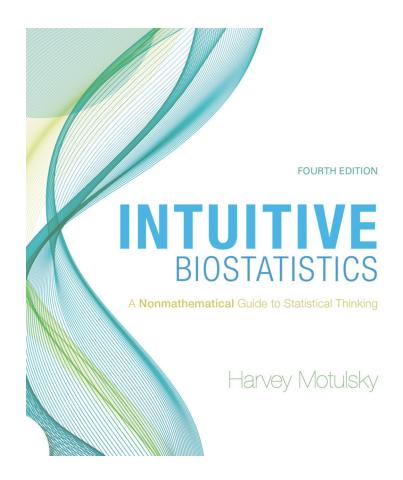
BIOSTATISTICS FAQ



find answers to those questions you dare not ask

biostatistics: find answers to those questions you dare not ask

This FAQ can help you find concrete tips in the excellent book "Intuitive biostatistics".

You can borrow it from the secretariat or the CMU library.

How do I know how many samples I need?

The appropriate tool is power analysis.

How should I deal with outliers?

precise rules and methods.

Read chapters 20 & 26

Read chapter 25

Can I run statistics on technical replicates?

Statistics must be computed at the level of biological replicates. Read pages 83 & 84

Why should I care about independence?

Lack of independence can bias your results. Mice of the same litter or cells of different wells in a plate are not independent. Read page 51

Removing an outlier is not cheating if based on

Normality tests are not as useful as they appear. Read chapter 24

Should I test for normality?

Should I transform my data? Transformations can have a huge impact on When should I not use a t-test?

Each test depends on assumptions. What to do if some are violated? Read chapter 30

High p-value = no effect?

A p-value > 0.05 means your data are not surprising, but not necessarily that there is no effect.

Read chapters 16, 18 & 19

your conclusions. Be careful. Read page 71-72

How to avoid p-hacking?

All statistical analyses should be planned, and all planned analyses should be conducted and reported.

*Read chapters 22 & 23**

No time? Have a look at least at Chapters 16, 18, 19, 44 & 45

Why not multiple t-tests, but ANOVA?

Such multiple comparisons increase your chances of false positives. Read chapter 39

Which post-hoc test after an ANOVA?

The choice depends on the goals of your study. $Read\ chapter\ 40$

How do I know an effect is important?

The appropriate tool is effect size.

Read chapter 26 & page 468

What should represent my error bars?

SD, SE, SEM or CI? It depends on your aim. $Read\ chapter\ 14$

ADDITIONAL RESOURCES

To perform a power analysis, you can use R or some specific free softwares, e.g. G*Power or www.powerandsamplesize.com Want some more tips? Browse the collection "What is wrong with this picture?" from BMC Biology.