# Scientific Computing – Statistics

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# Day 3 – study design: choosing n

Day 3 – study design: choosing n choosing n for confidence intervals

# general theme

- 1. make an educated guess about the true parameters
- 2. state how accurate/powerful you want to be
- 3. select n based on that

# estimating a single mean standard error and $\alpha$

- Assume you have an estimate *s* of the standard deviation from the literature.
- The 95% confidence interval is given by

$$\underbrace{\left|\tilde{\mu}-\mu_{0}\right|}_{=:\delta} \geq t_{97.5\%,\nu} \frac{s}{\sqrt{n}}$$

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• How should we choose *n* to get a confidence interval of a particular size  $\pm \delta$ ?

We should set n to be

$$n \ge \left(\frac{t_{97.5\%,\nu} \cdot s}{\delta}\right)^2$$

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#### exercise

#### choosing n

Example from last lecture: Literature value of thymus gland weights is 34.3g. The estimate of the standard deviation from the literature is s = 10g.

The equation for n is

$$n \ge \left(\frac{t_{97.5\%,\nu} \cdot s}{\delta}\right)^2$$

- Assume we want to sacrifice as few animals as possible. We say we are fine with a confidence interval of size  $\pm \delta = 5$ , how should we choose *n*?
- What *n* should we choose for *n* if we want  $\pm \delta = 2$ ?

Extend your bootstrapping script from yesterday to check that the equation is correct.

# How to interrupt for/while loops

- Sometimes you want to stop a for/while loop early.
- The command for that is break

#### Example

```
1
   % silly way to find a random number larger than .8
2
   for i = 1:2000
3
     u = rand();
4
     if u >= .8
5
       disp('Found it!');
6
       break
7
     end
8
   end
```

## winner's curse

#### Why it is important to estimate *n* beforehand

Use the thymus gland dataset to repeat the following procedure

- 1. Randomly select n = 10 numbers from the whole dataset.
- 2. Perform a one-sample ttest (ttest) to test against the mean of 34.3g.
- 3. If the p-value is smaller than 0.05, stop the loop and print the mean of the 10 datapoints. Also print the mean of the entire thymus gland dataset.
- 4. Why is it better to use a for instead of a while loop?
- 5. What can you observe? Why does that tell you that choosing *n* is important?

# solution

```
1
    load thymusglandweights.dat
2
3
    n = 10;
4
5
    x = thymusglandweights;
6
7
    for i = 1:5000
8
        idx = randi(length(x), n,1);
9
        y = x(idx);
10
        [h,p] = ttest(y, 34.3);
11
12
        if h == 1
13
            disp(['p-value: ', num2str(p)]);
14
            disp(['mu: ', num2str(mean(y))]);
            disp(['mu total: ', num2str(mean(x))]);
15
16
            break
17
        end
18
    end
```

# Day 3 – study design: choosing n

Day 3 – study design: choosing n choosing n for confidence intervals power

### test nomenclature



#### You want:

- large power
- small type I & II error probability  $(\alpha \text{ and } \beta)$

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#### power

#### estimating power with bootstrapping

- Take the script from yesterday in which we simulated the null distribution of the means.
- Extend it such that it plots the bootstrapped distribution of the means as well (use the same bins for both histograms by using hist for computing the histogram and bar for plotting).
- Use logical indexing to find all means that correspond to true positives (using the 95% decision boundaries computed yesterday). Estimate the power by computing the fraction of true positive bootstrapped means.
- What is the probability that you get a false negative?
- If you have time, plot the histogram of true positives in a different color.

#### summary

- Proper study design is important to avoid statistical problems like the winner's curse.
- You should choose a test with high power.
- There are also equations to select *n* for type I error <u>and</u> power (see book by Zar).

# Overview

- minimizing/maximizing a function numerically (optimization) is ubiquitous in science (curve fitting, maximum likelihood, ...)
- today we will look at the basic elements of optimization and apply it to curve fitting
- tomorrow, we will apply it to maximum likelihood

## plotting surfaces

```
1 range = linspace(-1,1,20);
2 [X,Y] = meshgrid(range, range);
3 
4 surf(X,Y, (X.^2 + Y.^2));
5 colormap('winter');
```

## linear least squares



- The most common curve fitting problem is linear least squares.
- Its goal is to predict a set of output values  $y_1, ..., y_n$  from their corresponding input values  $x_1, ..., x_n$  with a line  $f_{a,b}(x) = ax + b$ .
- How is the line chosen?



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- How is the line chosen?
   By minimization of the mean squared error



$$g(a,b) = \sum_{i=1}^{n} (y_i - f_{a,b}(x_i))^2$$

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### error surface

#### plotting the error surface

Write a function lserr that takes 2-dimensional parameter vector (slope and offset), an array of inputs x, and an array of corresponding outputs y.

Day 4-5 - curve fitting and maximum likelihood

# That's it.

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