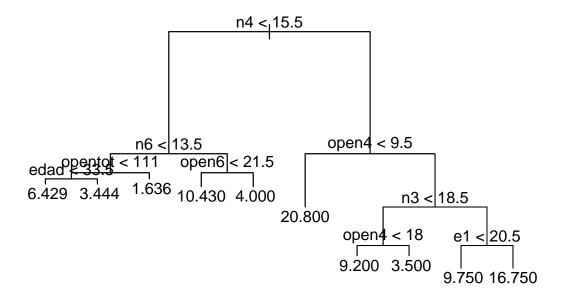
## Predicting depression based on big five personality scales

Load dataset and select training data:

```
library(foreign)
cardata <- read.spss("data Carillo et al.sav", to.data.frame = TRUE)
set.seed(42)
train <- sample(1:112, 80)</pre>
```

#### Regression tree for predicting depression

```
library(tree)
car.tree <- tree(bdi ~ ., data = cardata[train,])
plot(car.tree)
text(car.tree, pretty = 0)</pre>
```



```
car.pred <- predict(car.tree, cardata[-train,])
mean((cardata$bdi[-train] - car.pred)^2)

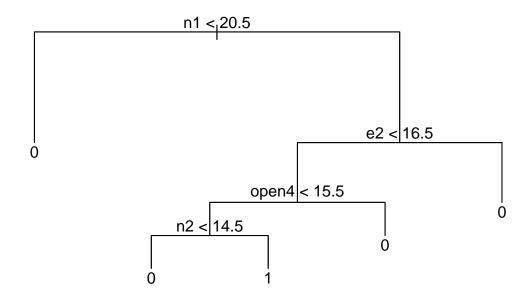
## [1] 87.56231

cor(cardata$bdi[-train], car.pred)

## [1] 0.4378805</pre>
```

### Classification tree for predicting depression

```
cardata2 <- cardata
cardata2$bdi <- factor(ifelse(cardata2$bdi > 16, 1, 0))
car.tree2 <- tree(bdi ~ ., data = cardata2[train,])
plot(car.tree2)
text(car.tree2, pretty = 0)</pre>
```



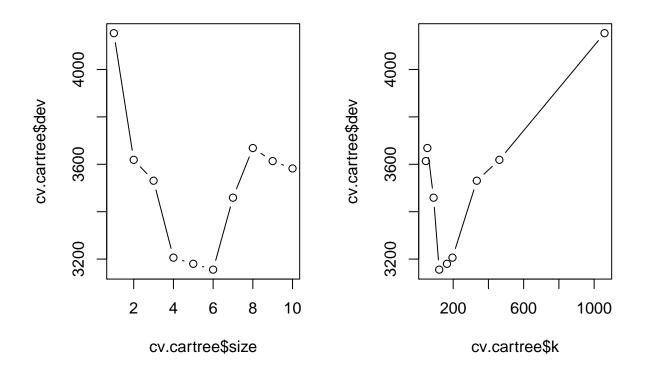
```
car.pred2 <- predict(car.tree2, cardata2[-train,])
prop.table(table(cardata2$bdi[-train], round(car.pred2[,2])))

##
## 0 1
## 0 0.75000 0.03125
## 1 0.09375 0.12500</pre>
```

### Pruning

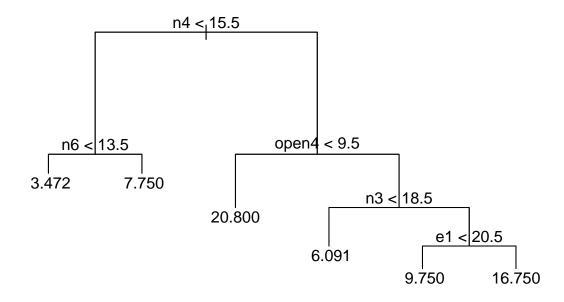
We are going to prune the regression tree. First, we determine the optimal tree size by 10-fold cross validation:

```
set.seed(3)
cv.cartree <- cv.tree(car.tree)
par(mfrow = c(1,2))
plot(cv.cartree$size, cv.cartree$dev, type = "b")
plot(cv.cartree$k, cv.cartree$dev, type= "b")</pre>
```



