

WisconsinBreastCancer

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- a. Major predictors of diagnosis are concave_points, concavity_points_mean, area mean, radius_mean, parameter_mean, area, radius, and perimeter. There are no missing values in the data set. There are outliers. The outliers have been eliminated using Z-score method.
- b. 5 leaves in the Decision Tree
- c. Major predictors suggested by our tree are: Perimeter, Concave_points_mean, texture
- d. Firstly, eliminated the outliers using Z-score and obtained 398 observations. Then to enhance the accuracy cp value has been changed. When cp = 0.04, accuracy of test was 87.5%, when cp was changed to 0.01, Accuracy for Train = 94.49%
- e. Accuracy for Train: 94.49% Accuracy for Test: 94.64%
- f. Initially, we clean the data for outliers by using the z-score method. Then we combine the trainx and trainy variable to obtain a matrix with the diagnosis. For this new combined variable we applied the decision tree classification. From this, we applied the testx and testy data to the created model and obtain the result. To support the same we have plotted a bivariate analysis and the importance of variable bar graph.
- g. From the importance of variable graph, we get the most important variables from which the tree can be affected. Here we used, perimeter_lv, radius_lv, area_lv and perimeter.

```
library("rpart")
library("ggplot2")
library("tidyverse")
```

```
## — Attaching packages ————— tidyverse 1.3.2 —
—
## ✓ tibble 3.1.8      ✓ dplyr 1.0.10
## ✓ tidyr 1.2.0      ✓ stringr 1.4.1
## ✓ readr 2.1.2      ✓ forcats 0.5.2
## ✓ purrr 0.3.4
## — Conflicts ————— tidyverse_conflicts() —
—
## ✖ dplyr::filter() masks stats::filter()
## ✖ dplyr::lag() masks stats::lag()
```

```
library("psych")
```

```
##
## Attaching package: 'psych'
##
## The following objects are masked from 'package:ggplot2':
##
##      %+%, alpha
```

```
library("corrplot")
```

```
## corrplot 0.92 loaded
```

```
library("RColorBrewer")
```

```
#Load Dataset
```

```
trainx=read.csv("/Users/abhinavram/Documents/IDS572 Data Mining/Assignment 1/train
X.csv")
trainy=read.csv("/Users/abhinavram/Documents/IDS572 Data Mining/Assignment 1/train
Y.csv")
testx=read.csv("/Users/abhinavram/Documents/IDS572 Data Mining/Assignment 1/testX.
csv")
testy=read.csv("/Users/abhinavram/Documents/IDS572 Data Mining/Assignment 1/testY.
csv")
```

```
#Column Names
```

```
colnames(trainx)<-c('radius','texture','perimeter','area','smoothness','compactnes
s','concavity','no_of_concave_contour','symmetry','fractal_dim',
                    'radius_sd','texture_sd','perimeter_sd','area_sd','smoothness_sd'
,'compactness_sd','concavity_sd','no_of_concave_contour_sd','symmetry_sd','fract
al_dim_sd',
                    'radius_lv','texture_lv','perimeter_lv','area_lv','smoothness_lv'
,'compactness_lv','concavity_lv','no_of_concave_contour_lv','symmetry_lv','fract
al_dim_lv')
colnames(testx)<-c('radius','texture','perimeter','area','smoothness','compactness
','concavity','no_of_concave_contour','symmetry','fractal_dim',
                  'radius_sd','texture_sd','perimeter_sd','area_sd','smoothness_sd'
,'compactness_sd','concavity_sd','no_of_concave_contour_sd','symmetry_sd','fract
al_dim_sd',
                  'radius_lv','texture_lv','perimeter_lv','area_lv','smoothness_lv'
,'compactness_lv','concavity_lv','no_of_concave_contour_lv','symmetry_lv','fract
al_dim_lv')
```

Summary of dataset

```
summary(trainx)
```

```
##      radius      texture      perimeter      area
## Min.   : 6.981   Min.    : 9.71   Min.     : 43.79   Min.    : 143.5
## 1st Qu.:11.613   1st Qu.:16.21   1st Qu.: 74.69   1st Qu.: 412.5
```

