

# Prediction of California Hospital Quality Ratings

Maria P. Frushicheva

August 15, 2016

## Introduction

**Importance:** Using hospital quality ratings, patients are able to make a better decision in what hospital they want to be treated and where the best care is available in state of California, based on overall hospital performance or based on particular medical condition or procedure.

**Question:** Can we predict hospital quality ratings based on risk adjusted mortality rates, number of deaths, number of cases, medical procedures performed and medical conditions treated for 2012-2013?

## Description of Data Set

**Dataset:** is available from [California Hospital Inpatient Mortality Rates and Quality Ratings, 2012-2013](#).

**Description of dataset:** The dataset contains risk-adjusted mortality rates, and number of deaths and cases for 6 medical conditions treated (Acute Stroke, Acute Myocardial Infarction, Heart Failure, Gastrointestinal Hemorrhage, Hip Fracture and Pneumonia) and 6 procedures performed (Abdominal Aortic Aneurysm Repair, Carotid Endarterectomy, Craniotomy, Esophageal Resection, Pancreatic Resection, Percutaneous Coronary Intervention) in California hospitals for 2012 and 2013. This dataset does not include conditions treated or procedures performed in outpatient settings.

## Description, Analysis and Cleaning of Variables in the Data Set

**Load the data from csv file.**

```
setwd("C:/Users/postdoc/Dropbox (Personal)/SpringBoard Fund/Rprojects/")
data <- read.csv("California_Hospital_Inpatient_Mortality_Rates_and_Quality_Ratings__2012-2013.csv", sep=";", header=TRUE)
df <- tbl_df(data)
```

**Dataset:** 11169 observations and 12 variables.

**Variables with missing values:**

- Risk Adjusted Mortality Rate: The Risk Adjusted Mortality Rates (RAMR) presented here adjusts the observed mortality rates. This statistical methodology takes into account pre-existing health problems that put some patients at greater risk of death to

level the playing field and allow fair comparisons across hospitals; **4754** missing values.

- Number of Deaths: Number of patients that died in this hospital; **4926** missing values.
- Number of Cases: Number of patients that had this medical procedure or condition in this hospital; **5004** missing values.

**Remove missing values, because number of missing values consists of half of dataset.**

```
df_clean <- df[which(is.na(df$X.of.Cases)==F),]
```

**Clean Dataset: 6165 observations and 12 variables.**

**Variables with no missing values:**

- Year: **3100** values for 2012 year and **3065** values for 2013 year.
- County: **55** counties.
- Hospital: **341** hospitals.
- OSHPDID: A unique number established by the Office of Statewide Health Planning and Development (OSHPD) for identifying facilities and used in the Licensed Facility Information System (LFIS). The first three numbers identify the type of facility, the next two represent the county number, and the last five are randomly assigned within each county. 570261 unique codes.
- Longitude: **Longitude** of hospital.
- Latitude: **Latitude** of hospital.
- location1: **333** levels.
- Hospital Ratings: Comparison rating based on a 95% Confidence Interval (CI). If a hospitals upper CI is less than the statewide observed rate, it is designated as performing better than the average hospital. If a hospitals lower CI is greater than the state rate, it is designated as performing worse than the average state hospital. **3 levels of Hospital Ratings:** As Expected, Better and Worse.

```
summary(df_clean$Hospital.Ratings)
```

```
## As Expected      Better      Worse
##           5797           158           210
```

- Procedure.Condition: Procedure that was performed or condition that was treated. **6** medical **procedures** performed: Abdominal Aortic Aneurysm (AAA) Repair, Carotid Endarterectomy, Craniotomy, Esophageal Resection, Pancreatic Resection, Percutaneous Coronary Intervention. **6** medical **conditions** treated: Acute Stroke, Acute Myocardial Infarction, Heart Failure, Gastrointestinal Hemorrhage, Hip Fracture and Pneumonia. Clean dataset contains **17 levels, instead of 12.**

```
summary(df_clean$Procedure.Condition)
```

```
##           AAA Repair           Acute Stroke
##           283           617
## Acute Stroke Hemorrhagic Acute Stroke Ischemic
##           466           615
## Acute Stroke Subarachnoid           AMI
```

##	241	590
##	Carotid Endarterectomy	Craniotomy
##	404	298
##	Esophageal Resection	GI Hemorrhage
##	75	622
##	Heart Failure	Hip Fracture
##	616	426
##	Pancreatic Cancer	Pancreatic Other
##	142	130
##	Pancreatic Resection	PCI
##	190	299
##	Pneumonia	
##	151	

### Decoding Procedure.Condition variable.

According to the American Stroke Association (ASA), strokes can be classified into 2 main categories: **87%** are ischemic strokes, caused by blockage of an artery; **13%** are hemorrhagic strokes, caused by bleeding. Ischemic strokes are further divided into 2 groups: thrombotic and embolic strokes. Hemorrhagic strokes are divided into 2 main categories: intracerebral and subarachnoid hemorrhages.

Our clean dataset has four categories for Acute Stroke:

- Acute Stroke: 617 observations;
- Acute Stroke Hemorrhagic: 466 observations;
- Acute Stroke Ischemic: 615 observations;
- Acute Stroke Subarachnoid: 241 observations.

Within each hospital, there are different notations for Acute Stroke variable. It suggests that different doctor uses different notations for the condition. These four categories are combined in one: Acute Stroke.

```
df_clean$Procedure.Condition <- gsub("Acute Stroke .*", "Acute Stroke", df_clean$Procedure.Condition)
df_clean$Procedure.Condition <- factor(df_clean$Procedure.Condition)
```

Two additional categories are present in Procedure.Condition variable:

- Pancreatic Cancer: 142 observations;
- Pancreatic Other: 130 observations.

These categories are separate medical conditions and are not combined in one category.

