# Predicting PM10 concentration in Le Havre, France

## Describing the data set

Loading the data set:

```
library(VSURF)
A = VSURF::rep
str(A)
   'data.frame':
                    1096 obs. of
                                  18 variables:
##
                      13 22 29 25 23 17 20 17 32 31 ...
    $ PM10
               : int
##
    $ NO
                      15 25 31 36 72 53 31 28 47 44 ...
##
    $ NO2
                      25 32 42 36 56 54 45 41 47 50 ...
               : int
##
    $ SO2
               : int
                      5 3 10 4 15 NA 14 13 3 10 ...
##
    $ T.min
                      0.6 -0.6 -2 1.9 5.7 NA 6.3 7.4 7 7.3 ...
               : num
##
    $ T.max
                      8.4 7.6 1.2 6.5 8 NA 8.6 10.9 9.8 10.8 ...
               : num
                      5.025 3.333 -0.583 4.675 7.238 ...
##
    $ T.moy
               : num
    $ DV.maxvv : int
##
                      210 330 190 270 220 NA 190 190 250 220 ...
##
    $ DV.dom
               : num
                      338 45 180 225 NA ...
    $ VV.max
               : int
                      16 8 6 7 9 NA 11 22 14 8 ...
                      9.17 4.43 3.42 3.92 6.08 ...
##
    $ VV.moy
               : num
##
    $ PL.som
                      19 5 0 0 0 0 0 4 1 0 ...
               : int
##
    $ HR.min
               : int
                      79 81 74 86 94 NA 76 75 85 89 ...
                      99 93 90 99 97 NA 92 95 95 97 ...
##
    $ HR.max
               : int
##
    $ HR.moy
               : num
                      90.3 85.7 83 96.4 95.2 ...
##
    $ PA.moy
                      1010 1018 1024 1022 1021 ...
               : num
                      0.7 -0.5 2.4 2.7 2.4 0.6 -0.1 0 -0.1 1.2 ...
              : num
                      -0.3 0 0.1 -0.2 -0.2 -0.3 -0.5 NA -0.5 0.1 ...
    $ GTlehavre: num
```

According to the data set description, PM10 concentrations were measured in Le Havre, France, by Air Normand, with the associated weather data provided by Meteo France, from 2004 to 2006. The monitoring station is situated in an urban site, close to traffic, and considered the most polluted in the region. We have 1096 observations.

The data set description gives the following definitions for the 18 numeric variables in the data set:

- PM10: Daily mean concentration of PM10, in  $\mu g/m^3$ .
- NO, NO2, SO2: Daily mean concentration of NO, NO<sub>2</sub>, SO<sub>2</sub> in  $\mu g/m^3$ .
- T.min, T.max, T.moy: Daily minimum, maximum and mean temperature, in degree Celsius.
- DV.maxvv, DV.dom: Daily maximum speed and dominant wind direction in degree.
- VV.max, VV.moy: Daily maximum and mean wind speed, in m/s.
- PL.som: Daily rainfall in mm.
- HR.min, HR.max, HR.moy: Daily minimum, maximum and mean relative humidity, in %.
- PA.moy: Daily mean air pressure in hPa.

• GTrouen, GTlehavre: Daily temperature gradient in degree Celsius in the cities of Rouen and Le Havre, respectively.

PM10 is the target variable we want to predict using the other 17 variables.

We have missing data:

```
sum(is.na(A))
## [1] 269
There is data missing in most columns:
colnames(A)[ apply(A, 2, anyNA)]
    [1] "PM10"
                     "NO"
                                  "N02"
                                               "S02"
##
                                                            "T.min"
                                                                         "T.max"
    [7] "T.moy"
                     "DV.maxvv"
                                  "DV.dom"
                                               "VV.max"
                                                            "VV.moy"
                                                                         "HR.min"
                     "HR.moy"
## [13] "HR.max"
                                  "PA.moy"
                                               "GTlehavre"
I will substitute the missing values for the mean of each variable:
#Calculating the mean for each variable
mean_columns <- apply(A[,colnames(A)],2, mean, na.rm =</pre>
#Substituing the missing values by the mean
for (c in names(A)){
  A[,c] = ifelse(is.na(A[,c]), mean_columns[c], A[,c])
#Checking that there are no missing values
sum(is.na(A))
## [1] 0
```

