Attaching package: 'survival'

cluster

The following object is masked from 'package:caret':

```
In [1]:
   -----Loading of the Libraries-----Loading of the Libraries
library(pROC)
library(gbm)
library(randomForest)
library(caret)
library(readr)
library(rpart.plot)
library(caTools)
library(rpart)
library(plyr)
library(Hmisc)
Type 'citation("pROC")' for a citation.
Attaching package: 'pROC'
The following objects are masked from 'package:stats':
    cov, smooth, var
Loaded gbm 2.1.8
randomForest 4.6-14
Type rfNews() to see new features/changes/bug fixes.
Loading required package: lattice
Loading required package: ggplot2
Attaching package: 'ggplot2'
The following object is masked from 'package:randomForest':
   margin
Loading required package: rpart
Loading required package: survival
```

```
Loading required package: Formula
Attaching package: 'Hmisc'
The following objects are masked from 'package:plyr':
    is.discrete, summarize
The following objects are masked from 'package:base':
    format.pval, units
In [2]:
-----Loading of Dataset-----
credit card <- read.csv("C:\\Users\\13128\\Desktop\\creditcard.csv")</pre>
In [3]:
creditcard <- credit card</pre>
In [4]:
-----Checking any Missing values-----
apply(creditcard, 2, anyNA)
table(creditcard$Class)
Time: FALSE V1: FALSE V2: FALSE V3: FALSE V4: FALSE V5: FALSE V6: FALSE V7:
FALSE V8: FALSE V9: FALSE V10: FALSE V11: FALSE V12: FALSE V13: FALSE V14:
FALSE V15: FALSE V16: FALSE V17: FALSE V18: FALSE V19: FALSE V20: FALSE V21:
FALSE V22: FALSE V23: FALSE V24: FALSE V25: FALSE V26: FALSE V27: FALSE V28:
FALSE Amount: FALSE Class: FALSE
     0
            1
284315 492
In [5]:
-----Setting the Seed Value-----
set.seed(4495)
creditcard$Time <- NULL</pre>
creditcard[is.na(creditcard)] = -9999
```

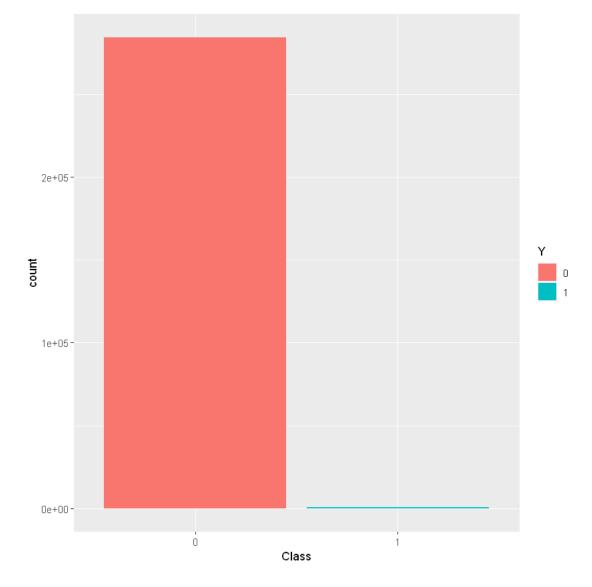
```
In [6]:
-----Replacing the NA values-----
replaceNAWithMean <- function(data) {</pre>
    for(i in 1:ncol(data)){
data[is.na(data[,i]), i] <- mean(data[,i], na.rm = TRUE)</pre>
}
replaceNAWithMean(creditcard)
In [7]:
-----Splitting of the Dataset-----
set.seed(4495)
t<-createDataPartition(p=0.5,y=creditcard$Class,list = F)
training<-creditcard[t,]</pre>
testing<-creditcard[-t,]
In [8]:
table(training$Class)
table(testing$Class)
    0
           1
142149
         255
    0
           1
142166
         237
In [9]:
-----Visualizations-----
library(ggplot2)
Y <- creditcard$Class
Y <- as.factor(Y)
ggplot(creditcard,aes(x = Y)) + geom_bar(aes(fill = Y)) + xlab('Class')
training$Class <- as.factor(training$Class)</pre>
ggplot(training,aes(x = training$Class)) + geom bar(aes(fill = training$Class)
) + xlab('Class')
(table(Y)[1]/length(Y))*100
(table(Y)[2]/length(Y))*100
```

"Use of `training\$Class` is discouraged. Use `Class` instead."

"Use of `training\$Class` is discouraged. Use `Class` instead."

