001 - Introduction to Inferential Statistics

EPIB 607 - FALL 2020

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slides compiled on September 1, 2020



Ohiectives

Visualize/Analyze/Interpret data using statistical methods with a computer

Gather data into analysis ready format

Learn regression

Understand the statistical results in a scientific paper

Learn the tools for creating reproducible analyses

Where does this course fit in my life?

Objectives 2/38.

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Objectives 3/38.

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Where does this course fit in my life?

Data is the new oil¹

Fuel of the future

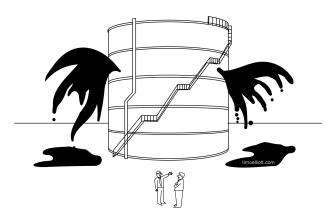
Data is giving rise to a new economy

How is it shaping up?



¹ https://www.economist.com/briefing/2017/05/06/data-is-giving-rise-to-a-new-economy Visualize/Analyze/Interpret data using statistical methods with a computer

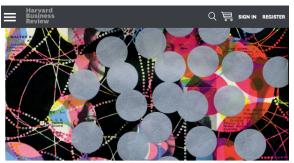
Danger²



"Data is the new oil? Absolutely—toxic if mishandled!..."

 $^{^2 \\ \}text{https://timoelliott.com/blog/2018/03/data-is-the-new-oil-yes-toxic-if-mishandled.html}$

Data science³



ARTWORK: TAMAR COHEN, ANDREW J BUBOLTZ, 2011, SILK SCREEN ON A PAGE FROM A HIGH SCHOOL YEARBOOK, 8.5" X 12"

DATA

Data Scientist: The Sexiest Job of the 21st Century

by Thomas H. Davenport and D.J. Patil

FROM THE OCTOBER 2012 ISSUE

³ https://hbr.org/2012/10/data-scientist-the-sexiest-job-of-the-21st-century Visualize/Analyze/Interpret data using statistical methods with a computer

Why R?

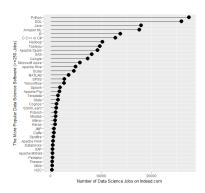


Figure: Data as of May 2019
http://r4stats.com/articles/popularity/

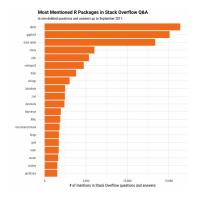


Figure: Popular R packages https://

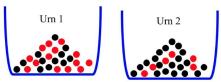
 ${\tt stackoverflow.blog/2017/10/10/impressive-growth-r/}$

First day in a statistics course

Example:

We have two urns. Urn 1 contains 14 red balls and 12 black balls. Urn 2 contains 6 red balls and 20 black balls.

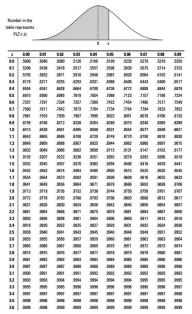
An Urn is selected at random and a ball is selected from that urn.



If the ball turns out to be red what is the probability that it came from the first urn?

	Sepal.Length [‡]	Sepal.Width [‡]	Petal.Length [‡]	Petal.Width [‡]	Species
1	5.1	3.5	1.4	0.2	setosa
2	4.9	3.0	1.4	0.2	setosa
3	4.7	3.2	1.3	0.2	setosa
4	4.6	3.1	1.5	0.2	setosa
5	5.0	3.6	1.4	0.2	setosa
6	5.4	3.9	1.7	0.4	setosa
7	4.6	3.4	1.4	0.3	setosa
8	5.0	3.4	1.5	0.2	setosa
9	4.4	2.9	1.4	0.2	setosa
10	4.9	3.1	1.5	0.1	setosa
11	5.4	3.7	1.5	0.2	setosa
12	4.8	3.4	1.6	0.2	setosa
13	4.8	3.0	1.4	0.1	setosa
14	4.3	3.0	1.1	0.1	setosa
15	5.8	4.0	1.2	0.2	setosa

Second day in a statistics course



Objectives

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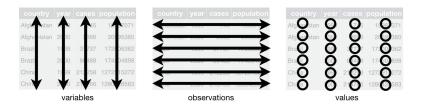
Understand the statistical results in a scientific paper

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Where does this course fit in my life?