# Methodology

We are interested in Regression. For variable selection, I will use Random Forests with the VSURF package. Linear, Decision Tree and Random Forest models will be constructed to make predictions. They will be compared by calculating the Root Mean Squared Error (RSME) and the Mean Absolute Error (MAE). The data set into Learning and Test sets before modelling.

# Splitting the data set

The models will be trained using the Learning set. Around 20% of the data will be kept separately on the Test set to test the models' predictions.

```
splitProb <- c(0.8,0.2)
splitNames <-c("Learning","Test")

n = nrow(x=A)

splitVector<- sample( x=splitNames, size=n, prob=splitProb, replace=TRUE )

table(splitVector)/n

## splitVector
## Learning Test
## 0.8029197 0.1970803</pre>
```

```
#getting the indeces
indeces<-list(
  learning = which(x=(splitVector=="Learning")),
  test=which(x=(splitVector=="Test"))
  )
#splitting the data set
learningSet<-A[indeces$learning,]
testSet<-A[indeces$test,]</pre>
```

#### Linear Model

## T.moy

## DV.dom

## VV.max

## VV.mov

## PL.som

## HR.min

## HR.max

## HR.moy

## PA.moy

## GTrouen

## GTlehavre

## DV.maxvv

Linear model using all explanatory variables:

-0.010066

-0.005504

-0.005122

-0.159936

0.303248

-0.243356

0.084768

0.010630

-0.062671

0.080378

-0.074847

1.004362

```
L = lm(PM10~., data = learningSet)
summary(L)
##
## Call:
## lm(formula = PM10 ~ ., data = learningSet)
## Residuals:
      Min
                10 Median
                                3Q
                                       Max
           -4.285 -0.875
##
  -17.380
                             3.303
                                    34.849
##
## Coefficients:
                Estimate Std. Error t value Pr(>|t|)
## (Intercept) -72.677640 31.750831 -2.289 0.02232 *
## NO
                0.119205
                           0.012941
                                       9.211 < 2e-16 ***
## NO2
                0.180255
                            0.021813
                                       8.264 5.30e-16 ***
## S02
                            0.018224
                                       9.138
                                             < 2e-16 ***
                0.166523
## T.min
                -0.244802
                            0.318731
                                     -0.768 0.44267
## T.max
                0.468910
                           0.355126
                                      1.320
                                             0.18705
```

-0.017

-2.029

-1.885

0.137807 -1.161 0.24613

2.655

-0.409

0.98668

0.04275 \*

0.05982 .

0.00807 \*\*

0.68299

3.164 0.00161 \*\*

1.623 0.10493

1.523 0.12814

0.136 0.89164

-0.609 0.54270

-4.537 6.52e-06 \*\*\*

```
##
## Residual standard error: 6.503 on 862 degrees of freedom
## Multiple R-squared: 0.5903, Adjusted R-squared: 0.5822
```