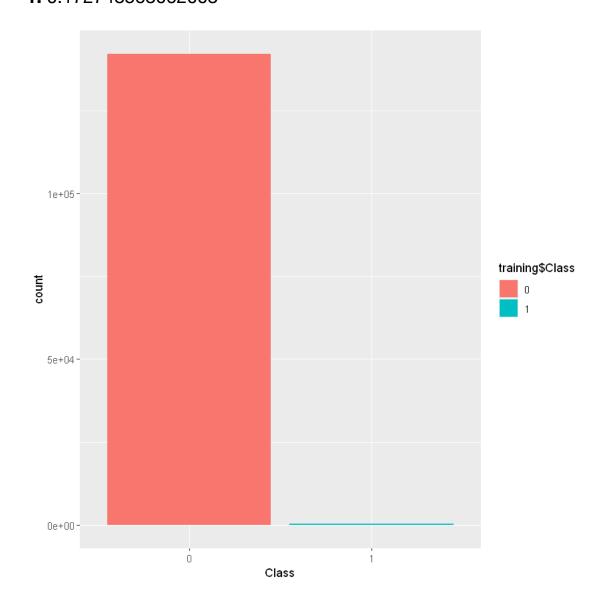
Warning message:

Warning message:



0: 99.827251436938

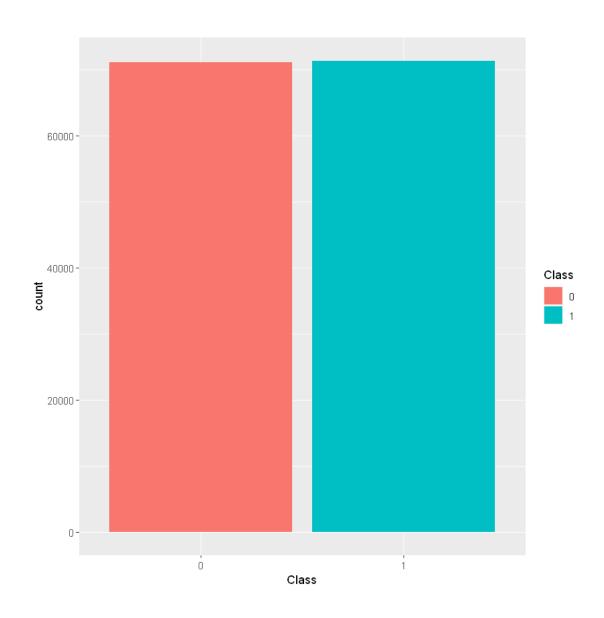
1: 0.172748563062003



In [10]:

```
library(ROSE)
attach(training)
set.seed(4495)
training_Rose <- ROSE(Class~.,data=training,seed = 4495)$data
training_Rose$Class <- as.factor(training_Rose$Class)
ggplot(training_Rose,aes(x = Class)) + geom_bar(aes(fill = Class))</pre>
```

Loaded ROSE 0.0-3



In [11]:

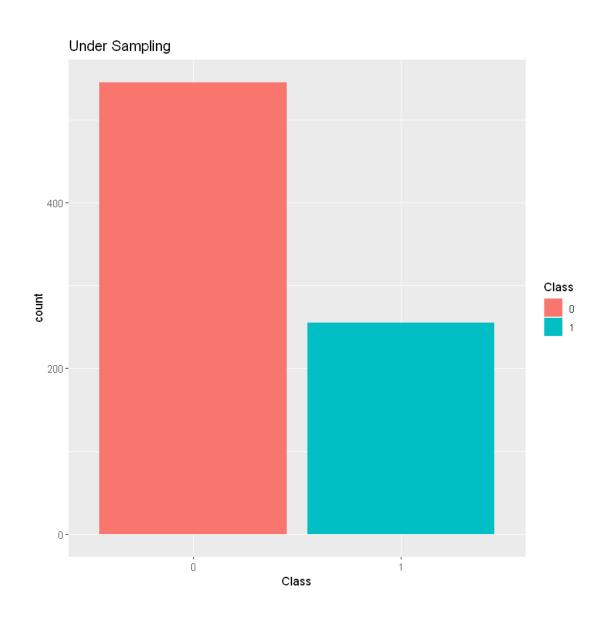
```
The following objects are masked from training (pos = 3):

Amount, Class, V1, V10, V11, V12, V13, V14, V15, V16, V17, V18,

V19, V2, V20, V21, V22, V23, V24, V25, V26, V27, V28, V3, V4,

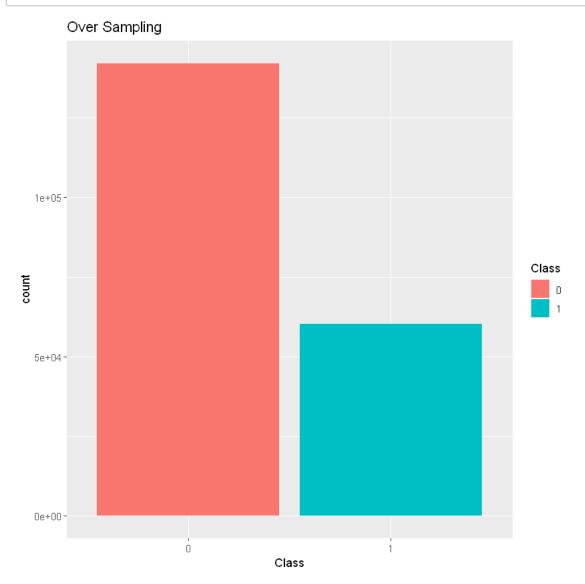
V5,

V6, V7, V8, V9
```



