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 Comparing a CART model to Random Forest (Part 1) (<https://www.analyticsvidhya.com/blog/2014/06/comparing-cart-random-forest-1/>)

# Comparing a CART model to Random Forest (Part 1)

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I created my first simple regression model with my father in 8th standard (year: 2002) or 8th grade. Obviously, my contribution in that model was minimal, but I really enjoyed the representation of the data. We tried validating all the assumptions etc. for this model. By

the exercise, we had 5 sheets of the simple regression model on 700 data points. The entire exercise was complex enough to confuse any person with average IQ level. When I look at my models which are built on millions of observations and utilize complex statistics behind the scenes, how machine learning with sophisticated tools (like SAS, SPSS, R) has made our life easy.

Having said that, many people in the industry do not bother about the complex statistics, 'behind the scene'. It becomes very important to realize the predictive power of each technique. No model is perfect in all scenarios. Hence, we need to understand the data and the surrounding system before coming up with a model recommendation.

In this article, we will compare two widely used techniques i.e. CART vs. Random forest. Random forest was covered in my last article (<https://www.analyticsvidhya.com/blog/2014/06/introduction-random-forest-simplified/>) take a case study to build a strong foundation of this concept and use R to do the computations. The dataset used in this article is an inbuilt dataset of R.

As the concept is pretty lengthy, we have broken down this article into two parts

### **Background on Dataset "Iris"**

Data set "iris" gives the measurements in centimeters of the variables : sepal length and petal length and width, respectively, for 50 flowers from each of 3 species of Iris. The data consists of 150 cases (rows) and 5 variables (columns) named Sepal.Length, Sepal.Width, Petal.Length, Petal.Width, Species. We intend to predict the Species based on the 4 flower characteristic variables.

We will first load the dataset into R and then look at some of the key statistics. Use the following codes to do so.

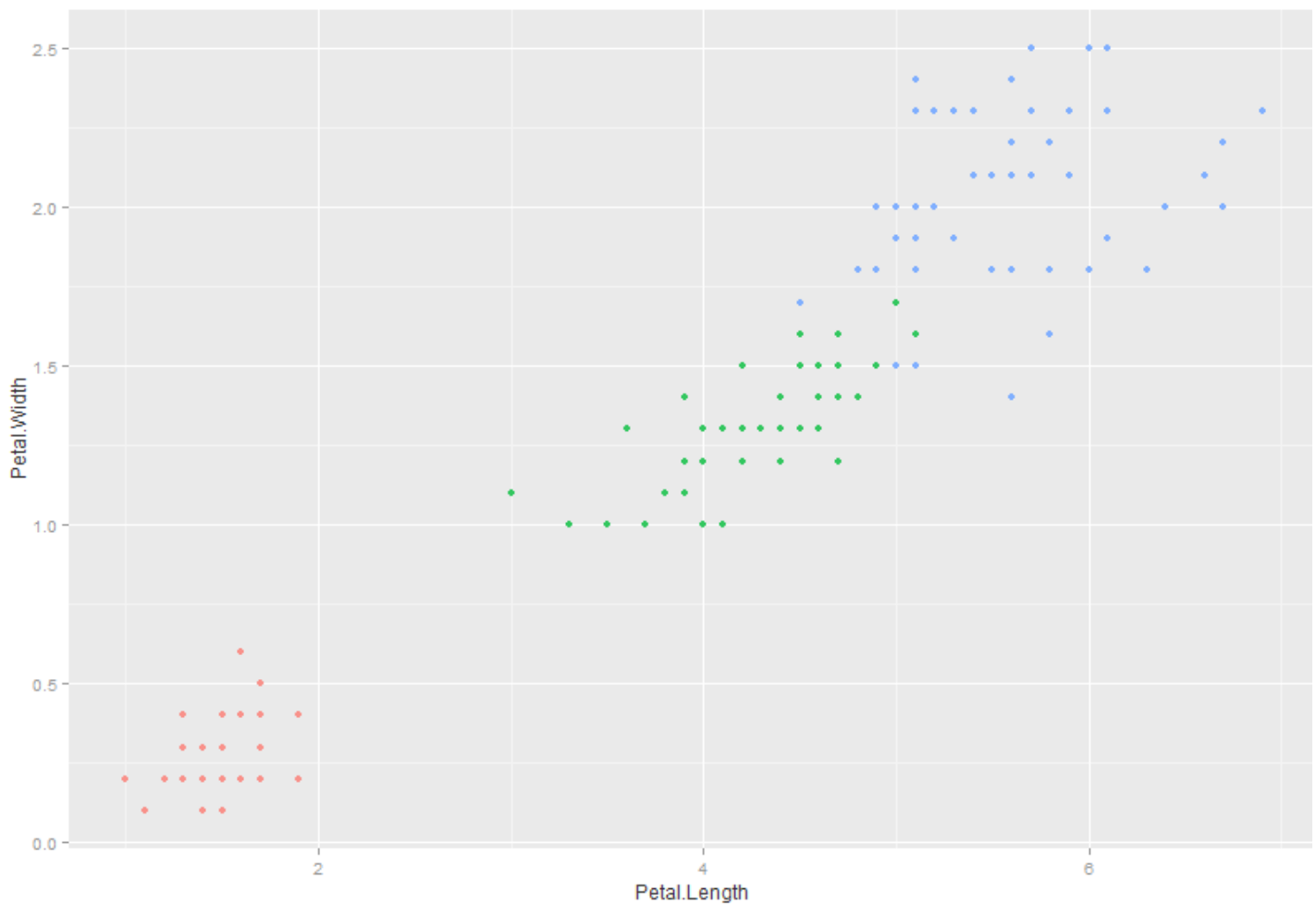
```
data(iris)
```

```
# look at the dataset
```

```
summary(iris)
```

```
# visually look at the dataset
```

```
qplot(Petal.Length,Petal.Width,colour=Species,data=iris)
```



