

# DATA HANDLING AND ASSUMPTIONS

Making the Most of Your Data



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## 1 Data Etiquettes

- Data Recording
- Data Storing
- Data Handling
- Data Mining
- Data Sharing

## 2 Statistical Assumptions

- Normality
- Independence
- Homogeneity of Variances

# Why Care?

*Biostatisticians often use 70% of their time to handle data and just 30% to actually analyse it.*

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- Proper data collection and data handling ensure accurate results
- Proper data collection cuts down on data handling time
- Proper data handling will make reproducing an analysis much easier

## What to consider?

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- What kind of data to record
- How data values are recorded/stored
- What kind of data values are feasible

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- When collecting categorical data, know what values the variables are allowed to take
- When collecting continuous data, know which range the variable values can fall into
- Make sure everyone involved in data collection is on the same page
- Make regular back-ups of your data set

## I recommend:

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  - Only allow pre-defined values to be entered
  - Need some excel macro writing
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# Common Issues

## The Decimals

Always use a *dot* to indicate decimals.

→ It is the standard in science.

## To NA Or Not To NA?

*Never enter NA values manually* into your data.

→ They cause problems in R.

## Redundancy Or Sparsity?

*Don't clutter data* with unnecessary data records.

→ Reduces storage space and chances for errors.



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# Data Storing

R works very well with:

- excel files (*.xls*, *.xlsx*, *.csv*)

- Easiest to handle outside of R, most storage-heavy

- Make sure to provide co-workers with a master file before data collection to avoid cell formatting issues on different computers

- text files (*.txt*)

- Difficult to handle outside of R, easy on storage

- I advise against using these, formatting issues are far too common

- RDS files (*.rds*)

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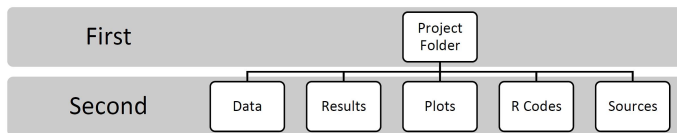
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I recommend a structure like the one below with at least two hierarchy levels.

The only files allowed in your first hierarchy level are:

- R master file
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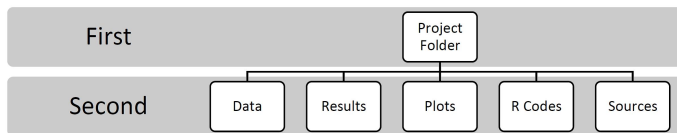
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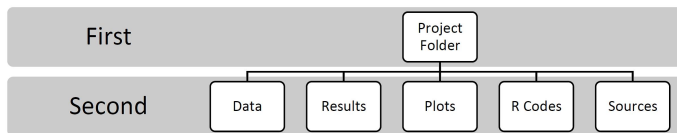
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Using the **README file**, one can identify what information is contained within the data set and thus decide:

- What type/class a data record should be of
- Which variables may be redundant
- Which data records exceed their variable-specific feasible thresholds
- Where to get comparative data sets from

**Data Mining** should then focus on:

- *Identifying problems* within the data records
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# Mining in R - Numbers or Visualisations?

For data mining, one may wish to enlist the use of Descriptive Statistics & Data Visualization:

## Descriptive Statistics:

- `summary()`
- `str()`
- iterative sub-setting and inspection

## Data Visualizations:

- Histograms (`hist()`)
- Scatter plots (`ggplot2` Package)

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**Documenting data recording** is just as important as proper data collection!

To do so, one usually uses a **README** file containing the following:

- Project Name and Summary
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- Your name and title (if you aren't the primary contact)
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Open science conduct is essential and you should (read *have to* as a student/employee of Aarhus University) share your data & coding to ensure **reproducibility** of your work:

## Peer-to-Peer:

- Raw data
- Code
- You may just as well point your peers to public repositories

## Public:

- Raw data
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- Html visualizations (`shiny`, `mapview`)
- Websites

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Statistical tests rely on individual *statistical assumptions*. Most prominent:

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- **Randomness:** Data are truly random
- **Independence:** Data are independent
- **Homogeneity of variances:** Data from separate groups have same variance
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## The QQ Plot In Theory

- Method for comparing two probability distributions by plotting their quantiles against each other
- If the two distributions being compared are similar, the plot will show the line  $y = x$ .
- Compare the data distribution to the normal distribution

# Independence

## Theory:

- Even the smallest dependence in your data can turn into heavily biased results (which may be undetectable).
- A dependence is a connection between/within the data.
- The assumption of independence relies on the absence of any connection in your data that haven't been accounted for in your approach (accounting for it is difficult).

## Independent data:

- *Between Groups*  
Groups of data records should be pulled from different individuals.
- *Within Groups*  
Data values within the same group are not to influence one another.
- *Within Individuals*  
Data values recorded for one individual should not influence each other. This is often an issue with repeated measurement approaches.

→ Fixing this *after data collection* is **almost impossible!**

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Data values within the same group are not to influence one another.
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Data values recorded for one individual should not influence each other. This is often an issue with repeated measurement approaches.

→ Fixing this *after data collection* is **almost impossible!**

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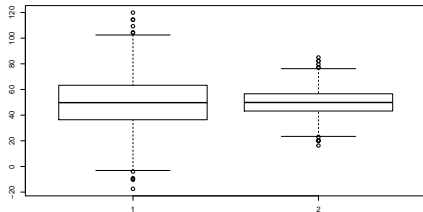
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# Homogeneity of Variances

Particularly important for t-Tests and ANOVAs

- **Assumption:** Data from separate groups have same variance
- **Test:** `leveneTest()` in the `car` package.

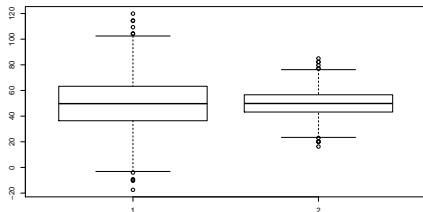


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## Levene's Test for Homogeneity of Variance (center = median)
##           Df F value Pr(>F)
## group      1      337 <2e-16 ***
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