##	Median :13.280	Median :18.82	Median : 86.04	Median : 545.6
##	Mean :14.107	Mean :19.38	Mean : 91.87	Mean : 655.0
##	3rd Qu.:15.832	3rd Qu.:21.91	3rd Qu.:103.78	3rd Qu.: 787.0
##	Max. :28.110	Max. :39.28	Max. :188.50	Max. :2501.0
##	smoothness	compactness	concavity	no_of_concave_contour
##	Min. :0.05263	Min. :0.01938	Min. :0.00000	Min. :0.00000
##	1st Qu.:0.08609	1st Qu.:0.06609	1st Qu.:0.03038	1st Qu.:0.01979
##	Median :0.09572	Median :0.09449	Median :0.06406	Median :0.03387
##	Mean :0.09620	Mean :0.10511	Mean :0.09011	Mean :0.04895
##	3rd Qu.:0.10460	3rd Qu.:0.13057	3rd Qu.:0.13205	3rd Qu.:0.07403
##	Max. :0.16340	Max. :0.34540	Max. :0.42680	Max. :0.20120
##	symmetry	fractal_dim	radius_sd	texture_sd
##	Min. :0.1060	Min. :0.04996	Min. :0.1115	Min. :0.3621
##	1st Qu.:0.1623	1st Qu.:0.05797	1st Qu.:0.2319	1st Qu.:0.8280
##	Median :0.1799	Median :0.06177	Median :0.3175	Median :1.1080
##	Mean :0.1808	Mean :0.06299	Mean :0.4022	Mean :1.2278
##	3rd Qu.:0.1957	3rd Qu.:0.06638	3rd Qu.:0.4702	3rd Qu.:1.4797
##	Max. :0.2906	Max. :0.09744	Max. :2.8730	Max. :4.8850
##	perimeter_sd	area_sd	smoothness_sd	compactness_sd
##	Min. : 0.757	Min. : 6.802	Min. :0.001713	Min. :0.002252
##	1st Qu.: 1.643	1st Qu.: 17.670	1st Qu.:0.005227	1st Qu.:0.013710
##	Median : 2.280	Median : 23.930	Median :0.006457	Median :0.021145
##	Mean : 2.857	Mean : 40.205	Mean :0.007163	Mean :0.026019
##	3rd Qu.: 3.309	3rd Qu.: 44.947	3rd Qu.:0.008391	3rd Qu.:0.032888
##	Max. :21.980	Max. :542.200	Max. :0.031130	Max. :0.135400
##	concavity_sd	no_of_concave_contour_sd	symmetry_sd	
##	Min. :0.00000	Min. :0.000000	Min. :0.007882	
##	1st Qu.:0.01569	1st Qu.:0.007735	1st Qu.:0.015220	
##	Median :0.02625	Median :0.011000	Median :0.018975	
##	Mean :0.03285	Mean :0.011880	Mean :0.020695	
##	3rd Qu.:0.04273	3rd Qu.:0.014897	3rd Qu.:0.023768	
##	Max. :0.39600	Max. :0.052790	Max. :0.078950	
##	fractal_dim_sd	radius_lv	texture_lv	perimeter_lv
##	Min. :0.0008948	Min. : 7.93	Min. :12.02	Min. : 50.41
##	1st Qu.:0.0022787	1st Qu.:12.97	1st Qu.:21.09	1st Qu.: 83.77
##	Median :0.0032335	Median :14.88	Median :25.43	Median : 97.39
##	Mean :0.0038669	Mean :16.22	Mean :25.77	Mean :106.95
##	3rd Qu.:0.0045708	3rd Qu.:18.71	3rd Qu.:29.89	3rd Qu.:125.78
##	Max. :0.0298400	Max. :36.04	Max. :49.54	Max. :251.20
##	area_lv	smoothness_lv	compactness_lv	concavity_lv
##	Min. : 185.2	Min. :0.07117	Min. :0.02729	Min. :0.0000
##	1st Qu.: 509.8	1st Qu.:0.11728	1st Qu.:0.14860	1st Qu.:0.1202
##	Median : 674.0	Median :0.13120	Median :0.21685	Median :0.2298
##	Mean : 877.9	Mean :0.13268	Mean :0.25810	Mean :0.2761
##	3rd Qu.:1063.5	3rd Qu.:0.14625	3rd Qu.:0.34190	3rd Qu.:0.3886
##	Max. :4254.0	Max. :0.22260	Max. :1.05800	Max. :1.2520
##	no_of_concave_contour_lv	symmetry_lv	fractal_dim_lv	
##	Min. :0.00000	Min. :0.1565	Min. :0.05504	
##	1st Qu.:0.06321	1st Qu.:0.2510	1st Qu.:0.07222	
##	Median :0.10160	Median :0.2826	Median :0.08042	
##	Mean :0.11484	Mean :0.2904	Mean :0.08456	
##	3rd Qu.:0.16608	3rd Qu.:0.3178	3rd Qu.:0.09217	
##	Max. :0.29030	Max. :0.6638	Max. :0.20750	