(<https://www.analyticsvidhya.com/blog/wp-content/uploads/2014/06/plot1.png>)

The three species seem to be well segregated from each other. The accuracy in borderline cases determines the predictive power of the model. In this case, we will install packages for making a CART model.

```
library(rpart)
```

```
library(caret)
```

After loading the library, we will divide the population in two sets: Training and validation. to make sure that we do not overfit the model. In this case, we use a split of 50-50 for training and validation. Generally, we keep training heavier to make sure that we capture the key characteristics. You can use the following code to make this split.

```
train.flag <- createDataPartition(y=iris$Species,p=0.5,list=FALSE)
```

```
training <- iris[train.flag,]
```

```
Validation <- iris[-train.flag,]
```

## Building a CART model

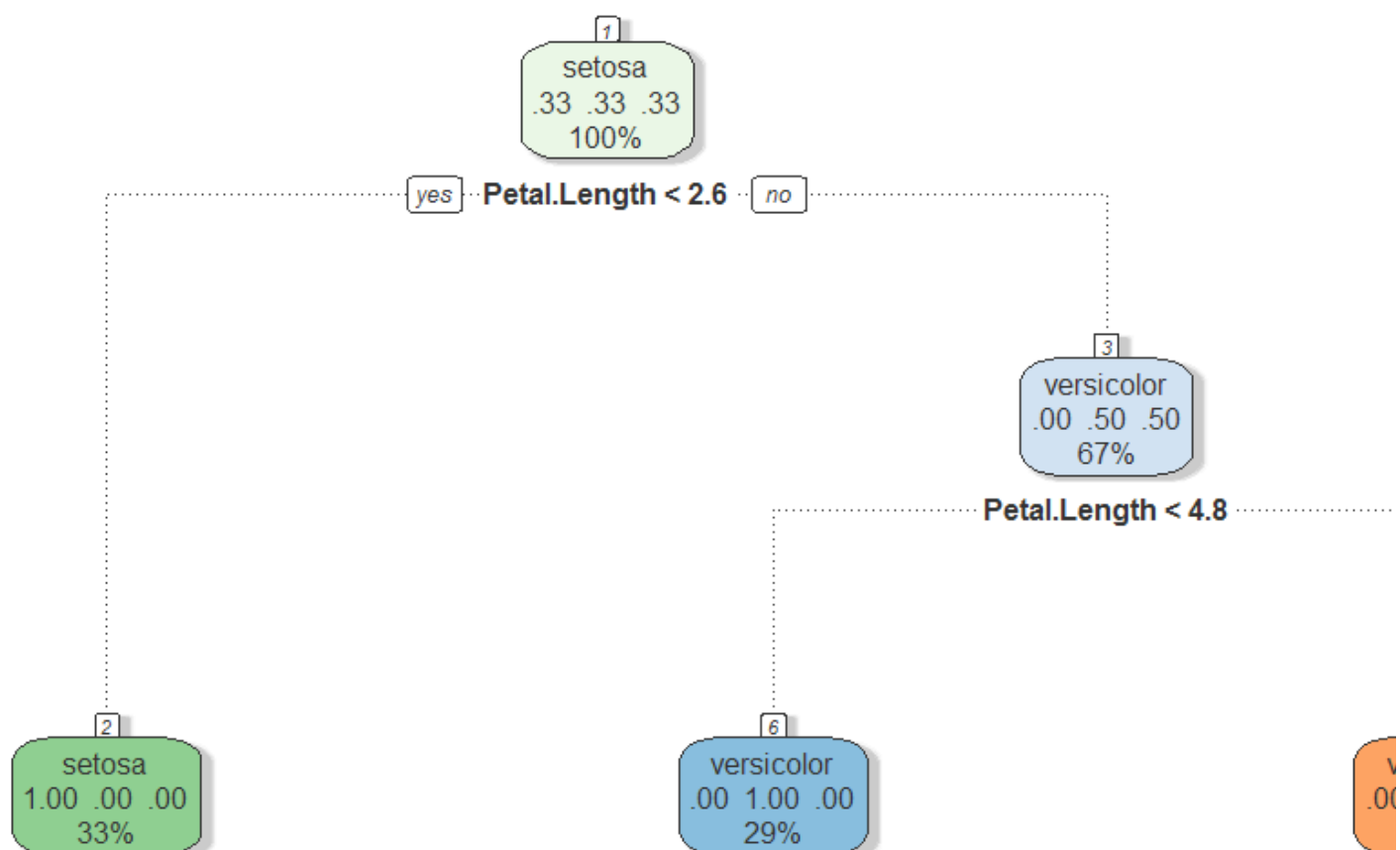
Once we have the two data sets and have got a basic understanding of data, we now build a model. We have used "caret" and "rpart" package to build this model. However, the representation of the CART model is not graphically appealing on R. Hence, we have used

