

Statistics in R

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goals

- introduce you to some basic statistics in R
- focus on linear models that you've met previously
- fitting simple linear models in R
- linear model validation techniques in R

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statistics in R

- many, many statistical tests available in R
- range from the simple to the highly complex
- many are included in standard installation
- you can extend the range of statistics by installing packages

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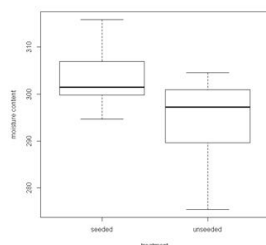
statistics in R

- example
 - does seeding clouds with dimethylsulphate alter the moisture content of clouds (can we make it rain!)
 - 10 random clouds were seeded and 10 random clouds unseeded
 - what's the null hypothesis?
 - no difference in mean moisture content between seeded and unseeded clouds

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statistics in R

- example
 - plot these data
 - interpretation?
 - what type of statistical test do you want to use?



```
str(clouds)
'data.frame': 20 obs. of 2 variables:
 $ moisture : num 301 302 299 316 307 ...
 $ treatment: Factor w/ 2 levels "seeded","unseeded":....
```

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statistics in R

- two sample t-test to compare the means of seeded group and unseeded group

```
t.test(clouds$moisture~clouds$treatment, var.equal=TRUE)
```

Two Sample t-test

```
data: moisture by treatment
t = 2.5404, df = 18, p-value = 0.02051
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 1.482679 15.657321
sample estimates:
mean in group seeded mean in group unseeded
      303.63           295.06
```

reject or fail to reject the null Hypothesis?

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statistics in R

- biological interpretation?
 - assumptions?
 - normality within each group
 - equal variance between groups
 - test for normality with Shapiro-wilk test for each group separately
- ```
shapiro.test(clouds$moisture[clouds$treatment=="seeded"])
shapiro.test(clouds$moisture[clouds$treatment=="unseeded"])
```

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## statistics in R

- null hypotheses?
- ```
Shapiro-Wilk normality test
data: moisture[treatment == "seeded"]
W = 0.9392, p-value = 0.544

shapiro.test(moisture[treatment=="unseeded"])

Shapiro-Wilk normality test
data: moisture[treatment == "unseeded"]
W = 0.8716, p-value = 0.1044
```
- I will give you a much better way of assessing normality later

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statistics in R

- equal variance using an F test
 - null hypothesis?
- ```
var.test(clouds$moisture~clouds$treatment)

F test to compare two variances

data: moisture by treatment
F = 0.5792, num df = 9, denom df = 9, p-value = 0.4283
alternative hypothesis: true ratio of variances is not equal to 1
95 percent confidence interval:
 0.1438623 2.3318107
sample estimates:
ratio of variances
 0.5791888
```
- I will give you a much better way of assessing equal variance later

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## linear models in R

- an alternative, but equivalent approach is to use a linear model to compare the means in each group
- general linear models are generally thought of as simple models, but can be used to model a wide variety of data and exp. designs
- traditionally statistics is performed (and taught) like using a recipe book (ANOVA, t-test, ANCOVA etc)
- general linear models provide a coherent and theoretically satisfying framework on which to conduct your analyses

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## what are linear models?

- thankfully, many of the statistical test you have learned so far are examples of linear models
- one sample t-test
  - two sample t-test
  - paired t-test
  - ANOVA
  - ANCOVA
  - correlation
  - linear regression
  - multiple regression
  - F-tests
  - etc...

} all cases of the (univariate)  
General Linear Model

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## model formulae

- general linear modelling is based around the concept of model formulae
 

response variable ~ explanatory variable(s) + error
  - literally read as 'variation in response variable modelled as a function of the explanatory variable(s) plus variation not explained by the explanatory variables'
  - it is the attributes of the response and explanatory variables that determines the type of linear model fitted
- $y \sim x$       if y and x are continuous then simple linear regression
- $y \sim \text{sex}$       If x is a categorical (nominal) variable then one-way ANOVA

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## linear modelling in R

- the function for carrying out linear regression in R is `lm`
- the response variable comes first, then the tilde `~` then the name of the explanatory variable

```
clouds.lm <- lm(moisture ~ treatment, data=clouds)
```

- how does R know that you want to perform a t-test (ANOVA)?

```
class(clouds$treatment)
[1] "factor"
```

- here the explanatory variable is a factor.