The Procedure.Condition variable contains 6 medical procedures and 8 medical conditions. To indicate what procedure was performed or what condition was treated, the Medical\_Category variable was added to the clean dataset.

```
df_clean <- df_clean %>%
  mutate(Medical_Category = ifelse(grepl("Repair", Procedure.Condition) | grep
```

```
1("Endarterectomy",Procedure.Condition) | grepl("Craniotomy",Procedure.Condition) | grepl("Resection",Procedure.Condition) | grepl("PCI",Procedure.Condition),
      "Procedure", "Condition"))
```

**Decoding Hospital.Ratings variable.**

```
df_clean <- df_clean %>% mutate(ratings =
  ifelse(grepl("As Expected",Hospital.Ratings),"0",
  ifelse(grepl("Better",Hospital.Ratings),"1",
  ifelse(grepl("Worse",Hospital.Ratings),"-1",NA))))
df_clean$ratings <- as.numeric(df_clean$ratings)
```

**Combine Acute Stroke repetitions for each hospital, so each hospital has one unique value for Procedure.Condition variable.**

```
df_clean_original <- df_clean
df_clean <- df_clean %>% group_by(Year,Hospital,Procedure.Condition,
      Longitude,Latitude,Medical_Category) %>%
  summarise(ratings = sum(ratings),
    X..of.Deaths = sum(X..of.Deaths),
    X..of.Cases = sum(X..of.Cases),
    Risk.Adjusted.Mortality.Rate = sum(Risk.Adjusted.Mortality.Rate)) %>%
  mutate(Hospital.Ratings =
    ifelse(ratings > 0,"Better",
    ifelse(ratings < 0, "Worse", "As Expected")))
df_clean$Hospital.Ratings <- as.factor(df_clean$Hospital.Ratings)
```

## Explanatory Data Analysis

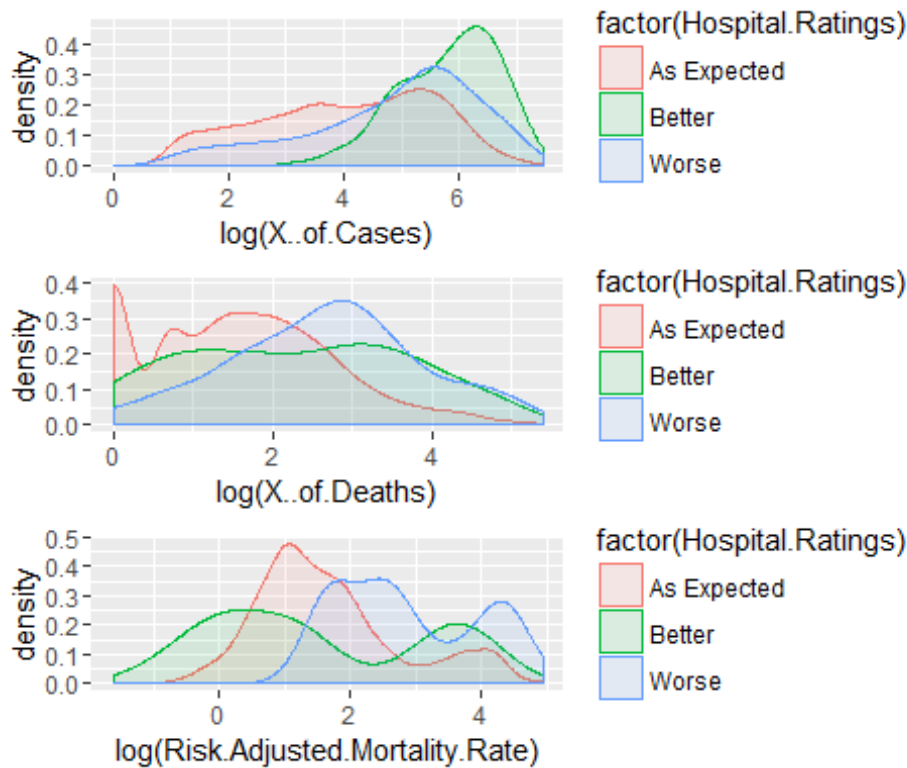
**Density Plots for # of Cases, # of Deaths and Risk Adjusted Mortality Rate by Hospital Ratings.**

```
p1 <- ggplot(df_clean,aes(log(X..of.Cases),fill=factor(Hospital.Ratings),colour=factor(Hospital.Ratings)))+
  geom_density(alpha = 0.1)

p2 <- ggplot(df_clean,aes(log(X..of.Deaths),fill=factor(Hospital.Ratings),colour=factor(Hospital.Ratings)))+
  geom_density(alpha = 0.1)

p3 <- ggplot(df_clean,aes(log(Risk.Adjusted.Mortality.Rate),fill=factor(Hospital.Ratings),colour=factor(Hospital.Ratings)))+
  geom_density(alpha = 0.1)

grid.arrange(p1, p2, p3, ncol=1)
```



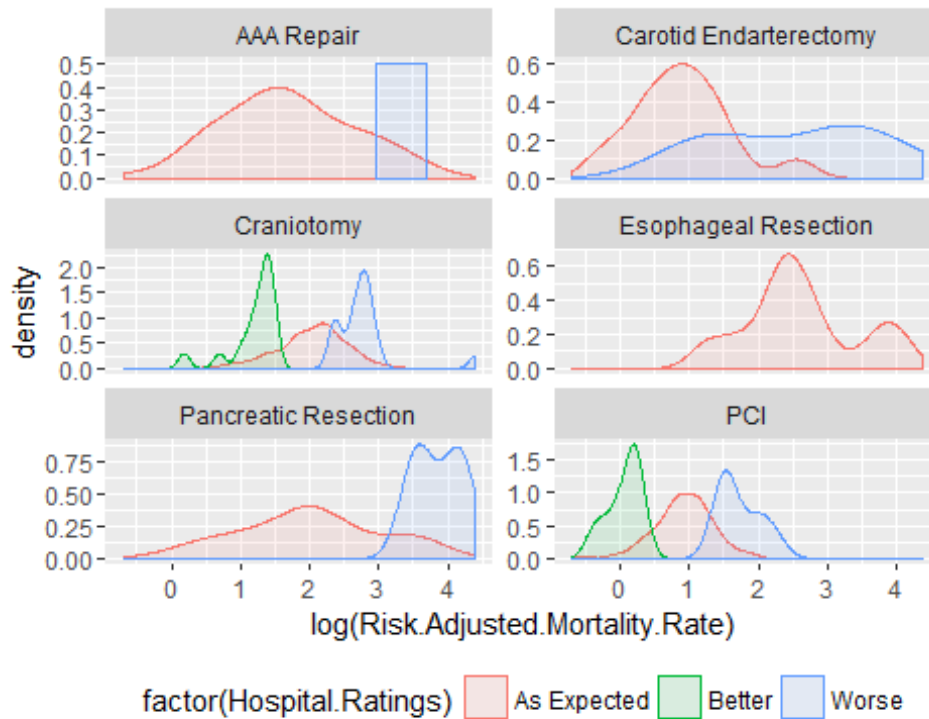
### Conclusions 1:

- Distributions between 2012 and 2013 years look similar (graphs are not shown).
- There are no associations between variables for number of deaths and number of cases.
- There is a possible **association** between the risk adjusted mortality rate and hospital ratings.
- Lower the risk adjusted mortality rate, better the hospital ratings.
- Higher the risk adjusted mortality rate, worse the hospital ratings.