0.602870

0.002713

0.002718

0.186830

0.053640

0.055661

0.078008

0.102914

0.030269

0.183215

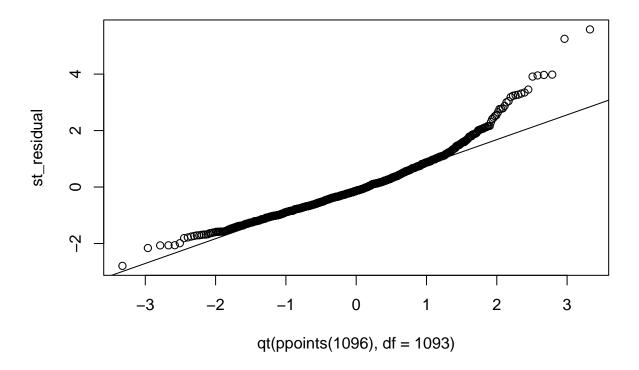
0.317393

## Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.05 '.' 0.1 ' ' 1

## F-statistic: 73.06 on 17 and 862 DF, p-value: < 2.2e-16

To check the gaussian assumption of the noise, I will use a QQ-plot of the studentized residuals:

```
#Studentized residuals - QQ plot
st_residual=rstudent(L)
#n=1096 #degrees of freedom n-3=1093
qqplot(qt(ppoints(1096), df=1093), st_residual)
qqline(st_residual, distribution=function(p){qt(p,df=1093)})
```



The QQ-plot is skewed and highly-tailed, so the noise does not appear to be Gaussian.

Using the Kolmogorov-Smirnov test to assess if the residuals have a student distribution.

```
#Smirnov test
#n=1096 #degrees of freedom n-3=1093
#'pt' is for student distribution
ks.test(st_residual, 'pt', 1093)

##
## Asymptotic one-sample Kolmogorov-Smirnov test
##
## data: st_residual
## D = 0.073788, p-value = 0.0001378
## alternative hypothesis: two-sided
```

The p-value is very small.

The noise does not seem to have a Gaussian distribution. This means that most parameters given in the linear model summary are not meaningful. We still can look at its  $R^2$  score, use the linear model for prediction and calculate errors, but we can't do much more.

#### Variable Selection

##

VSURF selected:

Many of the explanatory variables are highly dependent on each other.

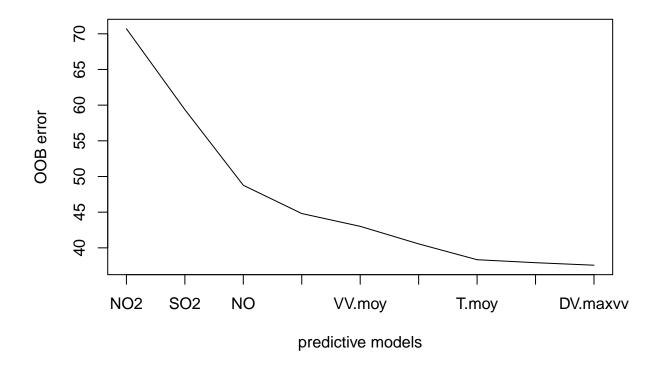
For example, the temperature variables:

```
cor(learningSet[,c("T.min","T.max","T.moy")])
##
             T.min
                       T.max
## T.min 1.0000000 0.9311867 0.9773227
## T.max 0.9311867 1.0000000 0.9829908
## T.moy 0.9773227 0.9829908 1.0000000
Or humidity variables:
cor(learningSet[,c("HR.min","HR.max","HR.moy")])
##
             HR.min
                       HR.max
                                  HR.moy
## HR.min 1.0000000 0.6042112 0.9102355
## HR.max 0.6042112 1.0000000 0.8157108
## HR.moy 0.9102355 0.8157108 1.0000000
And several others.
```

Since we cannot make the Gaussian assumption of the noise, we cannot use the p-values given by the linear model to perform variable selection. Instead, I will perform variable selection using Random Forest to detect the most relevant variables for prediction.

```
library('VSURF')
#Three steps variable selection procedure based on random forests for
#supervised classification and regression problems.
#First step ("thresholding step") is dedicated to eliminate irrelevant
#variables from the dataset. Second step ("interpretation step")
#aims to select all variables related to the response for interpretation purpose.
#Third step ("prediction step") refines the selection by eliminating redundancy in
#the set of variables selected by the second step, for prediction purpose.
set.seed(221921186)
Vy<-VSURF(PM10~.,data=learningSet, nmj=1)</pre>
## Thresholding step
## Estimated computational time (on one core): 52.4 sec.
## Interpretation step (on 17 variables)
## Estimated computational time (on one core): between 22.1 sec. and 115.6 sec.
##
## Prediction step (on 13 variables)
## Maximum estimated computational time (on one core): 68.9 sec.
## Warning in VSURF.formula(PM10 ~ ., data = learningSet, nmj = 1): VSURF with a formula-type call outp
## which are indices of the input matrix based on the formula:
## you may reorder these to get indices of the original data
summary(Vy)
##
   VSURF computation time: 2.9 mins
##
```

```
## 17 variables at thresholding step (in 1.3 mins)
## 13 variables at interpretation step (in 1.2 mins)
## 9 variables at prediction step (in 25.1 secs)
plot(Vy,step="pred",var.names=TRUE)
```



