Lecture 1: Course Overview

Ciaran Evans

"Statistical computing" vs. "computational statistics"

➤ Statistical computing: programming languages and computing tools for working with statistics (think: storing and accessing data, data transformations and wrangling, some data visualization, etc.)

➤ Computational statistics: the use of computational algorithms to implement statistical methods (think: simulating data from a distribution, fitting a regression model, performing hypothesis testing)

Key points

- ► Focus on *how* to statistical methods work. E.g., how do we fit a logistic regression model? What is R actually doing "under the hood"?
- ► Focus on *implementation* much of your work will be turning statistical methods and algorithms into code
- Focus on efficiency
 - Efficient algorithms (e.g. dynamic programming approaches)
 - Efficient approximations (e.g. integral and Hessian approximations)
 - ► Efficient languages (particularly the use of C++)
- Focus on iteration

Tentative course outline

- Simulation (simulation studies, generating random numbers, simulating from a distribution)
- ► Model fitting (linear and generalized linear models, maximum likelihood, Newton and quasi-Newton methods)
- Missing data and EM algorithm (Gaussian mixtures, Hidden Markov Models)
- Integration (numerical integration, Monte Carlo integration, MCMC)
- Bootstrapping

Roadmap for the first few weeks

- Motivation: simulation studies
 - Today: using simulations to answer questions about hypothesis tests
- Simulation: behind the scenes
 - Generating random numbers
 - Transformation methods
 - Acceptance/rejection sampling
 - Computing topics: iteration, functions, C++
 - Simulation for linear models and multivariate normal distributions
- Following unit: how do you actually fit a regression model? (linear, GLMs, GEEs, etc.)

Suppose we have a sample $X_1,...,X_n \stackrel{iid}{\sim} N(\mu,1)$. Want to test the hypotheses

$$H_0: \mu = 0 \quad H_A: \mu \neq 0$$

How do I test these hypotheses? (What is my test statistic, and how do I make a decision?)

Suppose we have a sample $X_1,...,X_n \stackrel{\it iid}{\sim} N(\mu,1)$. Want to test the hypotheses

$$H_0: \mu = 0 \quad H_A: \mu \neq 0$$

If actually $\mu = 0$, do we want to reject H_0 , or fail to reject?

- ightharpoonup Either H_0 is true or false (usually we don't know)
- \blacktriangleright We either reject or fail to reject H_0

Possible outcomes of a hypothesis test:

	H ₀ is true	H ₀ is false
fail to reject reject	correct decision type I error	type II error correct decision

Usual goal: Minimize type II error, subject to control of type I error rate (i.e. want $P(\text{reject } H_0|H_0 \text{ true}) <= \alpha)$

In R

```
mu x <- 0
n <- 20
x <- rnorm(n, mean=mu x, sd=1)
X
   [1] 0.6797336 -0.8548404 -1.3210164 -0.4273575 -1.020
##
## [7] -0.4315816 -0.3921952 1.0212251 -0.1325580 -1.0810
## [13] 0.7787329 0.5623875 0.1579840 2.0061821 0.155°
## [19] -0.3435500 -0.2215555
t.test(x, alternative="two.sided", mu=0)
##
##
    One Sample t-test
##
## data: x
## t = 0.33834, df = 19, p-value = 0.7388
## alternative hypothesis: true mean is not equal to 0
```

In R

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t.test(x, alternative="two.sided", mu=0)
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   One Sample t-test
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## data: x
## t = 0.33834, df = 19, p-value = 0.7388
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## -0.3297764 0.4569528
## sample estimates:
## mean of x
## 0.06358819
```

Here we fail to reject H_0 . Does this give us our type I error rate?

In R

t.test(x, alternative="two.sided", mu=0)

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##
    One Sample t-test
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## data:
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```

Here we fail to reject H_0 . Does this give us our type I error rate?

No! Type I error rate = $P(\text{reject } H_0|H_0 \text{ true})$. Need more than one observation to estimate a probability.

Repeating many times

```
set.seed(379)
n < -20
mu x <- 0
nsim <- 1000 # number of times to repeat
test results <- rep(NA, nsim) # vector to store the test
for(i in 1:nsim){
  x \leftarrow rnorm(n, mean=mu x, sd=1)
  test results[i] <- t.test(x, alternative="two.sided", must
head(test results)
```

[1] 0.8213916 0.2525886 0.3818531 0.3760550 0.7342035 0

What is this code doing?

Repeating many times

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```

[1] 0.8213916 0.2525886 0.3818531 0.3760550 0.7342035 0

How do I find the fraction of times we rejected H_0 ?

Repeating many times

```
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n < -20
mu x <- 0
nsim <- 1000 # number of times to repeat
test results <- rep(NA, nsim) # vector to store the test
for(i in 1:nsim){
  x \leftarrow rnorm(n, mean=mu x, sd=1)
  test results[i] <- t.test(x, alternative="two.sided", must
}
mean(test_results < 0.05)</pre>
```

[1] 0.045

Your turn: Practice questions

Practice questions on the course website:

https://sta379-s25.github.io/practice_questions/pq_1.html

- Experiment with changing μ and n. How does the probability of rejecting change?
- ▶ Start in class. You are welcome to work with others
- Practice questions are to help you practice. They are not submitted and not graded
- Solutions are posted on the course website