Checking for missing values

```
sapply(trainx, function(x) sum(is.na(x)))
```

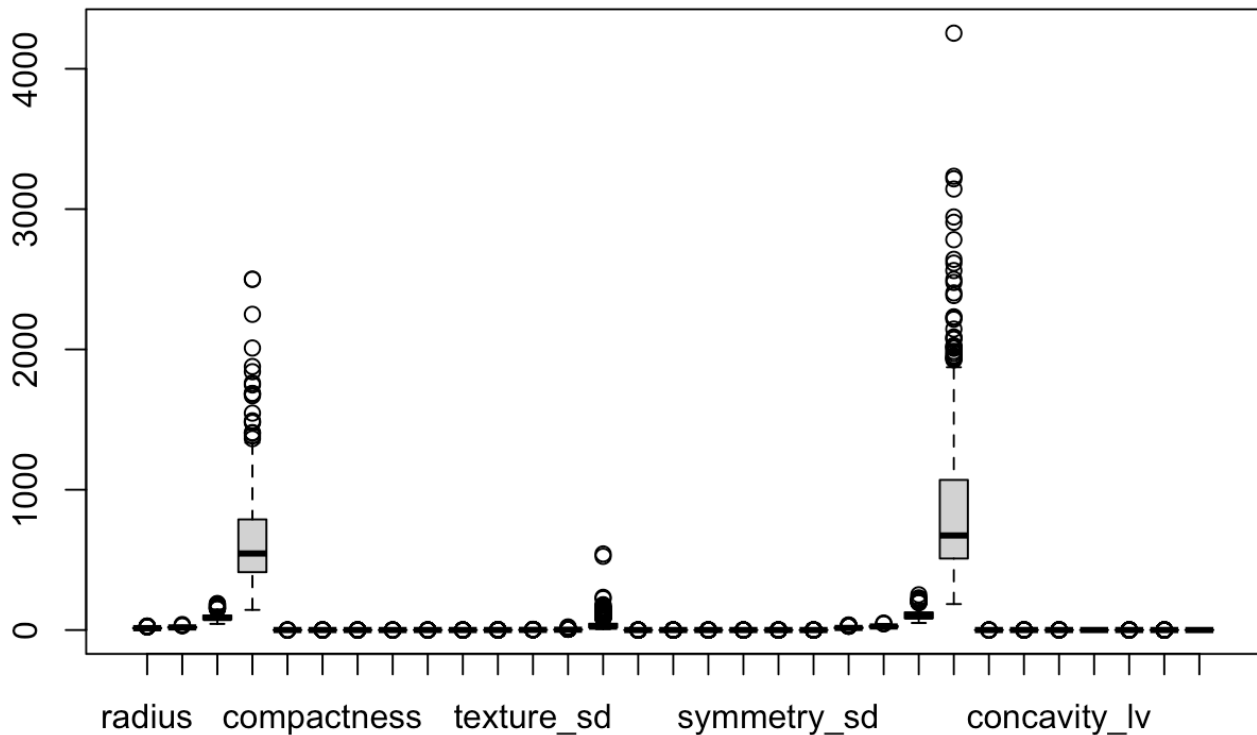
```
##           radius           texture           perimeter
##           0             0             0
##           area           smoothness       compactness
##           0             0             0
##           concavity      no_of_concave_contour      symmetry
##           0             0             0
##           fractal_dim      radius_sd       texture_sd
##           0             0             0
##           perimeter_sd      area_sd       smoothness_sd
##           0             0             0
##           compactness_sd      concavity_sd no_of_concave_contour_sd
##           0             0             0
##           symmetry_sd      fractal_dim_sd      radius_lv
##           0             0             0
##           texture_lv      perimeter_lv      area_lv
##           0             0             0
##           smoothness_lv      compactness_lv      concavity_lv
##           0             0             0
## no_of_concave_contour_lv      symmetry_lv      fractal_dim_lv
##           0             0             0
```

Combine trainx and trainy

```
train_x_y <- cbind(trainx, trainy)
colnames(train_x_y)<-c('radius','texture','perimeter','area','smoothness','compact
ness','concavity','no_of_concave_contour', 'symmetry', 'fractal_dim',
                      'radius_sd','texture_sd','perimeter_sd','area_sd','smoothness_
sd','compactness_sd','concavity_sd','no_of_concave_contour_sd', 'symmetry_sd', 'fr
actal_dim_sd',
                      'radius_lv','texture_lv','perimeter_lv','area_lv','smoothness_
lv','compactness_lv','concavity_lv','no_of_concave_contour_lv', 'symmetry_lv', 'fr
actal_dim_lv','diagnosis')
```