### Density Plots for Risk Adjusted Mortality Rate by Procedures Performed and Hospital Ratings.

```
df_p <- df_clean[which(df_clean$Medical_Category=="Procedure"),]

p6 <- ggplot(df_p, aes(log(Risk.Adjusted.Mortality.Rate), fill=factor(Hospital.
Ratings), colour=factor(Hospital.Ratings)))+
  geom_density(alpha = 0.1)+
  theme(legend.position='bottom')+
  facet_wrap(~ Procedure.Condition, ncol=2, scales="free_y")
p6
```

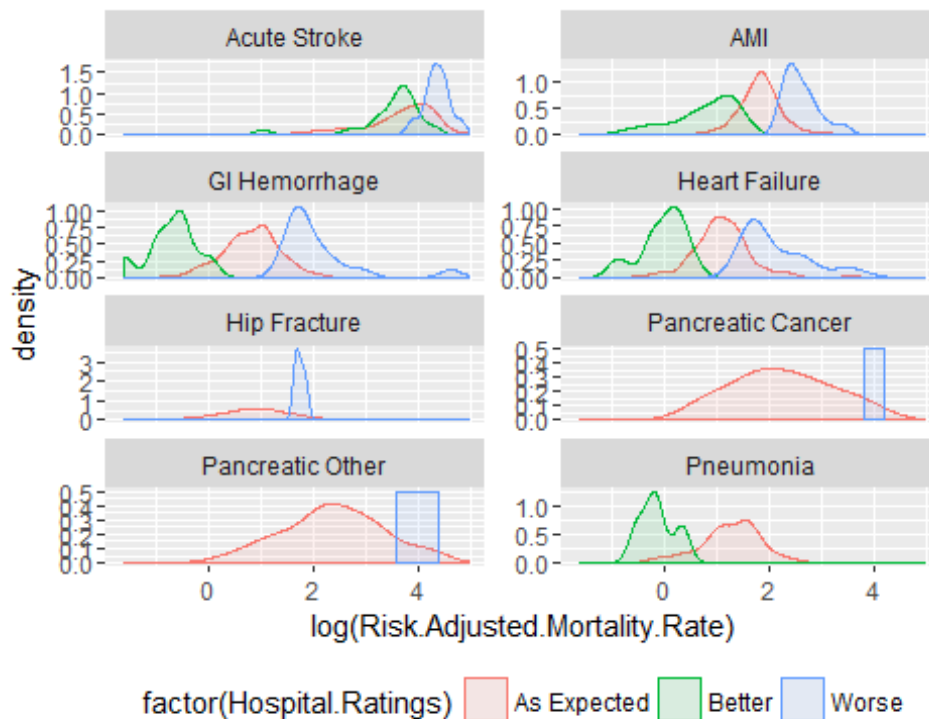


- The highest mortality rates are for Craniotomy and Pancreatic Resection procedures.
- Better and worst ratings are for Craniotomy and PCI procedures.
- There is **association** between the risk adjusted mortality rate and hospital ratings.

### Density Plots for Risk Adjusted Mortality Rate by Conditions Treated and Hospital Ratings.

```
df_c <- df_clean[which(df_clean$Medical_Category=="Condition"),]

p9 <- ggplot(df_c, aes(log(Risk.Adjusted.Mortality.Rate), fill=factor(Hospital.
Ratings), colour=factor(Hospital.Ratings)))+
  geom_density(alpha = 0.1)+
  theme(legend.position='bottom')+
  facet_wrap(~ Procedure.Condition, ncol=2, scales="free_y")
p9
```



- The highest mortality rates are for Acute Stroke, AMI and Heart Failure conditions.
- Better and worse ratings are for Acute Stroke, AMI, GI Hemorrhage and Heart Failure conditions.
- There is **association** between the risk adjusted mortality rate and hospital ratings.

**Associations between medical procedures or conditions with hospital ratings, number of cases, number of deaths and risk adjusted mortality rate.**

### Procedures.

```
df_p_all <- df_p %>%
  group_by(Procedure.Condition) %>%
  summarise(all_cases = sum(X..of.Cases),
            all_deaths = sum(X..of.Deaths),
            all_mortality_rate = sum(Risk.Adjusted.Mortality.Rate))
df_p_all
```

## # A tibble: 6 x 4

	Procedure.Condition	all_cases	all_deaths	all_mortality_rate
## 1	AAA Repair	4927	59	508.5
## 2	Carotid Endarterectomy	12478	60	290.2
## 3	Craniotomy	30164	2159	2354.6
## 4	Esophageal Resection	619	28	436.9
## 5	Pancreatic Resection	3356	93	1002.8
## 6	PCI	78660	2028	793.6

## Conditions.

```
df_c_all <- df_c %>%
  group_by(Procedure.Condition) %>%
  summarise(all_cases = sum(X..of.Cases),
            all_deaths = sum(X..of.Deaths),
            all_mortality_rate = sum(Risk.Adjusted.Mortality.Rate))
df_c_all
```

```
## # A tibble: 8 x 4
##   Procedure.Condition all_cases all_deaths all_mortality_rate
##   <fctr>             <int>      <int>          <dbl>
## 1      Acute Stroke   217956    20461      26582.9
## 2             AMI     93594     5731       3863.4
## 3      GI Hemorrhage  94804     2099       1597.8
## 4      Heart Failure 155066     4778       2200.0
## 5      Hip Fracture   32245      744        945.8
## 6 Pancreatic Cancer   1787        43        632.0
## 7 Pancreatic Other    1425        41        590.0
## 8      Pneumonia     20630     1019       552.5
```

- The highest number of cases is for PCI and Craniotomy procedures, Acute Stroke, Heart Failure, AMI and GI Hemorrhage conditions.
- The highest number of deaths is for Craniotomy and PCI procedures, Acute Stroke, AMI and Heart Failure conditions.
- The highest mortality rates is for Craniotomy and Pancreatic Resection procedures, Acute Stroke, AMI and Heart Failure conditions.
- The lowest number of cases is for Esophageal Resection procedure, Pancreatic Cancer and Pancreatic Other conditions.
- The lowest number of deaths is for Esophageal Resection procedure, Pancreatic Cancer and Pancreatic Other conditions.
- The lowest mortality rates is for Carotid Endarterectomy procedure, Pancreatic Other and Pneumonia conditions.

