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Figure gbm - Gradient Boosting Machine (gbm) Learning technique
> fitModel2<-trainControl(method="cv", number=5, verbose=T)
> gbmfit<-train(classe~.,data=dataTrain, method="gbm", trControl=fitModel2, v</pre>
erbose=F)
Loading required package: gbm
Loading required package: survival
Attaching package: 'survival'
The following object is masked from 'package:caret':
    cluster
Loading required package: splines
Loading required package: parallel
Loaded gbm 2.1.1
Loading required package: plyr
+ Fold1: shrinkage=0.1, interaction.depth=1, n.minobsinnode=10, n.trees=150
- Fold1: shrinkage=0.1, interaction.depth=1, n.minobsinnode=10, n.trees=150
+ Fold1: shrinkage=0.1, interaction.depth=2, n.minobsinnode=10, n.trees=150
- Fold1: shrinkage=0.1, interaction.depth=2, n.minobsinnode=10, n.trees=150
+ Fold1: shrinkage=0.1, interaction.depth=3, n.minobsinnode=10, n.trees=150
- Fold1: shrinkage=0.1, interaction.depth=3, n.minobsinnode=10, n.trees=150
+ Fold2: shrinkage=0.1, interaction.depth=1, n.minobsinnode=10, n.trees=150
- Fold2: shrinkage=0.1, interaction.depth=1, n.minobsinnode=10, n.trees=150
+ Fold2: shrinkage=0.1, interaction.depth=2, n.minobsinnode=10, n.trees=150
- Fold2: shrinkage=0.1, interaction.depth=2, n.minobsinnode=10, n.trees=150
+ Fold2: shrinkage=0.1, interaction.depth=3, n.minobsinnode=10, n.trees=150
Fold2: shrinkage=0.1, interaction.depth=3, n.minobsinnode=10, n.trees=150
+ Fold3: shrinkage=0.1, interaction.depth=1, n.minobsinnode=10, n.trees=150
- Fold3: shrinkage=0.1, interaction.depth=1, n.minobsinnode=10, n.trees=150
+ Fold3: shrinkage=0.1, interaction.depth=2, n.minobsinnode=10, n.trees=150 - Fold3: shrinkage=0.1, interaction.depth=2, n.minobsinnode=10, n.trees=150
+ Fold3: shrinkage=0.1, interaction.depth=3, n.minobsinnode=10, n.trees=150
- Fold3: shrinkage=0.1, interaction.depth=3, n.minobsinnode=10, n.trees=150
+ Fold4: shrinkage=0.1, interaction.depth=1, n.minobsinnode=10, n.trees=150
- Fold4: shrinkage=0.1, interaction.depth=1, n.minobsinnode=10, n.trees=150
+ Fold4: shrinkage=0.1, interaction.depth=2, n.minobsinnode=10, n.trees=150
- Fold4: shrinkage=0.1, interaction.depth=2, n.minobsinnode=10, n.trees=150 + Fold4: shrinkage=0.1, interaction.depth=3, n.minobsinnode=10, n.trees=150 - Fold4: shrinkage=0.1, interaction.depth=3, n.minobsinnode=10, n.trees=150
+ Fold5: shrinkage=0.1, interaction.depth=1, n.minobsinnode=10, n.trees=150
- Fold5: shrinkage=0.1, interaction.depth=1, n.minobsinnode=10, n.trees=150
+ Fold5: shrinkage=0.1, interaction.depth=2, n.minobsinnode=10, n.trees=150
- Fold5: shrinkage=0.1, interaction.depth=2, n.minobsinnode=10, n.trees=150
+ Fold5: shrinkage=0.1, interaction.depth=3, n.minobsinnode=10, n.trees=150
- Fold5: shrinkage=0.1, interaction.depth=3, n.minobsinnode=10, n.trees=150
Aggregating results
Selecting tuning parameters
Fitting n.trees = 150, interaction.depth = 3, shrinkage = 0.1, n.minobsinnode
= 10 on full training set
> gbmfit$finalModel
A gradient boosted model with multinomial loss function.
150 iterations were performed.
There were 51 predictors of which 42 had non-zero influence.
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```
> class(gbmfit)
[1] "train"
                          "train.formula"
> predgbm<-predict(gbmfit, newdata=dataTest)</pre>
> confusionMatrix(predgbm, dataTest$classe)
Confusion Matrix and Statistics
```

Reference Prediction A 1651 52 0 0 0 14 1040 30 2 12 В C 8 40 977 35 10 D 3 919 13 1 16 F 0 4 8 1047 3

Overall Statistics

Accuracy: 0.9573 95% CI: (0.9519, 0.9624) No Information Rate: 0.2845 P-Value [Acc > NIR] : < 2.2e-16

Kappa: 0.946 Mcnemar's Test P-Value : NA

Statistics by Class:

	class: A	Class: B	Class: C	Class: D	Class: E
Sensitivity	0.9863	0.9131	0.9522	0.9533	0.9677
Specificity	0.9877	0.9878	0.9809	0.9933	0.9969
Pos Pred Value	0.9695	0.9472	0.9131	0.9653	0.9859
Neg Pred Value	0.9945	0.9793	0.9898	0.9909	0.9927
Prevalence	0.2845	0.1935	0.1743	0.1638	0.1839
Detection Rate	0.2805	0.1767	0.1660	0.1562	0.1779
Detection Prevalence	0.2894	0.1866	0.1818	0.1618	0.1805
Balanced Accuracy	0.9870	0.9504	0.9666	0.9733	0.9823

> predtrain<-predict(gbmfit, newdata=dataTrain)</pre> > confusionMatrix(predtrain, dataTrain\$classe)

Confusion Matrix and Statistics

Reference Prediction A B C D Ε A 3873 63 1 0 1 В 21 2547 57 15 6 C 9 41 2312 67 17 D 2 1 22 2168 19

F Overall Statistics

11 2473

Accuracy: 0.9735 95% CI: (0.9707, 0.9761) No Information Rate: 0.2843 P-Value [Acc > NIR]: < 2.2e-16

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Kappa: 0.9665 Mcnemar's Test P-Value: 1.539e-11

Statistics by Class:

	Class: A	Class: B	Class: C	Class: D	Class: E
Sensitivity	0.9916	0.9582	0.9649	0.9627	0.9794
Specificity	0.9934	0.9911	0.9882	0.9962	0.9980
Pos Pred Value	0.9835	0.9626	0.9452	0.9801	0.9912

Neg Pred Value	0.9966	0.9900	0.9926	0.9927	0.9954
Prevalence	0.2843	0.1935	0.1744	0.1639	0.1838
Detection Rate	0.2819	0.1854	0.1683	0.1578	0.1800
Detection Prevalence	0.2867	0.1926	0.1781	0.1610	0.1816
Balanced Accuracy	0.9925	0.9747	0.9766	0.9794	0.9887

> predtrain<-predict(gbmfit, newdata= dataTrain)
> confusionMatrix(predtrain, dataTrain\$classe)
Confusion Matrix and Statistics

Reference

Prediction	Α	В	C	D	Ε
Α	3873	63	1	0	1
В	21	2547	57	6	15
C	9	41	2312	67	17
D	2	1	22	2168	19
E	1	6	4	11	2473

Overall Statistics

Accuracy: 0.9735 95% CI: (0.9707, 0.9761) No Information Rate: 0.2843 P-Value [Acc > NIR]: < 2.2e-16

Kappa: 0.9665 Mcnemar's Test P-Value: 1.539e-11

Statistics by Class:

	Class: A	Class: B	class: c	Class: D	Class: E
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