Box Plot for Outliers and cleaning the data

```
boxplot(train_x_y)
```



```
outliers<- as.data.frame(sapply(train_x_y, function(train_x_y) (abs(train_x_y- mea
n(train_x_y))/ sd(train_x_y))))

train_x_y_new <- train_x_y[!rowSums(outliers>3), ]

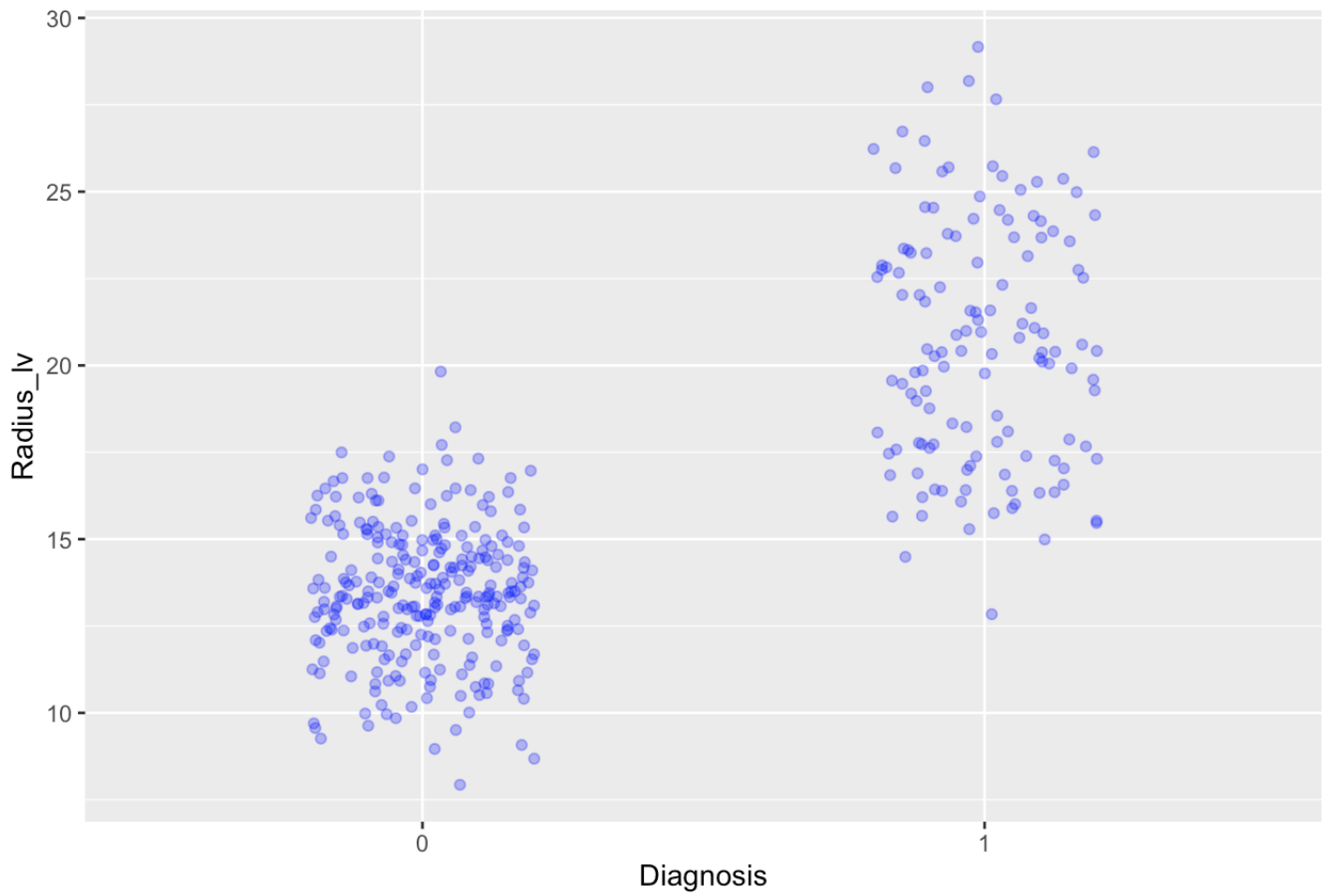
trainx_new = select(train_x_y_new, -diagnosis)
train_x_y_new$diagnosis = as.factor(train_x_y_new$diagnosis)
```