In [12]:

```
training_over <- ovun.sample(Class~.,data = training,method = "over",
N=202404,seed=4495)$data
ggplot(training_over,aes(x = Class)) + geom_bar(aes(fill = Class))+ggtitle("Over Sampling")</pre>
```



```
In [13]:
                 ------Logistic Regression Model-----
attach(training)
log <- glm(Class~., data = training,family=binomial)</pre>
The following objects are masked from training (pos = 3):
    Amount, Class, V1, V10, V11, V12, V13, V14, V15, V16, V17, V18
    V19, V2, V20, V21, V22, V23, V24, V25, V26, V27, V28, V3, V4,
V5,
    V6, V7, V8, V9
The following objects are masked from training (pos = 4):
    Amount, Class, V1, V10, V11, V12, V13, V14, V15, V16, V17, V18
    V19, V2, V20, V21, V22, V23, V24, V25, V26, V27, V28, V3, V4,
V5,
    V6, V7, V8, V9
Warning message:
"glm.fit: fitted probabilities numerically 0 or 1 occurred"
In [14]:
log2 <- glm(training Rose$Class~., data = training_Rose,family=binomial(logit)</pre>
Warning message:
"glm.fit: fitted probabilities numerically 0 or 1 occurred"
In [15]:
log3 <- glm(training under$Class~.,data = training under,family=binomial(logit)</pre>
))
Warning message:
"glm.fit: algorithm did not converge"
Warning message:
"glm.fit: fitted probabilities numerically 0 or 1 occurred"
In [16]:
log4 <- glm(training over$Class~.,data = training over,family=binomial(logit))</pre>
Warning message:
```

"glm.fit: fitted probabilities numerically 0 or 1 occurred"

```
In [17]:
----- Regression-----Prediction for Logistc Regression-----
pred <- predict(log4, testing,type="response")</pre>
pred <- round(pred)</pre>
In [18]:
-----Finding the Accuracy-----
accuracy <- (1-mean(pred != testing$Class))*100</pre>
accuracy
99.073053236238
In [19]:
 ----- Matrix of Logistic Regression-----
confusionMatrix(table(pred, testing$Class))
mat <- as.matrix(confusionMatrix(table(pred, testing$Class)))</pre>
Confusion Matrix and Statistics
pred 0
              1
  0 140879
              33
  1
    1287
             204
             Accuracy : 0.9907
               95% CI: (0.9902, 0.9912)
   No Information Rate: 0.9983
   P-Value [Acc > NIR] : 1
                Kappa : 0.2339
Mcnemar's Test P-Value : <2e-16
           Sensitivity : 0.9909
           Specificity: 0.8608
        Pos Pred Value: 0.9998
        Neg Pred Value: 0.1368
            Prevalence: 0.9983
        Detection Rate: 0.9893
  Detection Prevalence: 0.9895
     Balanced Accuracy: 0.9259
```

'Positive' Class: 0

In [20]:

```
------Getting AUROC values-----

print(roc(testing$Class,pred))

plot(roc(testing$Class,pred),main = "Logistic regression ROC curve(UnderSampling)")
```

Setting levels: control = 0, case = 1

Setting direction: controls < cases

Call:

roc.default(response = testing\$Class, predictor = pred)