Optimal tree size is 6. Now we prune the tree:

```
cartree.pruned <- prune.tree(car.tree, best = 6)
plot(cartree.pruned)
text(cartree.pruned, pretty = 0)</pre>
```



```
car.pruned.pred <- predict(cartree.pruned, cardata[-train,])
mean((cardata$bdi[-train] - car.pruned.pred)^2)

## [1] 88.03171

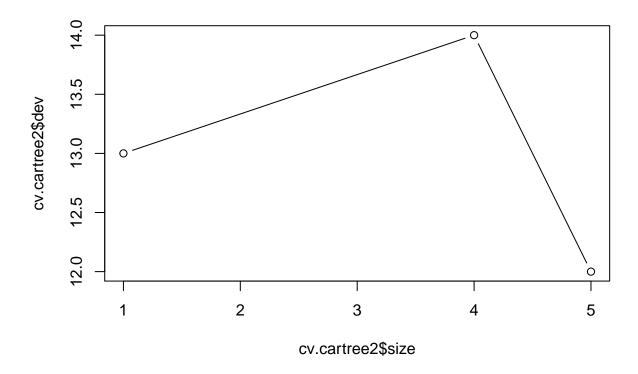
cor(cardata$bdi[-train], car.pruned.pred)

## [1] 0.4253286</pre>
```

We got a smaller tree, but in this case, pruning did not improve predictive accuracy on test data.

We also determine the optimal size for the classification tree:

```
cv.cartree2 <- cv.tree(car.tree2, FUN = prune.misclass)
plot(cv.cartree2$size, cv.cartree2$dev, type = "b")</pre>
```

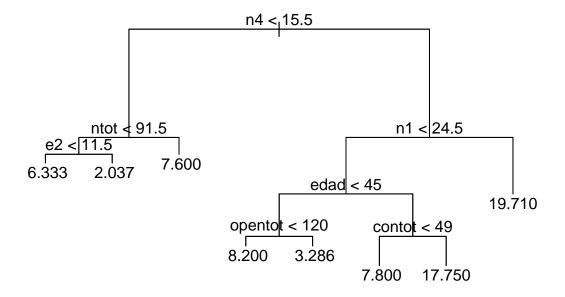


The classification tree does not need pruning.

## Tree instability

We grow a regression tree on a different sample of the same size, from the same data:

```
set.seed(46383)
train2 <- sample(1:112, 80)
car.tree3 <- tree(bdi ~ ., data = cardata[train2,])
plot(car.tree3)
text(car.tree3, pretty = 0)</pre>
```



```
car.pred3 <- predict(car.tree3, cardata[-train2,])
mean((cardata$bdi[-train2] - car.pred3)^2)

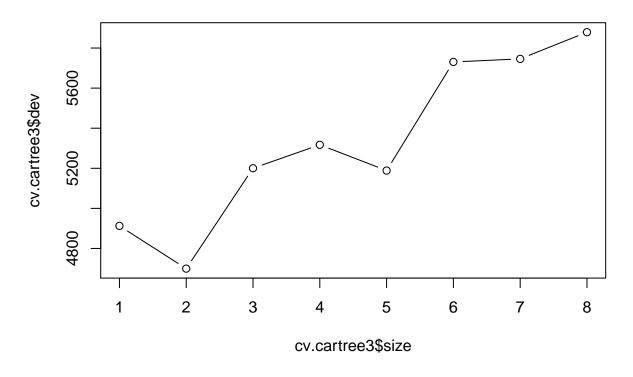
## [1] 69.03054

cor(cardata$bdi[-train2], car.pred3)

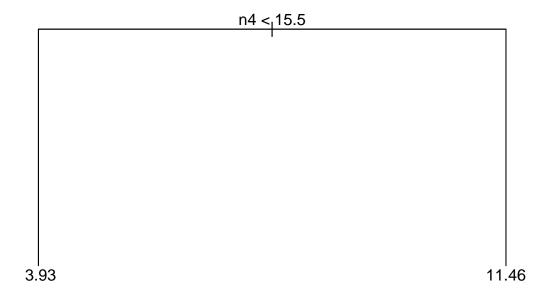
## [1] 0.5435759</pre>
```