Bivariate Analysis

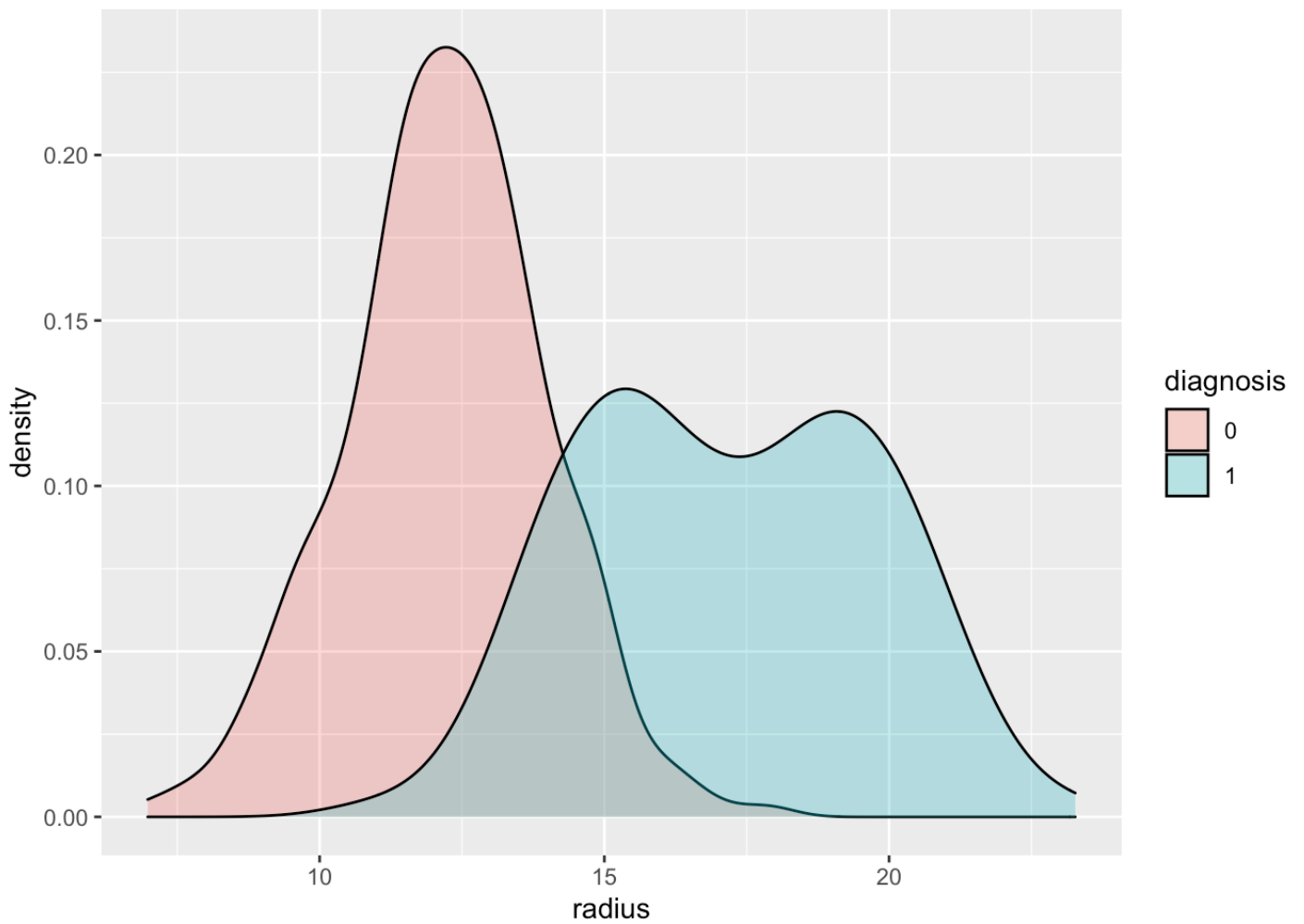
```
ggplot(data=train_x_y_new, aes(x=diagnosis, y=radius_lv, group = 1)) +

  geom_jitter(alpha=0.3,
              color =" blue",
              width = 0.2) +
  labs(title="Wisconsin Breast Cancer", x="Diagnosis", y="Radius_lv")
```

Wisconsin Breast Cancer

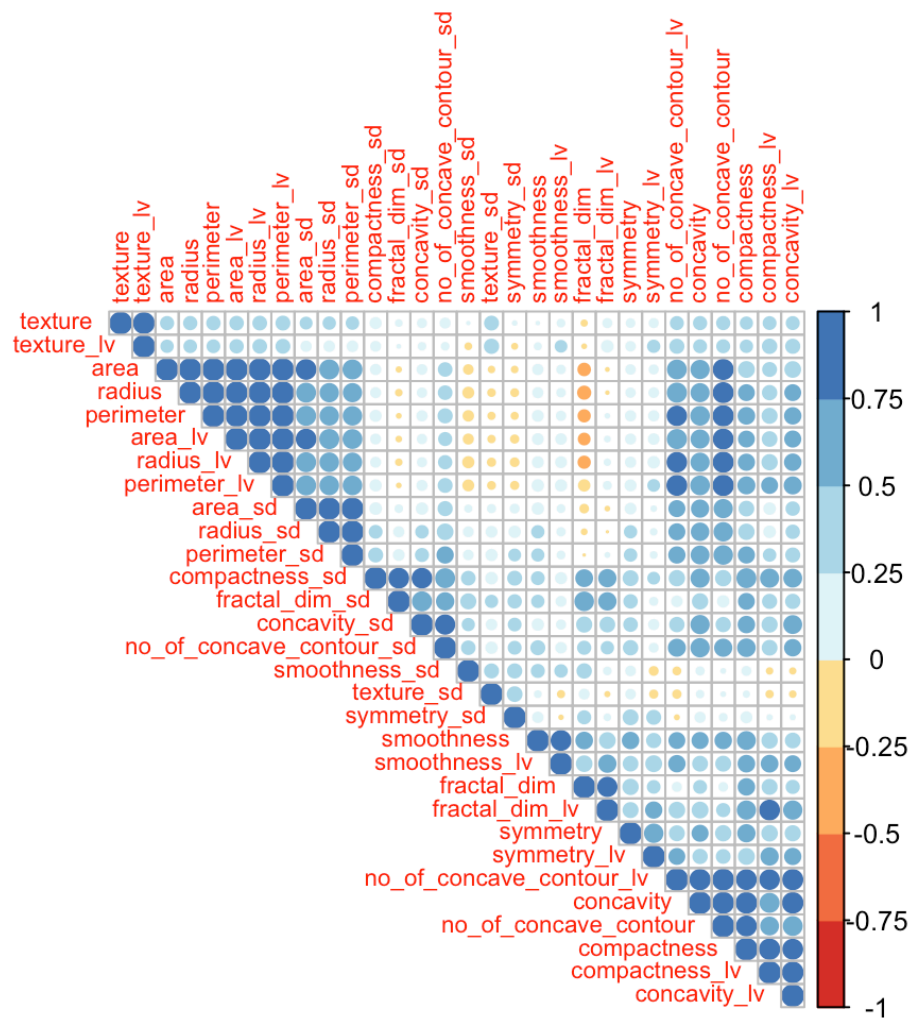


```
ggplot(data=train_x_y_new, aes(x=radius, fill=diagnosis)) +  
  geom_density(alpha=.3)
```