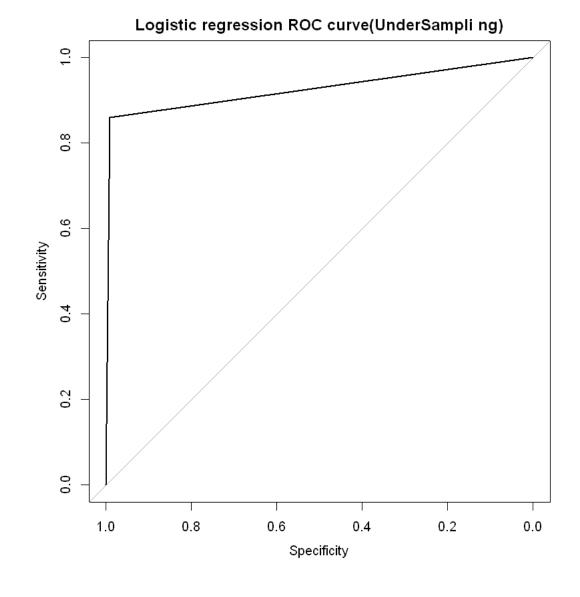
Data: pred in 142166 controls (testing\$Class 0) < 237 cases (testi

ng\$Class 1).

Area under the curve: 0.9259

Setting levels: control = 0, case = 1

Setting direction: controls < cases



In [21]:

In [22]:

logrgImp

A data.frame: 29 × 1

Overall

<dbl>

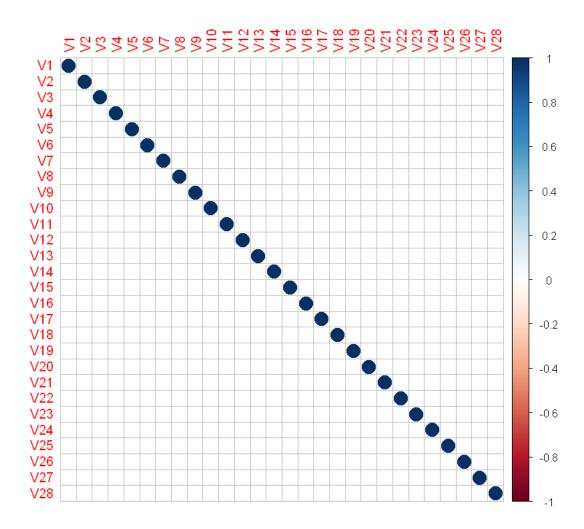
V1 2	0.7734055
-------------	-----------

- **V2** 9.1857692
- **V3** 16.7970361
- **V4** 55.7589186
- **V5** 20.1315767
- **V6** 12.3605734
- **V7** 12.1023691
- **V8** 18.5051715
- **V9** 15.2503371
- **V10** 30.2930195
- **V11** 24.5119709
- **V12** 34.2939813
- **V13** 30.5482485
- **V14** 41.9922865
- **V15** 3.3906752
- **V16** 24.8075535
- **V17** 17.7684252
- **V18** 3.8484204
- **V19** 4.5958506
- **V20** 21.6016873
- **V21** 10.1083491
- **V22** 35.3937924
- **V23** 2.0585542
- **V24** 0.5703025
- **V25** 3.9919142
- **V26** 29.9084167
- **V27** 9.4307256
- **V28** 6.1087944
- **Amount** 17.7063166

In [23]:

```
------Correlation Matrix------
library(corrplot)
correlations <- cor(creditcard[,1:28])
corrplot(correlations, method="circle")
```

corrplot 0.84 loaded



In [24]:

```
-----Random Forest Regression-----set.seed(4495)
library(e1071)
```

Attaching package: 'e1071'

```
The following object is masked from 'package:Hmisc':
impute
```