## Hospital Ratings.

```
prop.table(table(df_clean$Procedure.Condition,df_clean$Hospital.Ratings))*100
```

```
##
##           As Expected      Better      Worse
## AAA Repair      5.80218873  0.00000000  0.04129672
## Acute Stroke    10.81973983  1.01176956  0.90852777
## AMI             11.06752013  0.47491224  0.64009911
## Carotid Endarterectomy 8.19739831  0.00000000  0.14453851
## Craniotomy      5.40986992  0.37167045  0.37167045
## Esophageal Resection 1.54862688  0.00000000  0.00000000
## GI Hemorrhage   12.28577328  0.20648358  0.35102209
## Heart Failure   11.48048730  0.47491224  0.76398926
## Hip Fracture    8.71360727  0.00000000  0.08259343
```



##	Pancreatic Cancer	2.89077018	0.00000000	0.04129672
##	Pancreatic Other	2.64298988	0.00000000	0.04129672
##	Pancreatic Resection	3.84059467	0.00000000	0.08259343
##	PCI	5.78154037	0.12389015	0.26842866
##	Pneumonia	3.03530869	0.08259343	0.00000000

- Better ratings are for Craniotomy procedure, Acute Stroke, AMI and Heart Failure conditions.
- Worse ratings are for Craniotomy and PCI procedures, Acute Stroke, AMI, GI Hemorrhage and Heart Failure conditions.
- As Expected ratings are for Acute Stroke, AMI, GI Hemorrhage and Heart Failure conditions.

## Conclusions 2:

- There is **association** between the risk adjusted mortality rate and hospital ratings.
- Lower the risk adjusted mortality rate, better the hospital ratings.
- Higher the risk adjusted mortality rate, worse the hospital ratings.
- **Procedures:**
  - with severe outcomes: PCI, Craniotomy and Pancreatic Resection.
  - with good outcomes: Esophageal Resection and Carotid Endarterectomy.
- **Conditions:**
  - with severe outcomes: Acute Stroke, AMI, Heart Failure and GI Hemorrhage.
  - with good outcomes: Pancreatic Cancer, Pancreatic Other and Pneumonia.

## Mapping and summary of overall hospital quality ratings and mean mortality rate among all conditions and procedures.

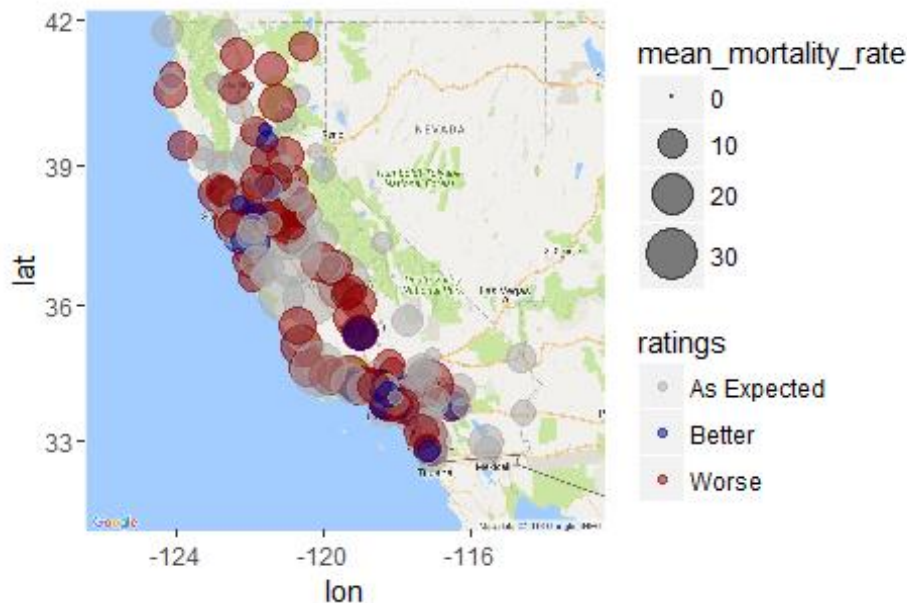
### Summary of hospital ratings over all conditions and procedures.

```
all_ratings <- df_clean %>%
  group_by(Hospital, Latitude, Longitude) %>%
  summarise(all_ratings = 0.5 * sum(ratings), # 0.5 to account
    for 2012 and 2013
    mean_mortality_rate = mean(Risk.Adjusted.Mortality.
Rate)) %>%
  mutate(ratings =
    ifelse(all_ratings > 0, "Better",
    ifelse(all_ratings < 0, "Worse", "As Expected")))
all_ratings$ratings <- as.factor(all_ratings$ratings)
all_ratings <- tbl_df(all_ratings)
```

### Mapping of overall hospital ratings and mean mortality rates.

```
CAmap <- get_map(location="California", source="google", maptype="roadmap", crop
=FALSE, zoom=6)
ggmap (CAmap) +
  geom_point(aes(x=Longitude, y=Latitude, size=mean_mortality_rate, colour=rating))
```

```
gs),data=all_ratings,alpha=0.5)+
  scale_colour_manual(values=c("Worse" = "darkred","Better" = "darkblue","As
Expected" = "darkgrey"))+
  scale_size(range = c(0, 10))
```



### Overall Hospital Ratings:

```
summary(all_ratings$ratings)
```

```
## As Expected      Better      Worse
##           172           69           99
```

- Top 5 hospitals with the **best** quality ratings:

```
all_ratings %>% arrange(desc(all_ratings)) %>% select(Hospital) %>% slice(1:5)
```

```
## # A tibble: 5 x 1
```

```
##                               Hospital
##                               <fctr>
## 1 Kaiser Foundation Hospital â Redwood City
## 2 Kaiser Foundation Hospital â Sunset
## 3 Centinela Hospital Medical Center
## 4 Scripps Green Hospital
## 5 Cedars Sinai Medical Center
```

- Top 5 hospitals with the **lowest** mean mortality rate:

```
all_ratings %>% arrange(mean_mortality_rate) %>% select(Hospital) %>% slice(1:5)
```

```
## # A tibble: 5 x 1
##               Hospital
##             <fctr>
## 1 Corcoran District Hospital
## 2 Eastern Plumas Hospital â Portola Campus
## 3           Glenn Medical Center
## 4 Good Samaritan Hospital â Bakersfield
## 5 Hoag Orthopedic Institute
```

- Top 5 hospitals with the **worst** quality ratings:

```
all_ratings %>% arrange(all_ratings) %>% select(Hospital) %>% slice(1:5)
```

```
## # A tibble: 5 x 1
##               Hospital
##             <fctr>
## 1 Palomar Health Downtown Campus
## 2 Los Angeles County/University of Southern California Medical Center
## 3 San Francisco General Hospital
## 4 Santa Barbara Cottage Hospital
## 5 Arrowhead Regional Medical Center
```

- Top 5 hospitals with the **highest** mean mortality rate:

```
all_ratings %>% arrange(desc(mean_mortality_rate)) %>% select(Hospital) %>% slice(1:5)
```

```
## # A tibble: 5 x 1
##               Hospital
##             <fctr>
## 1 Memorial Hospital Los Banos
## 2 Central Valley Specialty Hospital
## 3 Coalinga Regional Medical Center
## 4 Santa Ynez Valley Cottage Hospital
## 5 Biggs Gridley Memorial Hospital
```