called "rattle" to make this decision tree. "Rattle" builds a more fancy and clean trees, which are easily interpreted. Use the following code to build a tree and graphically check this tree:

```
modfit <- train(Species~.,method="rpart",data=training)
```

```
library(rattle)
```

```
fancyRpartPlot(modfit$finalModel)
```



(<https://www.analyticsvidhya.com/blog/wp-content/uploads/2014/06/tree.png>)

## Validating the model

Now, we need to check the predictive power of the CART model, we just built. Here, we are using a discordance rate (which is the number of misclassifications in the tree) as the decision metric. We will use the following code to do the same :

```
train.cart<-predict(modfit,newdata=training)
```

```
table(train.cart,training$Species)
```

```
train.cart   setosa versicolor virginica
```

```
setosa       25         0         0
```

```
versicolor   0         22         0
```

```
virginica     0         3        25
```

```
# Misclassification rate = 3/75
```

Only 3 misclassified observations out of 75, signifies good predictive power. In general, a misclassification rate less than 30% is considered to be a good model. But, the range of a good misclassification rate depends on the industry and the nature of the problem. Once we have built the model,

validate the same on a separate data set. This is done to make sure that we are not overfitting the model. In case we do overfit the model, validation will show a sharp decline in the predictive power. It is also recommended to do an out of time validation of the model. This will make sure that the model is not time dependent. For instance, a model built in festive time, might not hold in regular time. In simplicity, we will only do an in-time validation of the model. We use the following code to do in-time validation:

```
pred.cart<-predict(modfit,newdata=Validation)
```

```
table(pred.cart,Validation$Species)
```

```
pred.cart   setosa versicolor virginica
```

```
setosa      25         0         0
```

```
versicolor  0         22         1
```

```
virginica   0          3        24
```

```
# Misclassification rate = 4/75
```