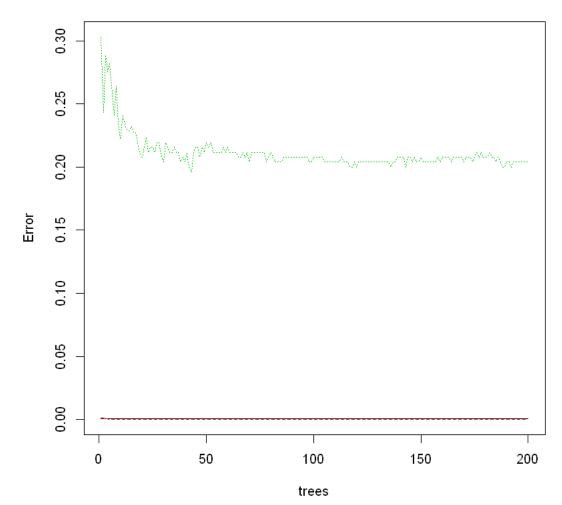
In [25]:

```
library(randomForest)
```

```
In [26]:
set.seed(4495)
system.time(rand model <- randomForest(Class~., data = training, ntree = 200))</pre>
         system elapsed
   user
 452.76
         14.23
                471.91
In [27]:
         -----Confusion Matrix-----
rand_model
Call:
 randomForest(formula = Class ~ ., data = training, ntree = 200)
               Type of random forest: classification
                     Number of trees: 200
No. of variables tried at each split: 5
        OOB estimate of error rate: 0.05%
Confusion matrix:
       0
           1 class.error
0 142133
         16 0.0001125579
      52 203 0.2039215686
In [28]:
```

```
plot(rand model)
```

rand_model



In [29]:

-----Variable Importance----importance(rand_model)

A matrix: 29×1 of type dbl

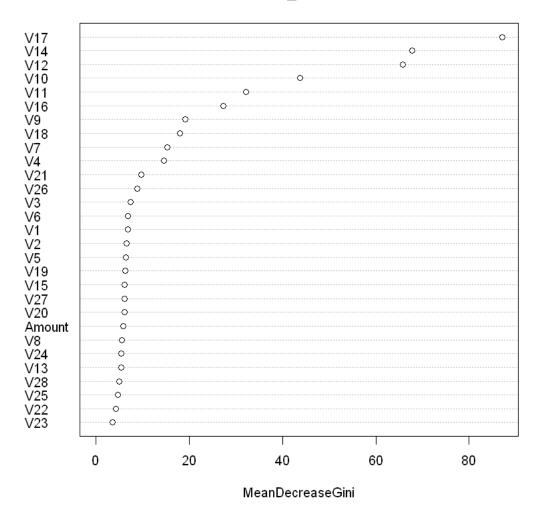
MeanDecreaseGini

	MeanDecreaseGini
V1	6.806659
V 2	6.614172
V 3	7.468481
V 4	14.529754
V 5	6.334535
V 6	6.859376
V 7	15.204911
V 8	5.526452
V 9	19.165079
V10	43.776896
V11	32.120139
V12	65.764325
V13	5.374850
V14	67.786611
V 15	6.183966
V 16	27.281721
V17	87.046434
V18	17.968638
V19	6.311618
V20	6.123274
V21	9.680153
V22	4.204290
V23	3.475204
V24	5.456588
V25	4.655896
V26	8.804864
V27	6.132533
V28	4.923242
Amount	5.818262

In [30]:

```
-----variable Importance Plot-----variable Importance Plot-----
```

rand_model



In [31]:

```
------
pred1 <- predict(rand_model,training,type = "class")
```

In [32]:

```
table(pred1,training$Class )
```

```
pred1 0 1
0 142149 0
1 0 255
```

In [33]:

```
pred2 <- predict(rand_model,testing,type = "class")</pre>
```

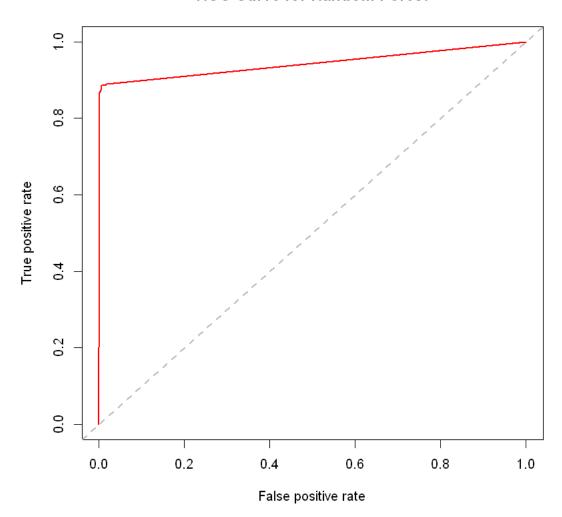
In [34]:

```
-------accuracy_rf <- mean(pred2 == testing$Class)
```

In [41]:

```
------
plot(pred3,main="ROC Curve for Random Forest",col=2,lwd=2)
abline(a=0,b=1,lwd=2,lty=2,col="gray")
```

ROC Curve for Random Forest



In [42]:

In [43]:

```
------
tree.model <- rpart(Class ~ ., data = training, method = "class", minbucket = 20)
```