Tidy data

- Each variable forms a column.
- Each observation forms a row.
- Each type of observational units forms a table
- Tidy data is ready for regression routines and plotting



Example: Does a full moon affect behaviour?

- Many people believe that the moon influences the actions of some individuals.
- A study of dementia patients in nursing homes recorded various types of disruptive behaviors every day for 12 weeks.
- Days were classified as moon days if they were in a 3-day period centered at the day of the full moon.
- For each patient, the average number of disruptive behaviors was computed for moon days and for all otherdays.

patient	moon_days	other_days
1	3.33	0.27
2	3.67	0.59
3	2.67	0.32
4	3.33	0.19
5	3.33	1.26
6	3.67	0.11
7	4.67	0.30

patient	moon_days	other_days
1	3.33	0.27
2	3.67	0.59
3	2.67	0.32

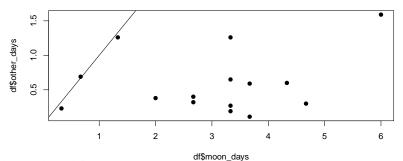
moon_days	other_days
3.33	0.27
3.67	0.59
2.67	0.32
	3.33 3.67

Question: Can I plot the data?

pat	ient	moon_days	other_days
	1	3.33	0.27
	2	3.67	0.59
	3	2.67	0.32

Question: Can I plot the data?

```
plot(df$moon_days, df$other_days, pch = 19)
abline(a=0,b=1)
```



patient	moon_days	other_days
1	3.33	0.27
2	3.67	0.59
3	2.67	0.32
4	3.33	0.19
5	3.33	1.26

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Question: Can I fit a meaningful regression model directly to the variables in the data?

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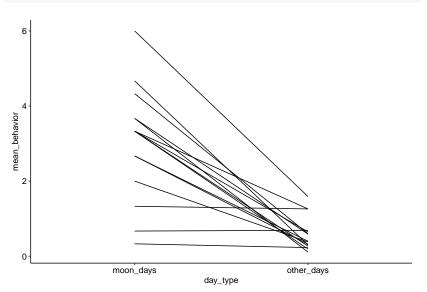
Question: Can I fit a meaningful regression model directly to the variables in the data?

```
## Call: lm(formula = moon_days ~ other_days, data = df)
##
## Coefficients:
## Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.56 0.66 3.9 0.002
## other_days 0.79 0.91 0.9 0.402
##
## Residual standard error: 1.5 on 13 degrees of freedom
## Multiple R-squared: 0.055,^^IAdjusted R-squared: -0.018
## F-statistic: 0.75 on 1 and 13 DF, p-value: 0.4
```

patient	day_type	mean_behavior
1	moon_days	3.33
1	other_days	0.27
2	moon_days	3.67
2	other_days	0.59
3	moon_days	2.67
3	other_days	0.32
4	moon_days	3.33
4	other_days	0.19
5	moon_days	3.33
5	other_days	1.26

Plotting with tidy data

ggplot(data = df_tidy, mapping = aes(x = day_type, y = mean_behavior, group = patient)) + geom_line()



Regression with tidy data

```
fit <- lme4::lmer(mean_behavior ~ day_type + (1|patient), data = df_tidy)
summary(fit)
## Linear mixed model fit by REML ['lmerMod']
## Formula: mean behavior ~ day type + (1 | patient)
     Data: df_tidy
##
## REML criterion at convergence: 90.3
##
## Scaled residuals:
       Min 10 Median 30
                                        Max
## -2.27236 -0.30142 -0.04023 0.48540 2.44753
##
## Random effects:
## Groups Name
                  Variance Std.Dev.
## patient (Intercept) 0.1563 0.3954
## Residual
                      1.0659 1.0324
## Number of obs: 30, groups: patient, 15
##
## Fixed effects:
##
                    Estimate Std. Error t value
## (Intercept) 3.0220 0.2854 10.587
## day_typeother_days -2.4327 0.3770 -6.453
##
## Correlation of Fixed Effects:
##
             (Intr)
## dy_typthr_d -0.660
```