In this sample, we find a different tree structure: only the first split in the tree is the same. Also, predictive accuracy in test data is different from that in the earlier sample, also indicating instability. Let's prune the tree to see if that yields a more stable tree:

```
set.seed(343545)
cv.cartree3 <- cv.tree(car.tree3)
plot(cv.cartree3$size, cv.cartree3$dev, type = "b")</pre>
```



```
cartree3.pruned <- prune.tree(car.tree3, best = 2)
plot(cartree3.pruned)
text(cartree3.pruned, pretty = 0)</pre>
```



```
car.pruned.pred3 <- predict(cartree3.pruned, cardata[-train2,])
mean((cardata$bdi[-train2] - car.pruned.pred3)^2)

## [1] 70.37825

cor(cardata$bdi[-train2], car.pruned.pred3)

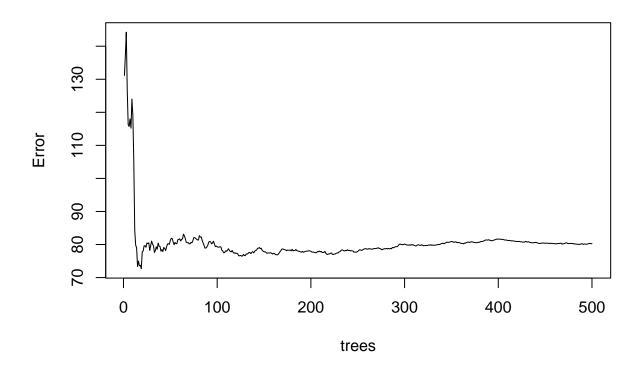
## [1] 0.4812077</pre>
```

Pruning does not completely eliminate the instability problem, although we do get a smaller tree and more similar predictive accuracy as with the full tree.

### Tree ensembles

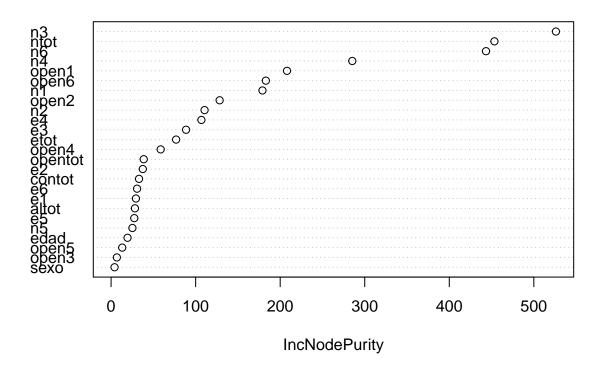
### Bagging

## **OOB** error estimates



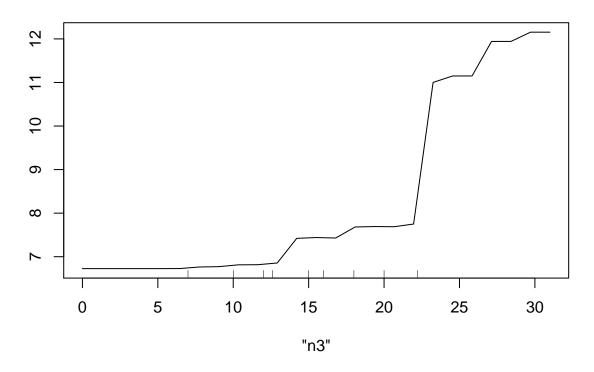
varImpPlot(bag.ens)

bag.ens



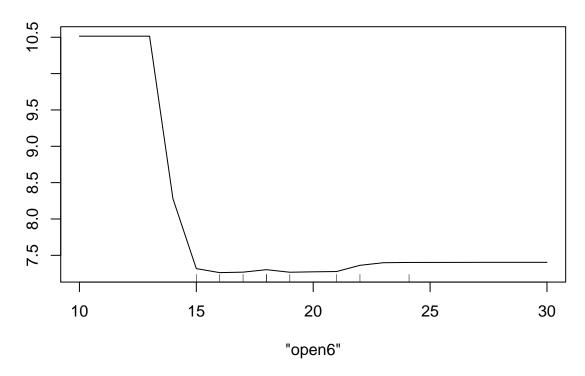
partialPlot(bag.ens, pred.data = cardata[train,], x.var = "n3")

# Partial Dependence on "n3"



partialPlot(bag.ens, pred.data = cardata[train,], x.var = "open6")