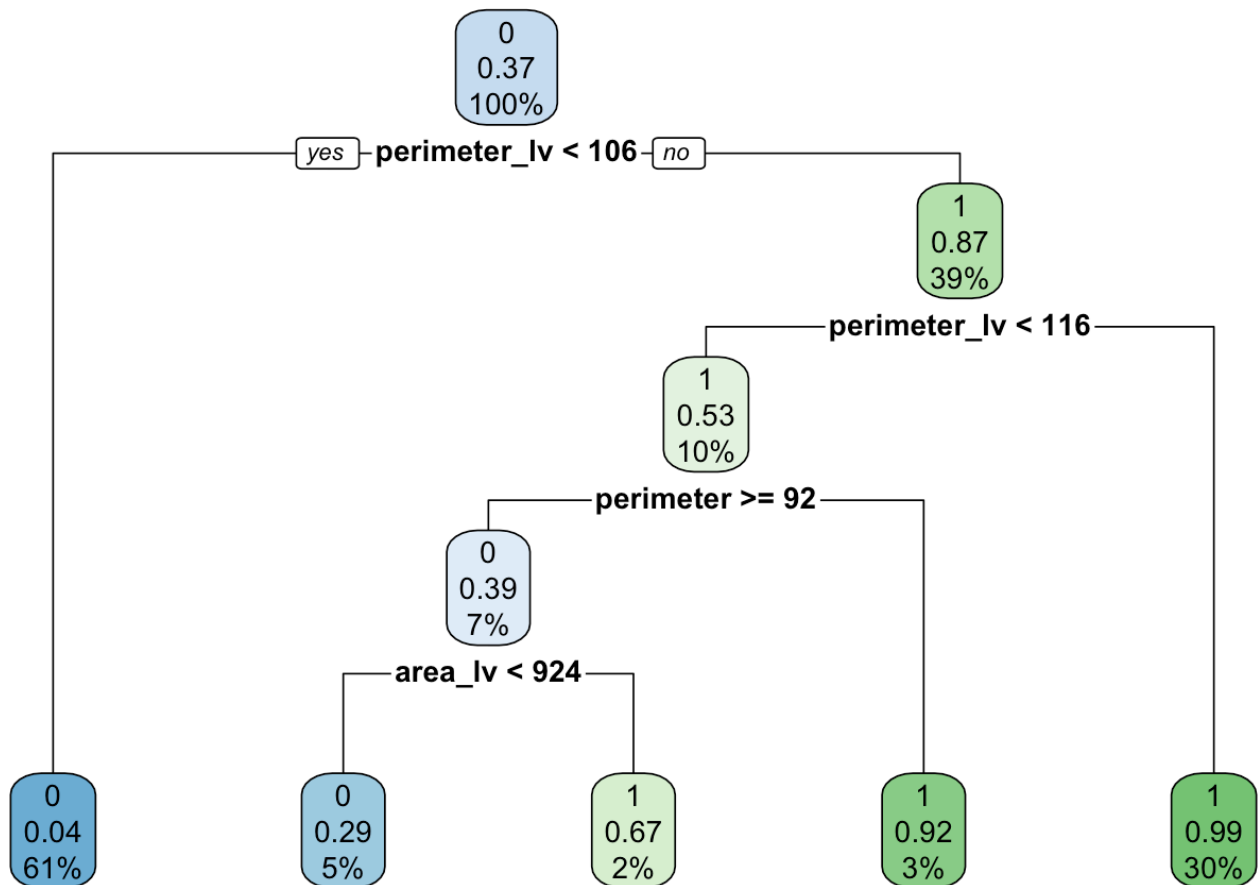
Correlation Plot

```
cor_graph <- cor(trainx)
corrplot(cor_graph, type="upper",order= "hclust", tl.cex = 0.7,col=brewer.pal(n=8,
name="RdYlBu"))
```