Not tidy vs. tidy data

patient	moon_days	other_days
1	3.33	0.27
2	3.67	0.59
3	2.67	0.32
4	3.33	0.19
5	3.33	1.26

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Not tidy vs. tidy data

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Not tidy

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4	other_days	0.19
5	moon_days	3.33
5	other_days	1.26

tidyr::pivot_longer()

patient	moon_days	other_days			
1	3.33	0.27			
2	3.67	0.59			
3	2.67	0.32			
4	3.33	0.19	_		
5	3.33	1.26			
				kěy	value
			patient	day_type	mean_behavior
			1	moon_days	3.33
					0.07

patient	udy_type	IIIcaii_bellavioi
1	moon_days	3.33
1	other_days	0.27
2	moon_days	3.67
2	other_days	0.59
3	moon_days	2.67
3	other_days	0.32
4	moon_days	3.33
4	other_days	0.19
5	moon_days	3.33
5	other_days	1.26

tidyr::pivot_longer(data = df, cols = -patient, names_to = "day_type", values_to = "mean_behavior")

Objectives

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Understand the statistical results in a scientific paper

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Where does this course fit in my life?

Learn regression 22/38.

Traditional stats textbook

CHAPTER 7

Hypothesis Testing: One-Sample Inference / 211

- 7.1 Introduction / 211
- 7.2 General Concepts / 211
- One-Sample Test for the Mean of a Normal Distribution: One-Sided Alternatives / 214
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- 7.6 The Power of a Test / 232
- 7.7 Sample-Size Determination / 239

- One-Sample χ² Test for the Variance of a Normal Distribution / 245
- One-Sample Inference for the Binomial Distribution / 249

 7.10 One-Sample Inference for the Poisson
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- Case Study: Effects of Tobacco Use on Bone-Mineral Density in Middle-Aged Women / 265
- 7.12 Derivation of Selected Formulas / 265 7.13 Summary / 267

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- Two-Sample t Test for Independent Sample with Unequal Variances / 298
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- Contrate
- Case Study: Effects of Lead Exposure on Neurologic and Psychological Function in Children / 305
- 8.9 Estimation of Sample Size and Power for Comparing Two Means / 307
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Regression and Correlation Methods / 457

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- 11.7 The Correlation Coefficient / 485
- 11.8 Statistical Inference for Correlation Coefficients / 490
- 11.9 Multiple Regression / 502
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Learn regression 23/38.

This course

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 - 7.11 Case Study: Effect co Use on Bone Mineral Density in Mi d Women / 2 7.12 Derivation of Selected
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 - Variances / 292 Two-S endent Samples
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- 8.8 Case Study: Effects of Lead Exposure on Neurologic and Psychological Function in
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Statistical concepts

RESULTS The total populations were 462 445 in the lowarborder counties and 272 385 in the Illinois border counties. Population density was higher in the lowa counties (114.2 people per square mile) than in the Illinois counties (78.2 people per square mile). Trends of cumulative COVID-19 cases per 10 000 residents for the lowa and Illinois border counties were comparable before the Illinois stayat-home order, which went into effect at 5:00 PM on March 21 (March 15 to March 21: 0.024 per 10 000 residents vs 0.026 per 10 000 residents). After that, cases increased more quickly in Iowa and more slowly in Illinois. Within 10, 20, and 30 days after the enactment of the stay-at-home order in Illinois, the difference in cases was -0.51 per 10 000 residents (SE, 0.09; 95% CI, -0.69 to -0.32; P < .001), -1.15 per 10 000 residents (SE, 0.49; 95% Cl, -2.12 to -0.18; P = .02), and -4.71 per 10 000 residents (SE, 1.99: 95% CI, -8.64 to -0.78: P = .02), respectively. The estimates indicate excess cases in the border lowa counties by as many as 217 cases after 1 month without a stay-at-home order. This estimate of excess cases represents 30.4% of the 716 total cases in those lowa counties by that date. Sensitivity analyses addressing differences in timing of closing schools and nonessential businesses and differences in county population density and poverty rates between the 2 states supported these findings.