# Partial Dependence on "open6"

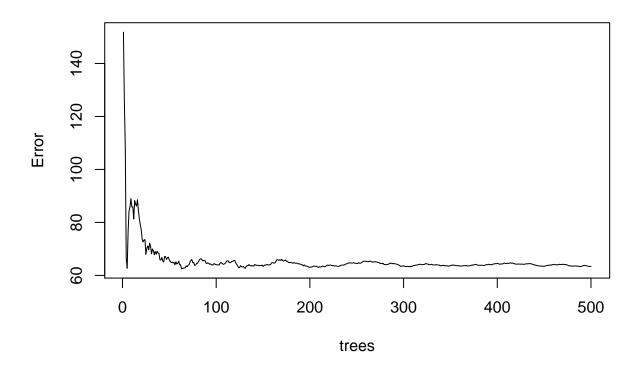


```
yhat.bag <- predict(bag.ens, newdata = cardata[-train,])
mean((yhat.bag - cardata$bdi[-train])^2)</pre>
```

## [1] 12.87469

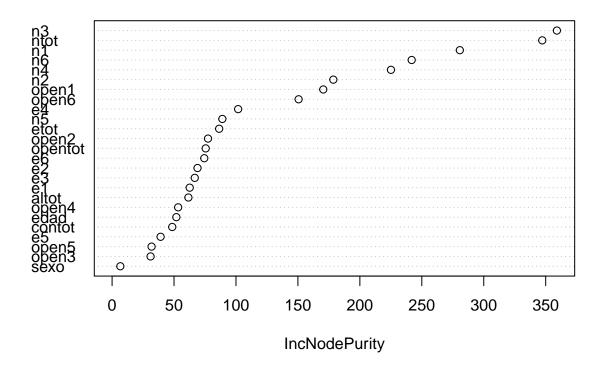
### Random forest

## **OOB** error estimates



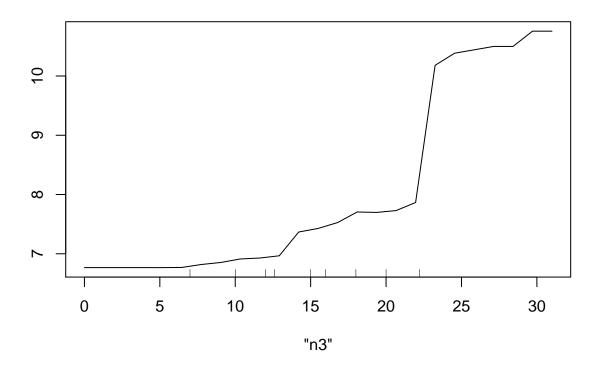
varImpPlot(rf.ens)

rf.ens



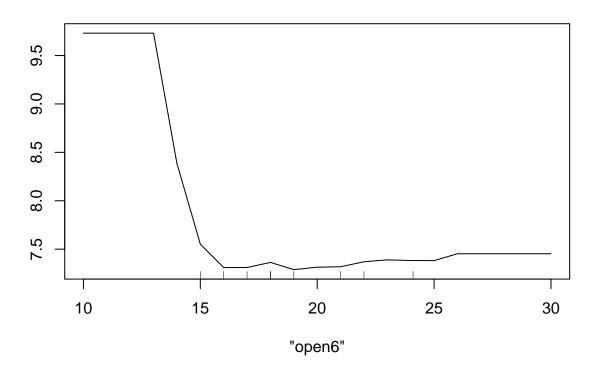
partialPlot(rf.ens, pred.data = cardata[train,], x.var = "n3")

# Partial Dependence on "n3"



partialPlot(rf.ens, pred.data = cardata[train,], x.var = "open6")

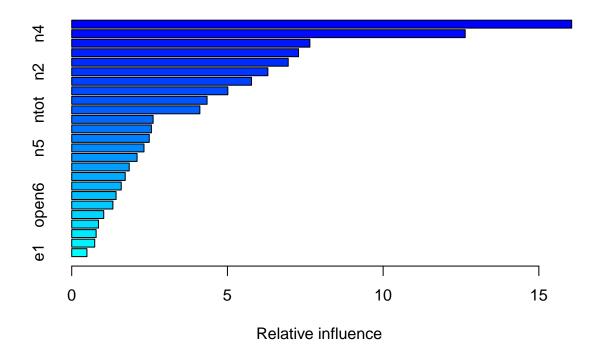
## Partial Dependence on "open6"



```
yhat.rf <- predict(rf.ens, newdata = cardata[-train,])
mean((yhat.rf - cardata$bdi[-train])^2)</pre>
```