The selected explanatory variables are:

```
variables = c()
for (i in Vy$varselect.pred){
  variables <- c(variables,colnames(learningSet)[i+1])</pre>
}
variables
## [1] "NO2"
                    "S02"
                                 "NO"
                                              "GTlehavre" "VV.moy"
                                                                        "PL.som"
## [7] "T.moy"
                    "PA.moy"
                                 "DV.maxvv"
Removing other explanatory variables from Learning and Test sets:
learningSet= learningSet[,c(variables, "PM10")]
testSet= testSet[,c(variables,"PM10")]
```

#### Back to the Linear Model

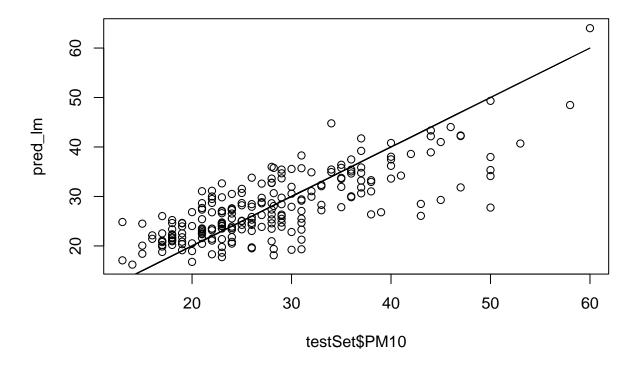
We can create a linear model using only the variables selected:

```
L2 = lm(PM10~., data = learningSet)
summary(L2)
```

```
##
## Call:
## lm(formula = PM10 ~ ., data = learningSet)
##
## Residuals:
##
               1Q Median
      Min
                                3Q
                                      Max
## -17.308 -4.415 -0.833
                            3.269
                                   34.647
##
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
## (Intercept) -72.791792 30.403155
                                     -2.394 0.01687 *
## NO2
                0.182061
                            0.021219
                                      8.580 < 2e-16 ***
## SO2
                0.177297
                            0.017455
                                     10.157
                                             < 2e-16 ***
                                      9.479 < 2e-16 ***
## NO
                0.111883
                            0.011803
## GTlehavre
                1.100519
                            0.243525
                                      4.519 7.07e-06 ***
## VV.moy
                0.064951
                            0.086556
                                      0.750 0.45323
## PL.som
                            0.051538
                                     -4.833 1.59e-06 ***
                -0.249072
## T.mov
                0.220000
                            0.045616
                                      4.823 1.67e-06 ***
## PA.moy
                0.083247
                            0.029544
                                      2.818 0.00495 **
## DV.maxvv
                -0.007919
                            0.002602
                                     -3.044 0.00241 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 6.519 on 870 degrees of freedom
## Multiple R-squared: 0.5844, Adjusted R-squared: 0.5801
## F-statistic: 135.9 on 9 and 870 DF, p-value: < 2.2e-16
```

The  $R^2$  didn't change much, but the model is simpler. We still have the problem that the noise is not gaussian, but we can do some predictions. Using the linear model to predict the PM10 values in the Test Set:

```
#predictions for Linear Model
pred_lm <- predict(L2, newdata = testSet)
#true values vs predictions
plot(testSet$PM10,pred_lm)
lines(testSet$PM10,testSet$PM10)</pre>
```



The Root Mean Squared Error (RMSE) of the predictions is:

```
RMSE_lm = sqrt(1 /nrow(testSet)*sum((testSet$PM10-pred_lm)**2))
RMSE_lm
## [1] 5.654754
The Mean Absolute Error (MAE) of the predictions is:
MAE_lm = 1 /nrow(testSet)*sum(abs(testSet$PM10-pred_lm))
MAE_lm
```

# ## [1] 4.348117

#### **Decision Tree**

Since the linear model might not be ideal in our case, we can try some non linear models.

