp.spread$mean.x
    t1 = join(x = tmp.spread, y = nyopth.zip, by = "npi")
    colnames(t1)[5] <- "ZIP"</pre>
    t2 = join(x = t1, y = agi)
    t2 = transform(t2, diff2 = diff/Indx)
    t2 = t2[complete.cases(t2),]
    #some kind of function that will give out points
    #depending on the quartile that the npi falls under
    q<-quantile(t2$diff2)</pre>
    t2$sc <- apply(t2, 1, FUN = scoreFunction,q = q, col = 7)
    for (j in 1:nrow(t2)){
        spread_scores[as.character(t2[j,1])] = spread_scores[as.character(t2[j,1])]+t2[j,8]
    #Dispense points based on number of beneficiaries
    tmp.linecnt <- ddply(tmp, ~npi, summarise, sum=sum(line_srvc_cnt))</pre>
    tmp.benecnt <- ddply(tmp, ~npi, summarise, sum=sum(bene_unique_cnt))</pre>
    tmp.cntdiff <- merge(tmp.linecnt,tmp.benecnt,by.x = "npi",by.y = "npi")</pre>
    tmp.cntdiff$diff = tmp.cntdiff$sum.x / tmp.cntdiff$sum.y
    q2 <- quantile(tmp.cntdiff$diff)</pre>
    tmp.cntdiff$sc <- apply(tmp.cntdiff,1,FUN = scoreFunction, q=q2, col = 4)</pre>
    for (j in 1:nrow(tmp.cntdiff)){
        bene\_scores[as.character(tmp.cntdiff[j,1])] = bene\_scores[as.character(tmp.cntdiff[j,1])] + tmp.cntdiff[j,1]) + tmp.
    }
}
## Joining by: ZIP
score_spread <- data.frame(keyName=names(spread_scores), value=spread_scores, row.names=NULL)</pre>
score_bene <- data.frame(keyName=names(bene_scores), value=bene_scores, row.names=NULL)</pre>
scores <- merge(x=score_spread,y=score_bene, by = "keyName")
colnames(scores) <- c("npi", "score_spread", "score_bene")</pre>
```

## summary(scores)

```
##
                  score_spread
                                 score_bene
          npi
## 1003018102: 1 Min. : 0.000 Min. : 0.00
## 1003058652: 1
                  1st Qu.: 2.000
                                 1st Qu.: 6.00
## 1003812454: 1
                  Median : 8.000
                                 Median :10.00
## 1003813247: 1 Mean : 9.234
                                 Mean :10.69
## 1003814682: 1
                  3rd Qu.:15.000
                                 3rd Qu.:15.00
## 1003815432: 1 Max. :30.000
                                 Max. :26.00
## (Other) :1608
```