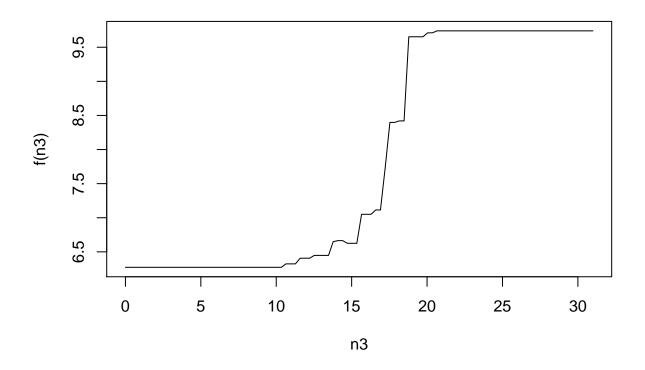
## [1] 12.50123

### Boosting

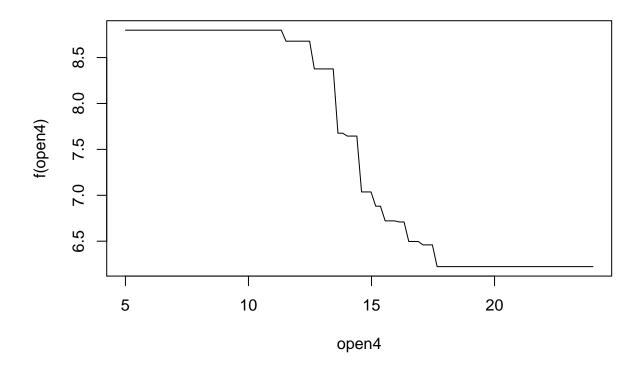


## rel.inf var ## n3 n3 16.0541454 ## n4 n4 12.6355765 ## open4 open4 7.6476624 ## n6 7.2831647 n6 ## open3 open3 6.9522351 ## n2 6.2984536 n2 ## e2 5.7735845 e2 ## n1 n1 5.0132382 ## e6 е6 4.3465808 ## ntot ntot 4.1149076 ## opentot opentot 2.6137953 ## etot etot 2.5589454 ## edad edad 2.4897261 ## n5 2.3205583 n5 ## open5 2.0987289 open5 ## open1 open1 1.8464558 ## altot altot 1.7182326 ## e5 e5 1.5898183 ## contot contot 1.4239046 ## open6 open6 1.3221283 ## open2 1.0278423 open2 ## sexo sexo 0.8591599 ## e4 e4 0.7841236 ## e3 еЗ 0.7386340 0.4883978 ## e1 e1

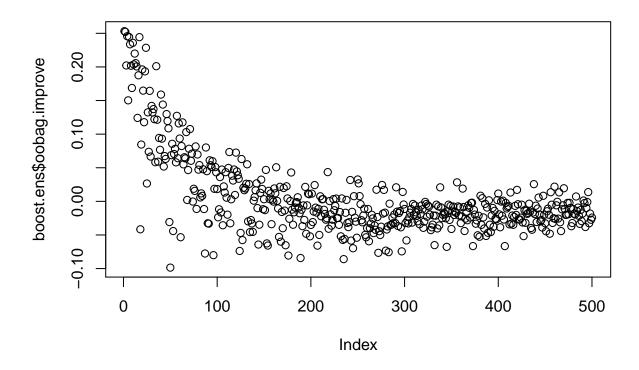
plot(boost.ens, i = "n3")



plot(boost.ens, i = "open4")



plot(boost.ens\$oobag.improve)



```
yhat.boost <- predict(boost.ens, newdata = cardata[-train,], n.trees = 500)
mean((yhat.boost - cardata$bdi[-train])^2)</pre>
```

#### ## [1] 65.23001

Note that I selected the parameter values above, after performing a cross validation using the train() function from package caret: