# Case Study 3: Basic model fitting: Leaf burning

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# The data

An experiment to determine the effect on the burn time of leaves on the concentration of Nitrogen, Chlorine and Potassium. To build a purely scientific model from first principles based on chemistry and physics alone, would be extremely challenging. So instead, we propose various statistical models and fit them to the observations.

```
filepath <- "https://www.maths.nottingham.ac.uk/personal/pmzrdw/LeafData.txt"
# Download the data from the internet
download.file(filepath, destfile = "LeafData.txt", method = "curl")
LeafData <- read.table(file='LeafData.txt', header=TRUE)
LeafData[1:10,]</pre>
```

```
##
      Nitrogen Chlorine Potassium log_burn_time
## 1
          3.05
                    1.45
                               5.67
                                              0.34
## 2
          4.22
                    1.35
                               4.86
                                              0.11
## 3
          3.34
                    0.26
                               4.19
                                              0.38
          3.77
                    0.23
                               4.42
                                              0.68
## 5
          3.52
                    1.10
                               3.17
                                              0.18
## 6
          3.54
                    0.76
                               2.76
                                              0.00
## 7
          3.74
                    1.59
                               3.81
                                              0.08
## 8
          3.78
                    0.39
                               3.23
                                              0.11
## 9
          2.92
                    0.39
                               5.44
                                              1.53
## 10
          3.10
                    0.64
                                              0.77
                               6.16
```

```
str(LeafData) # look at the data structure
```

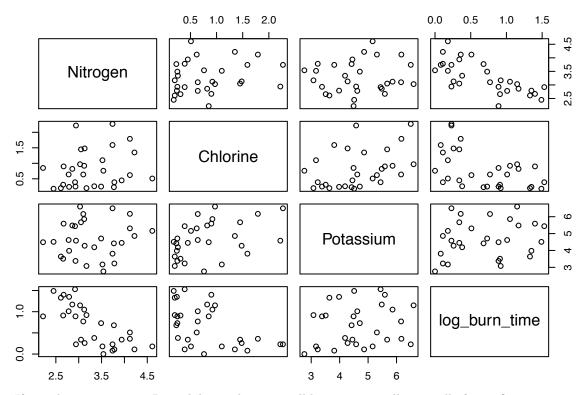
```
## 'data.frame': 30 obs. of 4 variables:
## $ Nitrogen : num 3.05 4.22 3.34 3.77 3.52 3.54 3.74 3.78 2.92 3.1 ...
## $ Chlorine : num 1.45 1.35 0.26 0.23 1.1 0.76 1.59 0.39 0.39 0.64 ...
## $ Potassium : num 5.67 4.86 4.19 4.42 3.17 2.76 3.81 3.23 5.44 6.16 ...
## $ log_burn_time: num 0.34 0.11 0.38 0.68 0.18 0 0.08 0.11 1.53 0.77 ...
```

The variables are

- Nitrogen %
- Chlorine %
- Potassium %
- log of leaf burn time

Lets start (as always), by visualising the data.

```
plot(LeafData)
```



If you choose not to use R markdown, then you will have to manually save all of your figures using a command such as

What questions might we want to ask?

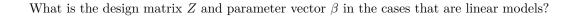
- How does the percentage of nitrogen, chlorine and potassium affect the leaf burn time?
- Which percentage is most important?

# Possible models

For the *i*th observation let  $y_i$  be the log of leaf burn time,  $x_{i,1}$  be the Nitrogen %,  $x_{i,2}$  be the Chlorine %, and  $x_{i,3}$  be the Potassium %. Let  $\beta = (a,b,c)^T$ .

Which of the following are linear models?

- $y_i = bx_{i,1} + \epsilon_i$ ,
- $y_i = a + bx_{i,1} + \epsilon_i$ ,
- $y_i = b^3 x_{i,1} + c x_{i,2} + \epsilon_i$ ,
- $y_i = a + b(x_{i,1} x_{i,2})^2 + \epsilon_i$ ,
- $y_i = a + 2bx_{i,1} + c\log(x_{i,2}) + \epsilon_i$ .
- $y_i = ax_{i,1} + bx_{i,1}^2 + \epsilon_i$ .