## Summary of hospital quality ratings and mortality rates for Acute Stroke, AMI and Heart Failure conditions, PCI, Craniotomy and Pancreatic Resection procedures.

```
df_as <- df_c[which(df_c$Procedure.Condition=="Acute Stroke"),]
```

```
df_as_all <- df_as %>%
  group_by(Hospital, Latitude, Longitude) %>%
  summarise(all_ratings = 0.5 * sum(ratings), # to account for
2012 and 2013
            mean_mortality_rate = mean(Risk.Adjusted.Mortality.
Rate)) %>%
  mutate(ratings =
         ifelse(all_ratings > 0, "Better",
```

```

      ifelse(all_ratings < 0, "Worse", "As Expected")),
      Procedure.Condition="Acute Stroke")

df_as_all$ratings <- as.factor(df_as_all$ratings)
df_as_all <- tbl_df(df_as_all)

df_as_best_rat <- df_as_all %>% arrange(desc(all_ratings)) %>% slice(1:50)
df_as_best <- df_as_best_rat %>% arrange(mean_mortality_rate) %>% slice(1:25)

```

### Top 5 hospitals for treatment of Acute Stroke condition.

```

df_as_best %>% slice(1:5) %>% select(Hospital)

## # A tibble: 5 x 1
##                               Hospital
##                               <fctr>
## 1 Pacific Alliance Medical Center, Inc.
## 2 Anaheim General Hospital
## 3 Olympia Medical Center
## 4 Encino Hospital Medical Center
## 5 Los Angeles County/Olive View â UCLA Medical Center

```

### Top 5 hospitals for treatment of AMI condition.

```

## # A tibble: 5 x 1
##                               Hospital
##                               <fctr>
## 1 Southern California Hospital at Hollywood
## 2 Sherman Oaks Hospital
## 3 Kaiser Foundation Hospital â Antioch
## 4 Encino Hospital Medical Center
## 5 La Palma Intercommunity Hospital

```

### Top 5 hospitals for treatment of Heart Failure condition.

```

## # A tibble: 5 x 1
##                               Hospital
##                               <fctr>
## 1 Adventist Medical Center â Reedley
## 2 Anaheim General Hospital
## 3 Sherman Oaks Hospital
## 4 Barstow Community Hospital
## 5 Centinela Hospital Medical Center

```

### Top 5 hospitals to perform the PCI procedure.

```

## # A tibble: 5 x 1
##                               Hospital
##                               <fctr>
## 1 Brotman Medical Center
## 2 Fresno Heart and Surgical Hospital
## 3 El Camino Hospital

```

```
## 4 Downey Regional Medical Center
## 5 Henry Mayo Newhall Memorial Hospital
```

### Top 5 hospitals to perform the Craniotomy procedure.

```
## # A tibble: 5 x 1
##           Hospital
##           <fctr>
## 1 Alhambra Hospital
## 2 Desert Valley Hospital
## 3 El Centro Regional Medical Center
## 4 Saint John's Health Center
## 5 El Camino Hospital
```

### Top 5 hospitals to perform the Pancreatic Resection procedure.

```
## # A tibble: 5 x 1
##           Hospital
##           <fctr>
## 1 Alameda County Medical Center
## 2 Community Hospital Monterey Peninsula
## 3 Community Hospital of The Monterey Peninsula
## 4 Community Memorial Hospital â San Buenaventura
## 5 Eden Medical Center
```

## Predictions

### Approach

- Predict hospital quality ratings using **random forests and classification decision trees**.
- Train the models and evaluate the model performances on 2012 training data.
- Test the model performances on 2013 test data.

### Hospital Ratings Prediction Using Random Forests for Dataset in Wide Format.

#### Cleanning the Data Set and converting to the wide format based on Procedure.Condition and Risk.Adjusted.Mortality.Rate variables.

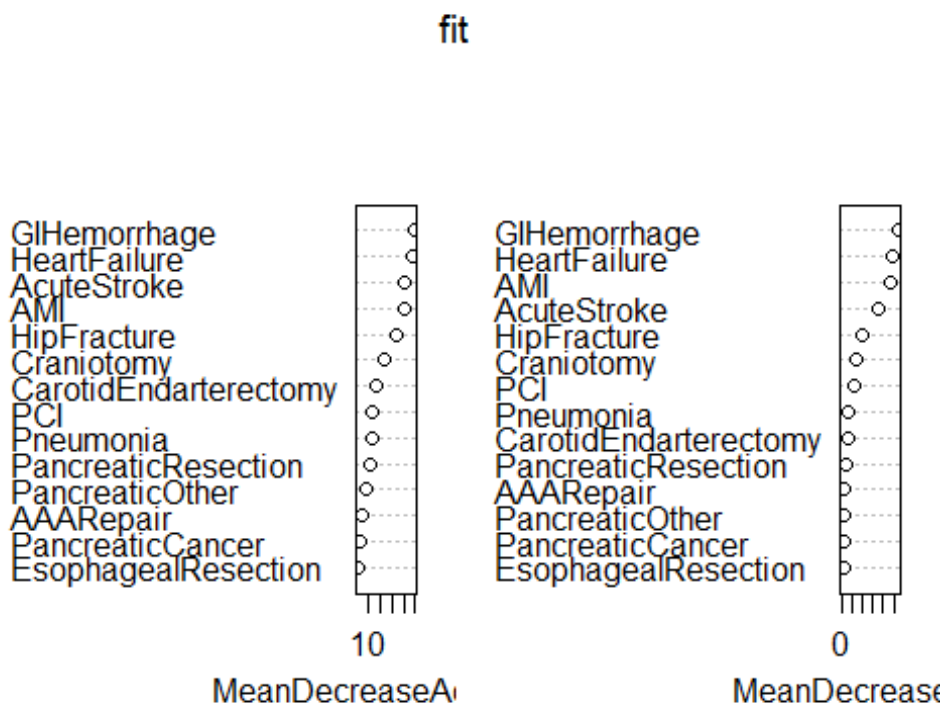
```
# convert data to the wide format
df_wide <- df_clean %>% select(Year,Hospital,Latitude,Longitude,Procedure.Condition,Hospital.Ratings,Risk.Adjusted.Mortality.Rate) %>% spread(Procedure.Condition,Risk.Adjusted.Mortality.Rate)
# remove white spaces from column names
colnames(df_wide) <- gsub(" ", "", colnames(df_wide))
# replace NA with 0, because some hospitals does not treat these conditions, thus mortality rate is zero.
df_wide[is.na(df_wide)] <- 0
```

### Split the Data Set into 2012 training and 2013 test sets.

```
train_wide <- df_wide[which(df_wide$Year==2012),]
test_wide_original <- df_wide[which(df_wide$Year==2013),]
test_wide <- subset(test_wide_original, select = -Hospital.Ratings)
```

## Feature Engineering with Random Forests

```
fit <- randomForest(Hospital.Ratings ~ AAARepair + AcuteStroke + AMI + Caroti
dEndarterectomy + Craniotomy + EsophagealResection + GIHemorrhage + HeartFail
ure + HipFracture + PancreaticCancer + PancreaticOther + PancreaticResection
+ PCI + Pneumonia, data=train_wide,importance=TRUE,ntree=1000)
varImpPlot(fit)
```



- **The most important variables are** Heart Failure, GI Hemorrhage, AMI and Acute Stroke, Hip Fracture **conditions**;
- **Procedure** variables are less important and thus are not included in classification.