Decision Tree

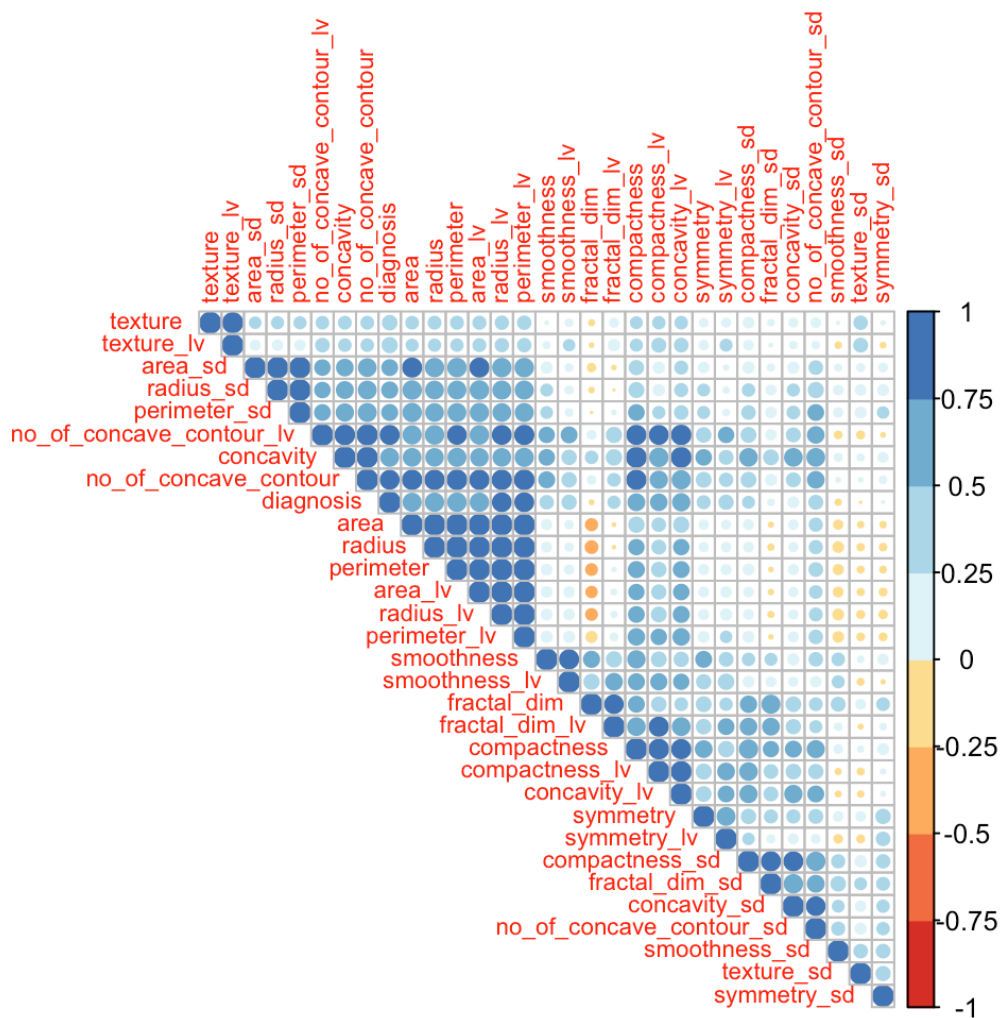
```
library(rpart.plot)
fit=rpart(diagnosis~perimeter_lv + radius_lv + area_lv + perimeter , data=train_x_
y, parms = list(split="information"), method = 'class')
rpart.plot(fit, extra = 106)
```

```

train_x_y$diagnosis <- as.numeric(train_x_y$diagnosis)
cor_graph <- cor(train_x_y)
corrplot(cor_graph, type="upper",order= "hclust",tl.cex = 0.7,col=brewer.pal(n=8,
name="RdYlBu" ))

```



Confusion Matrix for combined

```
t_pred=predict(fit, trainx, type='class')
table(train_x_y$diagnosis, t_pred)
```

```
##      t_pred
##           0    1
## 0 280     6
## 1  19 149
```

```
confusion_mat = table(train_x_y$diagnosis, t_pred)
acc = sum(diag(confusion_mat))/sum(confusion_mat)
print(confusion_mat)
```

