

Soil and climate model for predict admixture class of arabidopsis

Random forest based on all variables

Match admixture group for every sample base on the next simple rule, if component number N in vector > 0.7 , then this sample belong to the group N.

```
clear.just.parameters <- just.parameters

lbound <- .7
groups <- apply(all_data[, c(1:adm.count)], 1, function(current.row) {
  group <- ((current.row > lbound) * c(1:adm.count))[current.row > lbound]
  if (length(group) == 0) {
    return(NA)
  }
  return(group)
})
str(groups)
```

```
## int [1:1048] 6 6 6 NA NA NA 11 11 11 11 ...
```

```
clear.just.parameters <- just.parameters
prepared.data <- data.frame(group = groups)
prepared.data <- na.omit(cbind(prepared.data, clear.just.parameters))
prepared.data$group <- factor(prepared.data$group)
ncol(prepared.data)
```

```
## [1] 59
```

```
set.seed(666)
```

```
train.indexes <- sample(1:nrow(prepared.data), 0.8 * nrow(prepared.data))
train <- prepared.data[train.indexes, ]
test <- prepared.data[-train.indexes, ]
```

```
res.forest <- randomForest(group ~ ., data = train)
t_pred.forest <- predict(res.forest, test)
confMat <- table(test$group, t_pred.forest)
sum(diag(confMat)) / sum(confMat)
```

```
## [1] 0.8591549
```

```
important.dot.plot(res.forest)
```