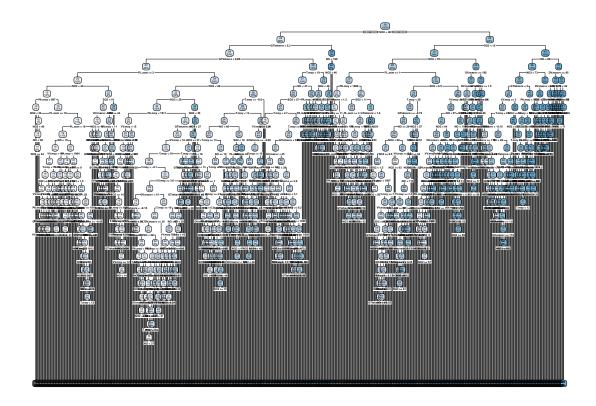
I will construct a decision tree using the explanatory variables selected before.

First, I obtain the maximal tree, maintaining the complexity parameter CP low.

The minimum number of observations in a node in order that an split is attempted is set as 2.

```
library(rpart)
library(rpart.plot)
#maximal tree
tree_max=rpart(PM10~.,data=learningSet, minsplit=2, cp = 10^(-9))
rpart.plot(tree_max)
```

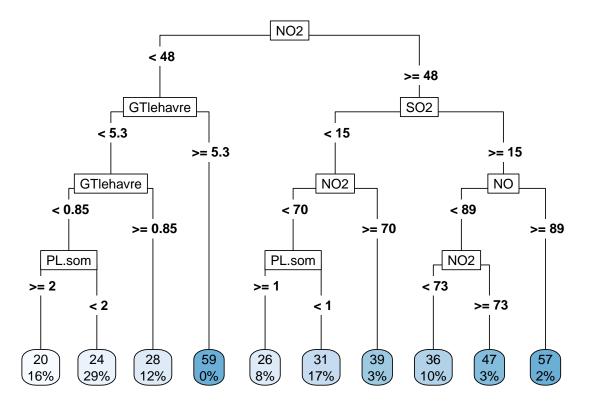
## Warning: labs do not fit even at cex 0.15, there may be some overplotting



Then, I will prune this tree using the 1-SE rule.

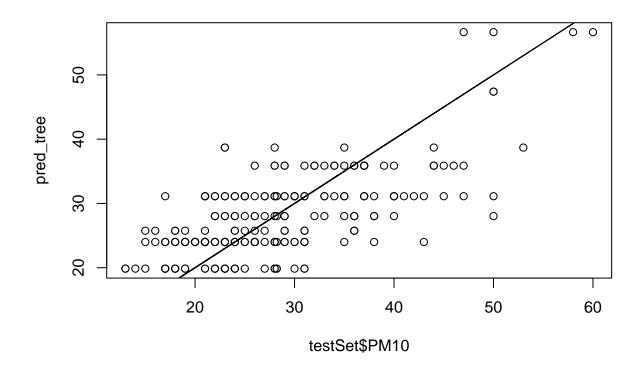
```
finalcart=function(T)
{
    P=printcp(T)
    CV=P[,4] #crossvalidation error
    a=which(CV==min(CV)) #finding the row with the smallest CV
    s=P[a,4]+P[a,5] #adding the standard deviation - the new threshold used in the 1SE rule
    ss=min(s) #in case s is a vector (several values are the min)
    b=which(CV<=ss)
    d=b[1] #selected value of cp
    Tf=prune(T,cp=P[d,1]) #pruning the maximal tree
    finalcart=Tf
}
tree = finalcart(tree_max)

rpart.plot(tree, type = 5)</pre>
```