# Simple linear regression

Let y be the log leaf burn time and x be the Nitrogen %. We can write our simple linear regression model as

$$y_i = a + bx_i + \epsilon_i, \qquad i = 1, \dots, 30.$$

What are  $\beta$ ,  $g(x)^T$  and Z here?

To find the sum of squares estimates for this model using R, we can do the following:

```
fit1 <- lm(log_burn_time~Nitrogen, data = LeafData)
fit1

##
## Call:
## lm(formula = log_burn_time ~ Nitrogen, data = LeafData)
##
## Coefficients:</pre>
```

```
## (Intercept) Nitrogen
## 2.6257 -0.5916
```

## coef(fit1) # explicitly gives the fitted coefficients

```
## (Intercept) Nitrogen
## 2.6257040 -0.5916137
```

#### deviance(fit1)

## [1] 3.243512

Let  $D_1$  denote the deviance of this first order linear fit.

### **Model Choice**

We can fit a quadratic model.

$$y_i = a + bx_i + cx_i^2 + \epsilon_i$$
  $i = 1, \dots, n.$ 

The vector of parameters is  $\beta = (a, b, c)^T$ ,

 $z_i^T = \begin{bmatrix} 1 & x_i & x_i^2 \end{bmatrix}$  and the design matrix is

$$Z = \begin{bmatrix} 1 & x_1 & x_1^2 \\ 1 & x_2 & x_2^2 \\ \vdots & \vdots & \vdots \\ 1 & x_n & x_n^2 \end{bmatrix}$$

NB: This is a linear model since it is linear in the parameters.

```
fit2 <- lm(log_burn_time~Nitrogen+I(Nitrogen^2), data =LeafData)
fit2</pre>
```

```
##
## Call:
## lm(formula = log_burn_time ~ Nitrogen + I(Nitrogen^2), data = LeafData)
##
## Coefficients:
## (Intercept) Nitrogen I(Nitrogen^2)
## 4.6985 -1.8525 0.1861
```

#### deviance(fit2)

## [1] 3.103197

Let  $D_2$  denote the deviance of this quadratic linear model.

When  $D_1 - D_2$  is large then the quadratic model is much better than the simple linear regression, i.e. the straight line model is significantly improved by adding a quadratic term.

Note the use of

```
I(Nitrogen^2)
```

in the model formula. This is needed to seperate the two terms Nitrogen and Nitrogen^2.

If we had used

```
lm(formula = log_burn_time ~ Nitrogen + Nitrogen^2, data=LeafData)
```

then we would be fitting the linear model

$$y_i = a + b(x_i + x_i^2) + \epsilon_i$$
  $i = 1, ..., n$ .

instead (try it!).

We can also fit the null model

$$y_i = a + \epsilon_i$$
  $i = 1, \dots, n$ .

```
fit0 <- lm(log_burn_time~1, data=LeafData)
fit0</pre>
```

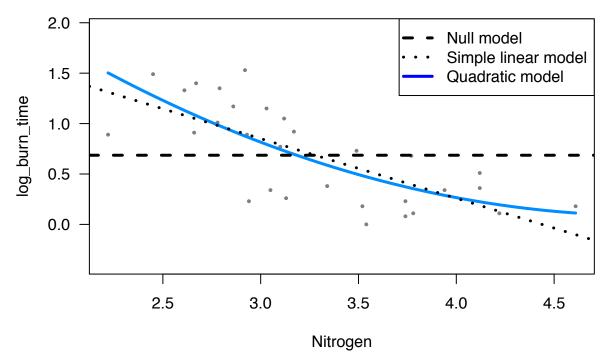
```
##
## Call:
## lm(formula = log_burn_time ~ 1, data = LeafData)
##
## Coefficients:
## (Intercept)
## 0.686
```

#### deviance(fit0)

```
## [1] 6.68952
```

We can plot these three models

```
library(visreg)
out = visreg(fit2, band=FALSE)
abline(fit1, lty=3, lwd=3)
abline(fit0, lty=2, lwd=3)
legend(x="topright", lty = c(2, 3, 1), c("Null model", "Simple linear model", "Quadratic model"), me
```



The plot and the deviances suggest that the linear model is probably sufficient. This is a subjective judgement based on eyeballing the data and fits - perhaps the most important aspect of model fitting. A more theoretical approach to model selection will be covered in Chapter 7.