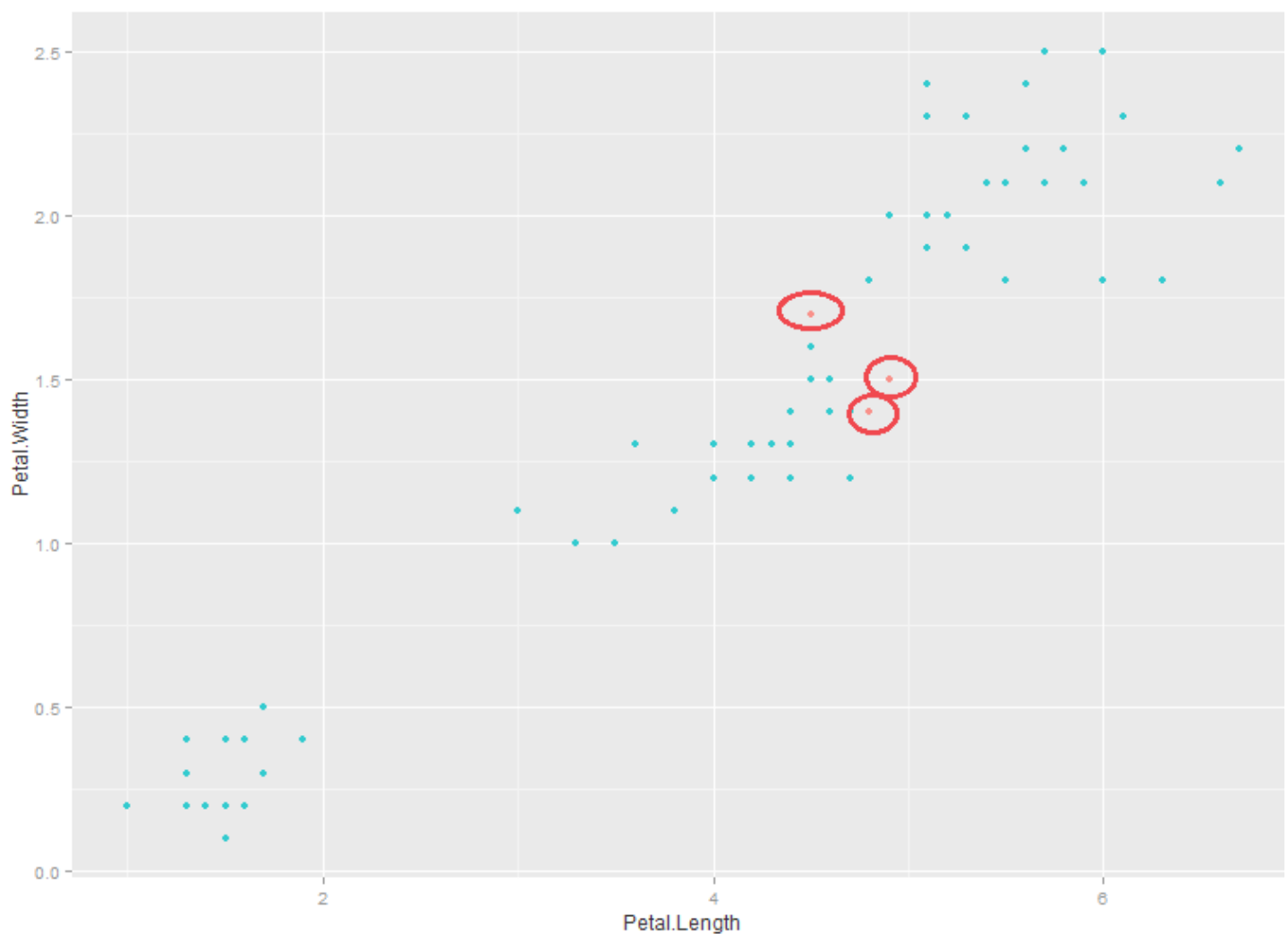
As we see from the above calculations that the predictive power decreased in validation as compared to training. This is generally true in most cases. The reason being, the model is trained on training data set, and just overlaid on validation training set. But, it hardly matters, if the predictive power is high in training and low in validation.

validation is lesser or better than training. What we need to check is that they are close enough. In this case, we do see the misclassification rate to be really close to each other. Hence, we select the CART model in this case study.

Let's now try to visualize the cases for which the prediction went wrong. Following is the code to find the same :

```
correct <- pred.cart == Validation$Species
```

```
qplot(Petal.Length, Petal.Width, colour=correct, data=Validation)
```



(<https://www.analyticsvidhya.com/blog/wp-content/uploads/2014/06/misclassify.png>)





As you see from the graph, the predictions which went wrong were actually those borderline cases. Before that these are the cases which make or break the confidence of the model. Most of the models will be able to categorize observation far away from the decision boundary. It takes a model to be sharp to distinguish these borderline cases.

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In random forest, the model will be able to make even better prediction for these borderline cases. It never generalizes the order of predictive power among a CART and a random forest or

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model on the data. We will see these observations graphically in the next article and take details on scenarios where random forest or CART comes out to be a better predictive model.

Did you find the article useful? Did this article solve any of your existing dilemmas? If you compared the two models in any of your projects? If you did, share with us your thoughts on

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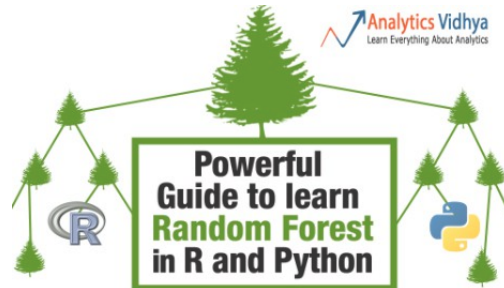
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I am Tavish Srivastava, a post graduate from IIT Madras in Mechanical Engineering. I have more than two years of work experience in Analytics. My experience ranges from hands on analytics in a developing country like India to convince banking partners with analytical solution in matured market like US. For last two and a half years I have contributed to various sales strategies, marketing strategies and Recruitment strategies in both Insurance and Banking industry.

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caret package, which has some other functions to split and train the data

A similar ex from caTools package in r -splitting the data according to binary c  
Dependent var has values as 0 and 1.

```
split=split(Dep var,SplitRatio=0.7)
train=subset(data,split==TRUE)
test=subset(data=split==FALSE)
```

---

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