Using the decision tree model to predict the PM10 values in the Test Set:

```
pred_tree<- predict(tree, newdata = testSet)
plot(testSet$PM10,pred_tree)
lines(testSet$PM10,testSet$PM10)</pre>
```



The RMSE error for this model:

```
RMSE_tree = sqrt(1 /nrow(testSet)*sum((testSet$PM10-pred_tree)**2))
RMSE_tree
## [1] 6.552845
The MAE error for this model:
MAE_tree = 1 /nrow(testSet)*sum(abs(testSet$PM10-pred_tree))
MAE_tree
```

## [1] 5.236443

## Random Forest

Finally, we can also use a Random Forest model.

```
library(randomForest)

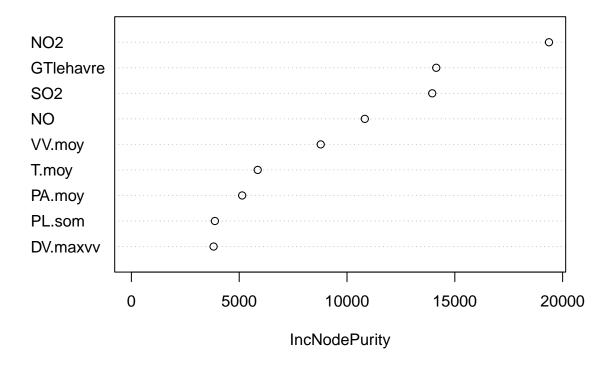
## randomForest 4.7-1.1

## Type rfNews() to see new features/changes/bug fixes.

RF=randomForest(PM10~.,data=learningSet)

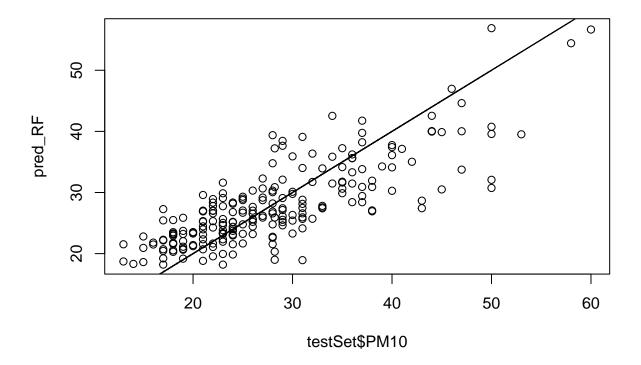
varImpPlot(RF)
```

## RF



Using the random forest model to predict the PM10 values in the Test Set:

```
pred_RF<- predict(RF, newdata = testSet)
plot(testSet$PM10,pred_RF)
lines(testSet$PM10,testSet$PM10)</pre>
```



The RMSE error for this model:

```
RMSE_RF = sqrt(1 /nrow(testSet)*sum((testSet$PM10-pred_RF)**2))
RMSE_RF

## [1] 5.430921
The MAE error for this model:
MAE_RF = 1 /nrow(testSet)*sum(abs(testSet$PM10-pred_RF))
MAE_RF
```

## [1] 4.260755

#### **Discussion and Conclusion**

```
results <- c(RMSE_lm, MAE_lm)
results <- rbind(results, c(RMSE_tree, MAE_tree))
results <- rbind(results,c(RMSE_RF, MAE_RF))
row.names(results) <- c("Linear Model", "Decision Tree", "Random Forest")
colnames(results) <- c("RMSE", "MAE")
results

## RMSE MAE
## Linear Model 5.654754 4.348117
## Decision Tree 6.552845 5.236443
## Random Forest 5.430921 4.260755</pre>
```

The Random Forest model has predictions with the smallest RSME and MAE, followed by the Linear Model. The Decision Tree has the worst performance, but it does provide some nice explainability.

The concentration of other pollutants such as NO,  $NO_2$  and  $SO_2$  in the atmosphere is one of the main predictors of the concentration of PM10. It is also case of the temperature gradient in Le Havre, where this particular monitoring station is located. When this temperature gradient is high, there seems to be a higher concentration of PM10.