# Simple linear model revisited, this time with matrices

Consider the simple linear model considered above. The design matrix is

$$Z = \begin{bmatrix} 1 & x_{1\,1} \\ 1 & x_{2\,1} \\ \vdots & \vdots \\ 1 & x_{30\,1} \end{bmatrix}.$$

We know that:

• The unbiased least squares estimator is

$$\hat{\beta} = (Z^T Z)^{-1} Z^T y = (2.63, -0.59)^T$$

• An unbiased estimator for  $\sigma^2$  is

$$s^{2} = \frac{1}{n-p} (y - Z\hat{\beta})^{T} (y - Z\hat{\beta}) = 0.12$$

• An estimate for  $Var(\hat{\beta})$  is given by

$$s^2 (Z^T Z)^{-1} = \begin{bmatrix} 0.130 & -0.04 \\ -0.04 & 0.012 \end{bmatrix}$$

These quantities are all available from R:

```
coef(fit1)
## (Intercept)
                   Nitrogen
     2.6257040 -0.5916137
fit1.sum <- summary(fit1)</pre>
fit1.sum$sigma^2
## [1] 0.1158397
vcov(fit1)
                (Intercept)
                               Nitrogen
                  0.1303385 -0.0385758
## (Intercept)
## Nitrogen
                 -0.0385758 0.0117657
The most useful R command for summarizing the model fit is
summary(fit1)
##
## Call:
## lm(formula = log_burn_time ~ Nitrogen, data = LeafData)
##
## Residuals:
        Min
                   1Q
                        Median
                                       ЗQ
                                                Max
## -0.65636 -0.27698 0.03712 0.27876 0.63181
##
## Coefficients:
                Estimate Std. Error t value Pr(>|t|)
##
## (Intercept)
                  2.6257
                              0.3610
                                      7.273 6.44e-08 ***
                 -0.5916
                              0.1085 -5.454 8.03e-06 ***
## Nitrogen
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.3404 on 28 degrees of freedom
## Multiple R-squared: 0.5151, Adjusted R-squared: 0.4978
## F-statistic: 29.75 on 1 and 28 DF, p-value: 8.025e-06
\mathbb{R}^2 and adjusted \mathbb{R}^2
Above we fitted the model
                                         y_i = a + bx_i + \epsilon_i
and found that s^2 = 0.12 and D = 3.24.
For the null model
                                           y_i = a + \epsilon_i
we find that s^2(\text{null}) = 0.23 and D_0 = 6.69.
```

#### Using R:

```
fit1.sum <- summary(fit1)
fit1.sum$r.squared ## gives the R^2 value</pre>
```

## [1] 0.5151353

```
fit1.sum$adj.r.squared ## gives the adjusted R^2 value
```

## [1] 0.4978187

Next, define a new input variate w to be 30 independent observations from a U(0,1) distribution and fit the model

$$y = a + bx + cw + \epsilon.$$

For this model we get that

$$R^2 = 0.516$$
 and  $R_{adj}^2 = 0.480$ .

Notice that  $R^2$  must improve when an input variate is added to the model. In this case, the input variate is unrelated to y and so  $R^2$  only improves by a very small amount. However  $R^2_{adj}$  has gone down: it reflects the fact that this new input variate is not providing any useful information and is not worth including in the model.

This shows why the adjusted R-squared is useful.

# Confidence intervals using R

```
confint(fit1)
                # gives 95% CI for both parameters
                   2.5 %
                             97.5 %
               1.886179 3.3652287
## (Intercept)
## Nitrogen
               -0.813804 -0.3694234
confint(fit1, level=0.99)
                            # gives 99% CI for both parameters
##
                    0.5 %
                              99.5 %
## (Intercept)
               1.6280992 3.6233088
               -0.8913442 -0.2918831
## Nitrogen
confint(fit1, level=0.99, parm="Nitrogen") ## supplies just the nitrogen CI.
##
                 0.5 %
                           99.5 %
## Nitrogen -0.8913442 -0.2918831
```