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## linear modelling in R

- to display the ANOVA table

```
anova(clouds.lm)
```

Analysis of Variance Table

Response: moisture

```
 Df Sum Sq Mean Sq F value Pr(>F)
treatment 1 367.22 367.22 6.4538 0.02051
Residuals 18 1024.20 56.90

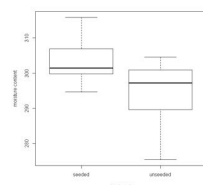
```

- do you notice anything about the  $p$  value?

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## linear modelling in R

- we have sufficient evidence to reject the null hypothesis (as before)
- therefore, there is a significant difference in the mean moisture content between clouds that were seeded and unseeded clouds
- do we accept this inference?
- what about assumptions?
- we could use Shapiro-wilks and  $F$  tests as before
- much better to assess visually by plotting the residuals



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## linear modelling in R

- because `clouds.lm` is a model object we can do stuff with it
- we can use the `plot()` function directly

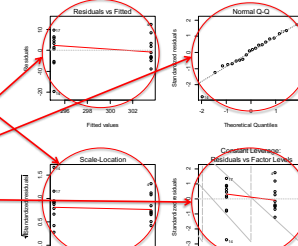
```
par(mfrow=c(2,2))
plot(clouds.lm)
```

assess equal variance assumption

assess normality assumption

assess any unusual observations

- look ok?



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## linear modelling in R

- the two sample t-test and a linear model with a categorical explanatory variable with 2 levels are equivalent
- this concept can easily be extended

| Traditional name     | Model formula                      | R code                                                               |
|----------------------|------------------------------------|----------------------------------------------------------------------|
| Bivariate regression | $Y \sim X_1$ (continuous)          | <code>lm(Y ~ X)</code>                                               |
| One way ANOVA        | $Y \sim X_1$ (categorical)         | <code>lm(Y ~ X)</code>                                               |
| Two-way ANOVA        | $Y \sim X_1$ (cat) + $X_2$ (cat)   | <code>lm(Y ~ X1 + X2)</code>                                         |
| ANCOVA               | $Y \sim X_1$ (cat) + $X_2$ (cont)  | <code>lm(Y ~ X1 + X2)</code>                                         |
| Multiple regression  | $Y \sim X_1$ (cont) + $X_2$ (cont) | <code>lm(Y ~ X1 + X2)</code>                                         |
| Factorial ANOVA      | $Y \sim X_1$ (cat) * $X_2$ (cat)   | <code>lm(Y ~ X1 * X2)</code> or <code>lm(Y ~ X1 + X2 + X1:X2)</code> |

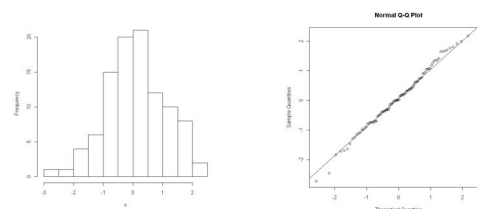
equivalent

difference?

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## extra detail of model validation in R

- normality of residuals
- use histograms and Q-Q plots of the residuals

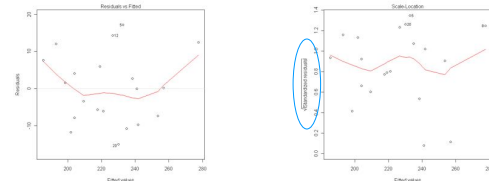


- normality is important, but linear models are fairly robust from smallish departures from normality

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## model validation in R

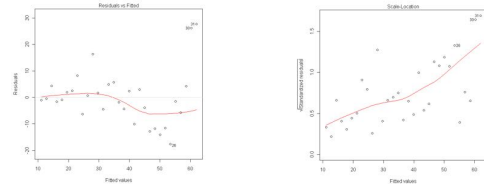
- equal variance (homogeneity, homoscedasticity)
  - plots of residuals (left) against fitted values and scale-location plots (right)
  - what you want to see is a complete scatter of points (sky at night)
  - what you don't want to see is any pattern or structure



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## model validation in R

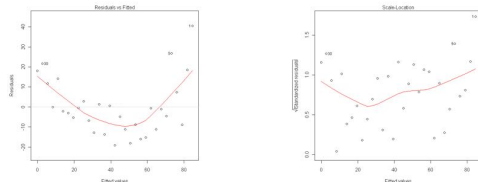
- a common problem is that the variance increases with the mean
- results in an expanding fan shaped pattern of residuals (left)
- an increase in scatter and value of standardised residuals (right)



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## model validation in R

- can also detect non linearity between response and explanatory variable(s) that has not been accounted for with the structure of the model
- residuals versus fitted clearer than scale-location plot



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