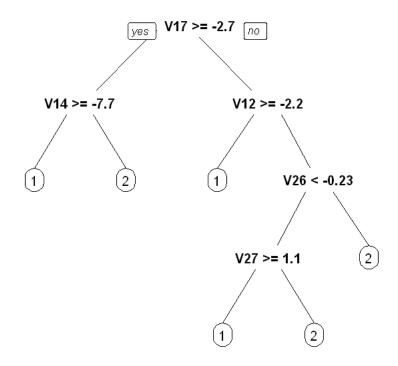
In [45]:

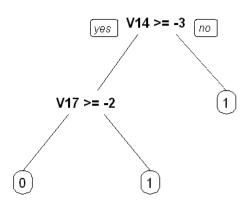
```
tree.model3 <- rpart(training_under$Class ~ ., data = training_under, method =
"class", minbucket = 20)</pre>
```

In [47]:

```
tree.model4 <- rpart(training_over$Class~.,data = training_over,method = "class",minbucket = 20)</pre>
```

```
In [48]:
------Generation of Decision Tree------
prp(tree.model)
prp(tree.model4)
```





```
In [49]:
-----Prediction and Accuracy------
tree.predict <- predict(tree.model4,testing,type = "class")</pre>
accuracy dt <-(1-mean(tree.predict != testing$Class))*100</pre>
In [50]:
accuracy dt
99.2212242719606
In [51]:
        ------Menerating Confusion Matrix-----Generating Confusion
```

```
confusionMatrix(table(tree.predict,testing$Class))
mat <- as.matrix(confusionMatrix(table(tree.predict,testing$Class)))</pre>
```

Confusion Matrix and Statistics

```
tree.predict
                         1
                       40
           0 141097
           1
               1069
                       197
               Accuracy : 0.9922
                 95% CI: (0.9917, 0.9927)
   No Information Rate: 0.9983
   P-Value [Acc > NIR] : 1
                  Kappa : 0.2601
Mcnemar's Test P-Value : <2e-16
```

Sensitivity: 0.9925 Specificity: 0.8312 Pos Pred Value: 0.9997 Neg Pred Value: 0.1556 Prevalence: 0.9983 Detection Rate: 0.9908 Detection Prevalence: 0.9911 Balanced Accuracy: 0.9119

'Positive' Class : 0

In [52]:

Setting levels: control = 0, case = 1

Setting direction: controls < cases

Call:

roc.default(response = testing\$Class, predictor = pred)

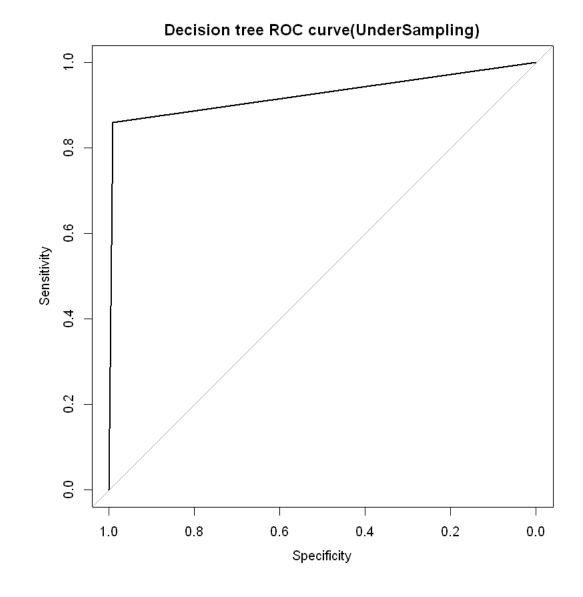
Data: pred in 142166 controls (testing\$Class 0) < 237 cases (testi

ng\$Class 1).

Area under the curve: 0.9259

Setting levels: control = 0, case = 1

Setting direction: controls < cases



In [53]:

```
------variable Importance-----variable Importance-----
```

Overall

	<dbl></dbl>
V 10	64393.638
V 11	55568.386
V12	56527.482
V 14	67253.315
V 17	60349.214
V20	1689.982
V 4	2176.235
V1	0.000
V 2	0.000
V 3	0.000
V 5	0.000
V 6	0.000
V 7	0.000
V 8	0.000
V 9	0.000
V13	0.000
V15	0.000
V 16	0.000
V 18	0.000
V 19	0.000
V21	0.000
V22	0.000
V23	0.000
V24	0.000
V25	0.000
V26	0.000
V27	0.000
V28	0.000
Amount	0.000