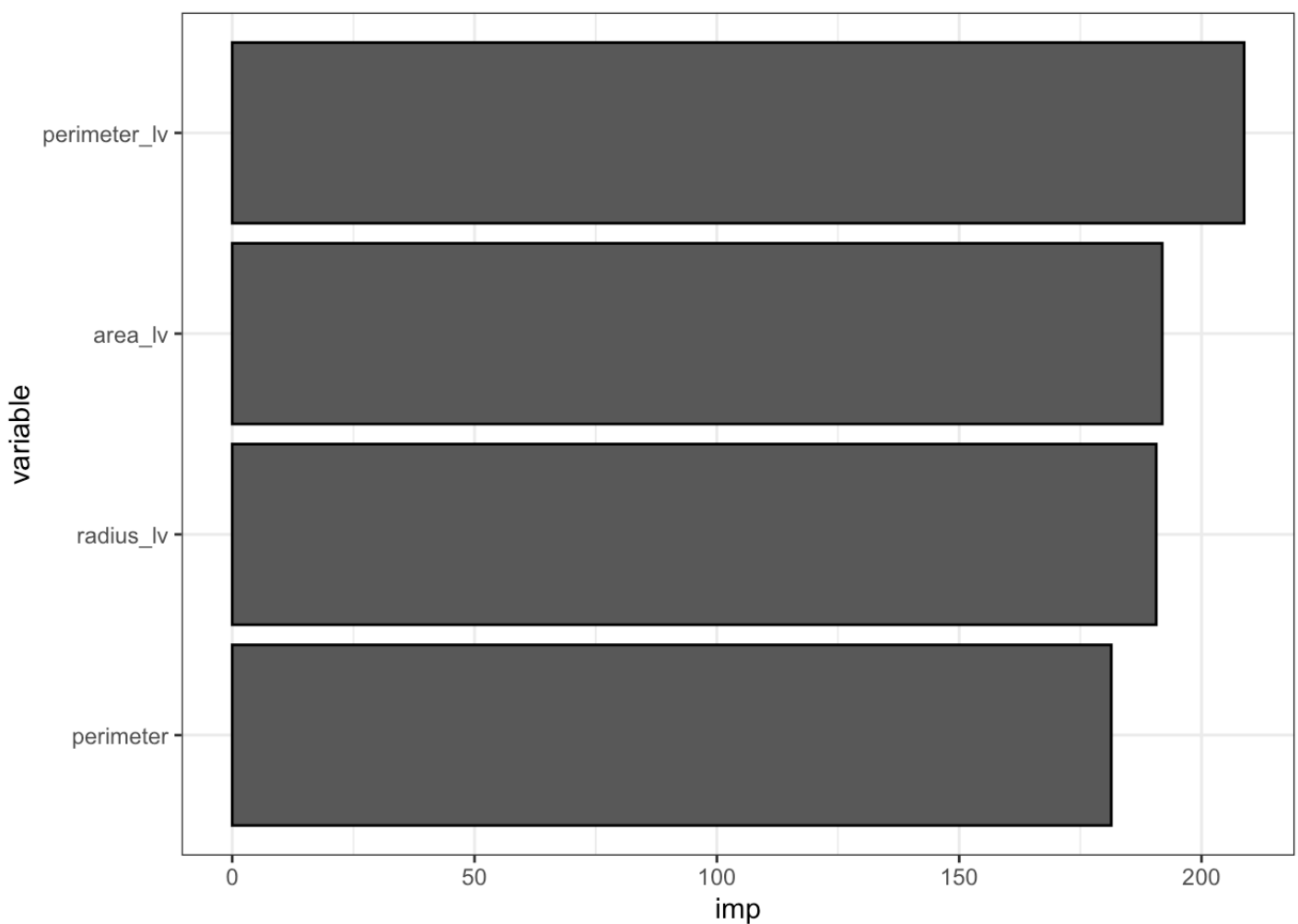
```
##      t_pred
##           0    1
## 0 280     6
## 1  19 149
```

```
print(acc)
```

```
## [1] 0.9449339
```

Bar graph imp vs variable

```
df=data.frame(imp=fit$variable.importance)
df2=df %>%
  tibble::rownames_to_column()%>%
  dplyr::rename("variable"= rowname)%>%
  dplyr::arrange(imp)%>%
  dplyr::mutate(variable = forcats::fct_inorder(variable))
ggplot2::ggplot(df2) +
  geom_col(aes(x=variable, y=imp), col="black", show.legend = F) +
  coord_flip() +
  scale_fill_grey() +
  theme_bw()
```



Confusion Matrix for test

```
t_pred=predict(fit, testx, type='class')
names(testy)=c("diagnosis")
```

Accuracy for Test

```
fit_test <- rpart(diagnosis~.,data=train_x_y_new, parms = list(split="information"
), method = 'class',control=rpart.control(minsplit=5,minbucket=3,cp=0.01))

t_pred=predict(fit_test, testx, type='class')
table(testy$diagnosis, t_pred)
```

```
##      t_pred
##           0  1
##    0 30   1
##    1  2 23
```

```
confusion_mat_test = table(testy$diagnosis, t_pred)
accTest = sum(diag(confusion_mat_test))/sum(confusion_mat_test)
print(confusion_mat_test)
```

```
##      t_pred
##           0  1
##    0 30   1
##    1  2 23
```

```
print(accTest)
```

```
## [1] 0.9464286
```