**Model performance on train\_wide dataset using all variables.**

```
# confusion matrix on train data
fit$confusion

##           As Expected Better Worse class.error
## As Expected          296      18      12  0.09202454
## Better                13      31       3  0.34042553
## Worse                 25       4      36  0.44615385
```

- Accuracy (how often is the classifier correct): 0.827108

- Error Rate (how often is the classifier wrong): 0.172892

**Predictions on test\_wide dataset using all variables.**

```
prediction <- predict(fit, test_wide)
# confusion matrix on test data
cm <- as.matrix(table(Actual = test_wide_original$Hospital.Ratings, Predicted = prediction))
cm

##               Predicted
## Actual           As Expected Better Worse
## As Expected           305         6    13
## Better                15        31     5
## Worse                 26         0    42

rf_a <- sum(diag(cm))/sum(cm)
rf_e <- 1 - sum(diag(cm))/sum(cm)
```

- Accuracy: 0.8532731
- Error Rate: 0.1467269

### Hospital Ratings Prediction Using Classification Decision Trees (CART).

**Model 1: All variables are included in tree construction.**

```
set.seed(34)
tree0 <- rpart(Hospital.Ratings ~ AAARepair + AcuteStroke + AMI + CarotidEndarterectomy + Craniotomy + EsophagealResection + GIHemorrhage + HeartFailure + HipFracture + PancreaticCancer + PancreaticOther + PancreaticResection + PCI + Pneumonia, data = train_wide, method = "class", control=rpart.control(cp=0.001))
printcp(tree0)

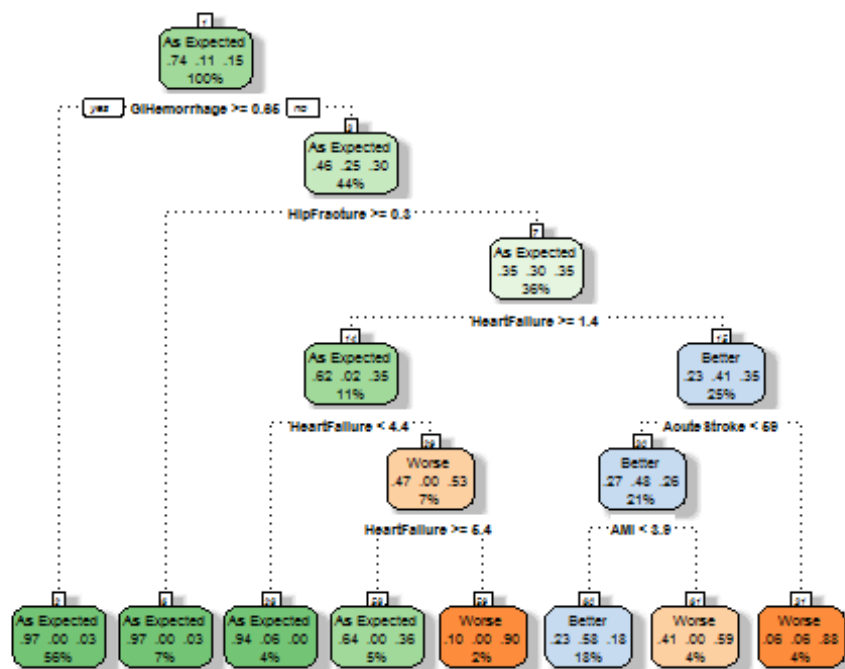
##
## Classification tree:
## rpart(formula = Hospital.Ratings ~ AAARepair + AcuteStroke +
##       AMI + CarotidEndarterectomy + Craniotomy + EsophagealResection +
##       GIHemorrhage + HeartFailure + HipFracture + PancreaticCancer +
##       PancreaticOther + PancreaticResection + PCI + Pneumonia,
##       data = train_wide, method = "class", control = rpart.control(cp = 0.001))
##
## Variables actually used in tree construction:
## [1] AcuteStroke AMI Craniotomy GIHemorrhage HeartFailure
## [6] HipFracture
##
## Root node error: 112/438 = 0.25571
##
## n= 438
##
```

```
##          CP nsplit rel error  xerror    xstd
## 1 0.0595238    0   1.00000 1.00000 0.081520
## 2 0.0357143    5   0.60714 0.70536 0.071847
## 3 0.0089286    7   0.53571 0.63393 0.068866
## 4 0.0059524    8   0.52679 0.67857 0.070763
## 5 0.0010000   11   0.50893 0.67857 0.070763

num <- which.min(tree0$cptable[, "xerror"])
tree0$cptable[num,]

##          CP      nsplit    rel error      xerror      xstd
## 0.008928571 7.000000000 0.535714286 0.633928571 0.068866360

cp.choice<-tree0$cptable[num,"CP"]
pruned.tree<-prune(tree0, cp=cp.choice)
fancyRpartPlot(pruned.tree)
```



Rattle 2016-Aug-15 21:40:32 postdoc

**Predictions on test\_wide dataset.**

```
# Make predictions on the test set
prediction <- predict(pruned.tree, test_wide, type = "class")
# confusion matrix
cm <- as.matrix(table(Actual = test_wide_original$Hospital.Ratings, Predicted
= prediction))
cm

##          Predicted
## Actual    As Expected Better Worse
```



```
## As Expected      287      24      13
## Better           4       44       3
## Worse            26      13      29
```

```
call_a <- sum(diag(cm))/sum(cm)
call_e <- 1 - sum(diag(cm))/sum(cm)
```

- Accuracy: 0.8126411
- Error Rate: 0.1873589

**Model 2: AMI, GIHemorrhage and HeartFailure variables are included in tree construction.**

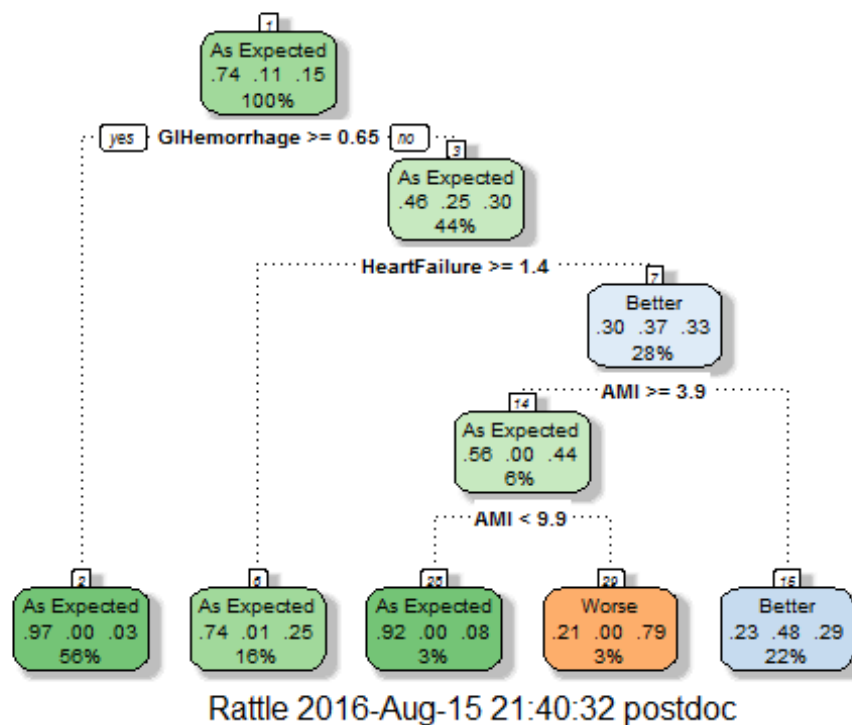
```
tree1 <- rpart(Hospital.Ratings ~ AMI + GIHemorrhage + HeartFailure, data = t
rain_wide, method = "class", control=rpart.control(cp=0.001)) # cp determines
when the splitting up of the decision tree stops
printcp(tree1)
```

```
##
## Classification tree:
## rpart(formula = Hospital.Ratings ~ AMI + GIHemorrhage + HeartFailure,
##       data = train_wide, method = "class", control = rpart.control(cp = 0.00
1))
##
## Variables actually used in tree construction:
## [1] AMI          GIHemorrhage HeartFailure
##
## Root node error: 112/438 = 0.25571
##
## n= 438
##
##      CP nsplit rel error  xerror    xstd
## 1 0.0714286      0  1.00000 1.00000 0.081520
## 2 0.0267857      4   0.71429 0.76786 0.074228
## 3 0.0089286      6   0.66071 0.80357 0.075502
## 4 0.0044643      7   0.65179 0.86607 0.077590
## 5 0.0010000      9   0.64286 0.85714 0.077303
```

```
num <- which.min(tree1$cptable[, "xerror"])
tree1$cptable[num,]
```

```
##      CP      nsplit rel error    xerror    xstd
## 0.02678571 4.00000000 0.71428571 0.76785714 0.07422761
```

```
cp.choice<-tree1$cptable[num, "CP"]
pruned.tree1<-prune(tree1, cp=cp.choice)
fancyRpartPlot(pruned.tree1)
```



**Predictions on test\_wide dataset.**

```
# Make predictions on the test set
prediction <- predict(pruned.tree1, test_wide, type = "class")
# confusion matrix
cm <- as.matrix(table(Actual = test_wide_original$Hospital.Ratings, Predicted = prediction))
cm

##               Predicted
## Actual         As Expected Better Worse
## As Expected      292      27     5
## Better           5      46     0
## Worse           29      30     9

c3_a <- sum(diag(cm))/sum(cm)
c3_e <- 1 - sum(diag(cm))/sum(cm)
```

- Accuracy: 0.7832957
- Error Rate: 0.2167043

### Conclusions 3:

- **Accuracy on the test data set using**
  - Random Forests with all variables: 0.8533
  - CART with all variables: 0.8126
  - CART with three variables: 0.7833

- **Random forests** gives the best performance, however is not good enough to predict hospitals with the best care in future.
- **Random forests** predicts that classification of **hospital ratings** depend on **conditions and not procedures** with the most severe patient outcomes.

## Future Work

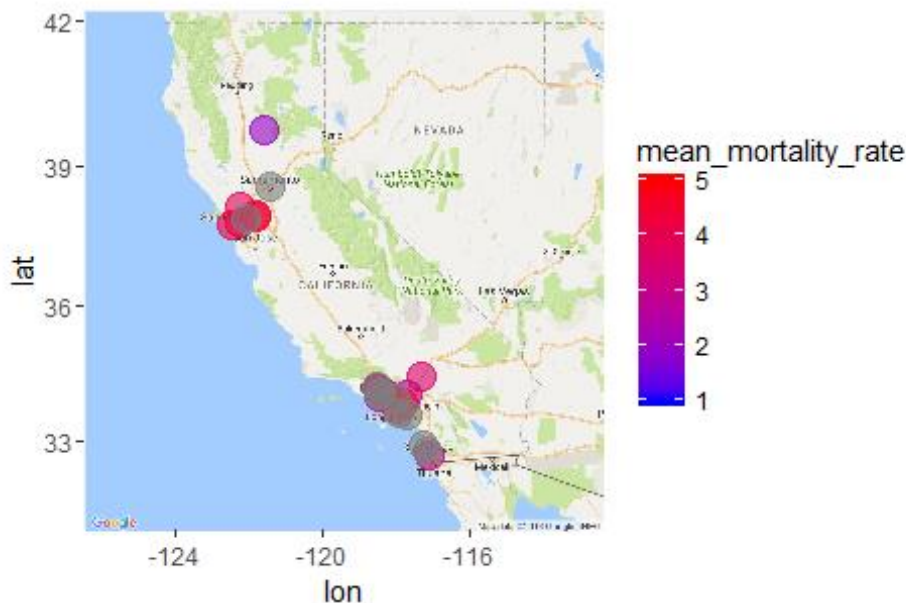
- Predict hospital quality ratings using **multinomial logistic regression**.
  - Train the model and evaluate the model performance on 2012 training data.
  - Test the model performance on 2013 test data.
- Compare three models: random forests, classification decision trees and multinomial logistic regression.
  - Summarize which model gives the best performance on 2012 training data and on 2013 test data.
  - Choose the best model and test its performance on [2014 test data](#).
- Recommend which hospitals will have the best care in future using predicted hospital ratings.

## Recommendations to Patients

### Top 25 hospitals with the best overall ratings and the lowest mean mortality rate in state of California.

```
best_ratings <- all_ratings %>% arrange(desc(all_ratings)) %>% slice(1:50)
best_lowest <- best_ratings %>% arrange(mean_mortality_rate) %>% slice(1:25)
# best_lowest$Hospital[duplicated(best_lowest$Hospital)]

CAmap <- get_map(location="California",source="google",maptype="roadmap",crop
=FALSE,zoom=6)
ggmap (CAmap) +
  geom_point(aes(x=Longitude,y=Latitude,colour=mean_mortality_rate),data=best
_lowest,size=5,alpha=0.6)+
  scale_colour_gradient(limits=c(1, 5), high="red", low="blue")
```



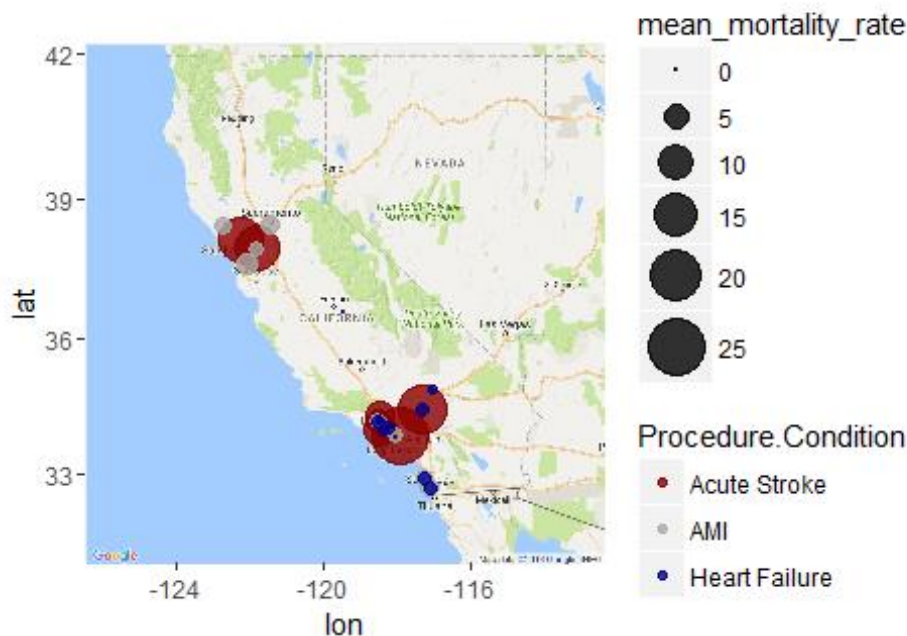
**Top 5 hospitals with the best overall ratings and the lowest mean mortality rate in state of California.**

```
best_lowest$Hospital[1:5]
## [1] Encino Hospital Medical Center Feather River Hospital
## [3] Chino Valley Medical Center Marina Del Rey Hospital
## [5] Paradise Valley Hospital
## 341 Levels: Adventist Medical Center ...
```

**Top hospitals with the best ratings and the lowest mean mortality rate for Acute Stroke, AMI and Heart Failure conditions.**

```
best_cond <- bind_rows(df_as_best[1:10,],df_ami_best[1:10,],df_hf_best[1:10,])

CMap <- get_map(location="California",source="google",maptype="roadmap",crop=FALSE,zoom=6)
ggmap (CMap) +
  geom_point(aes(x=Longitude,y=Latitude,size=mean_mortality_rate,colour=Procedure.Condition),data=best_cond,alpha=0.8)+
  scale_colour_manual(values=c("Acute Stroke"="darkred", "AMI"="darkgrey", "Heart Failure"="darkblue"))+
  scale_size(range = c(0, 10))
```



**There are 7 hospitals that have the best ratings and the lowest mortality rate for the most severe conditions.**

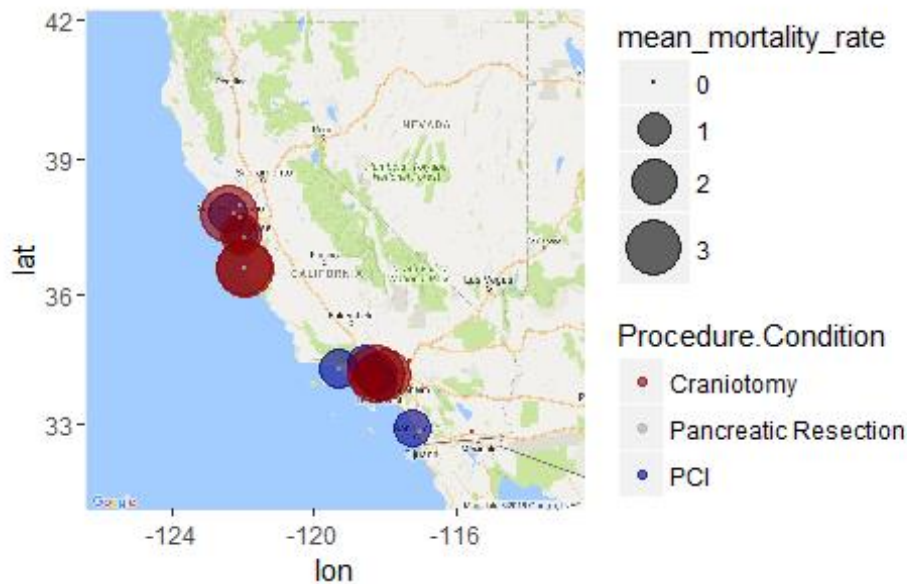
```
best_cond$Hospital[duplicated(best_cond$Hospital)]

## [1] Encino Hospital Medical Center Anaheim General Hospital
## [3] Sherman Oaks Hospital Encino Hospital Medical Center
## [5] Desert Valley Hospital Paradise Valley Hospital
## [7] Scripps Green Hospital
## 341 Levels: Adventist Medical Center ...
```

**Top hospitals with the best ratings and the lowest mean mortality rate for PCI, Craniotomy and Pancreatic Resection procedures.**

```
best_proc <- bind_rows(df_pci_best[1:10,],df_cr_best[1:10,],df_pr_best[1:10,])

CMap <- get_map(location="California",source="google",maptype="roadmap",crop=FALSE,zoom=6)
ggmap(CMap) +
  geom_point(aes(x=Longitude,y=Latitude,size=mean_mortality_rate,colour=Procedure.Condition),data=best_proc,alpha=0.6)+
  scale_colour_manual(values=c("PCI"="darkblue", "Craniotomy"="darkred", "Pancreatic Resection"="darkgrey"))+
  scale_size(range = c(0, 10))
```



**There are 7 hospitals that have the best ratings and the lowest mortality rate for the most severe procedures.**

```
best_proc$Hospital[duplicated(best_proc$Hospital)]

## [1] El Camino Hospital
## [2] California Pacific Medical Center â Pacific Campus
## [3] Glendale Adventist Medical Center â Wilson Terrace
## [4] Community Hospital Monterey Peninsula
## [5] Community Hospital of The Monterey Peninsula
## [6] Community Memorial Hospital â San Buenaventura
## [7] Fresno Heart and Surgical Hospital
## 341 Levels: Adventist Medical Center ...
```