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⁴ https://jamanetwork.com/journals/jamanetworkopen/fullarticle/2766229 Understand the statistical results in a scientific paper

Statistical concepts

Table 1. Difference-in-Differences Estimates of COVID-19 Cases Comparing Border Counties in Iowa With Those in Illinois Before and After the Stay-at-Home Order Was Issued in Illinois^a

Period	Difference in COVID-19 cases per 10 000 residents ^b	Heteroskedasticity robust SE (95% CI) ^c	P value	Excess cases in lowa border counties	Excess cases as proportion of total cases, %
3/22-3/26	-0.14	0.04 (-0.23 to -0.06)	.001	6	32.4
3/27-3/31	-0.51	0.09 (-0.69 to -0.32)	<.001	24	38.0
4/01-4/05	-0.41	0.17 (-0.74 to -0.07)	.02	19	15.2
4/06-4/10	-1.15	0.49 (-2.12 to -0.18)	.02	53	17.8
4/11-4/15	-3.35	1.19 (-5.70 to -0.99)	.006	154	30.0
4/16-4/20	-4.71	1.99 (-8.64 to -0.78)	.02	217	30.4

Abbreviation: COVID-19, coronavirus disease 2019.

5

^a The regression model was estimated separately for each of 5-day period. The regression was estimated using least squares weighted by the 2019 county population. The regression adjusted for county and day fixed effects. The number of county-day observations was 180 for each regression.

^b This indicates the estimated difference-in-differences association of a stay-at-home order with COVID-19 cases in a given period relative to March 15 to March 21 (ie, the period before the stay-at-home order in Illinois was enacted).

^c Heteroskedasticity robust SEs were estimated because homoscedasticity is rejected for all post-period regressions.

⁵https://jamanetwork.com/journals/jamanetworkopen/fullarticle/2766229
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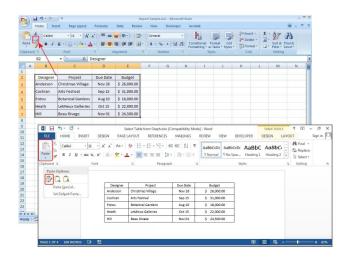
Learn regression

Understand the statistical results in a scientific paper

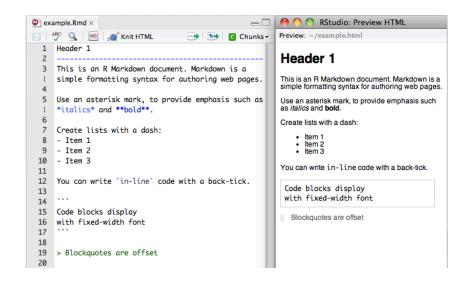
Learn the tools for creating reproducible analyses

Where does this course fit in my life?

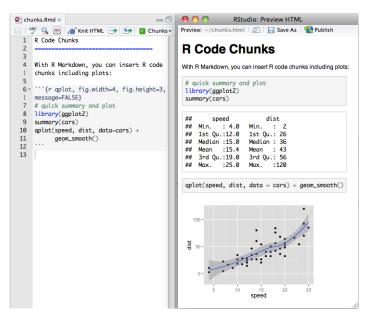
Copy paste ad nauseam



Markdown: HTML without knowing HTML

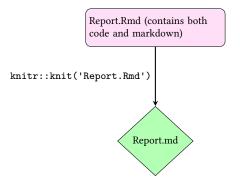


R + Markdown = RMarkdown



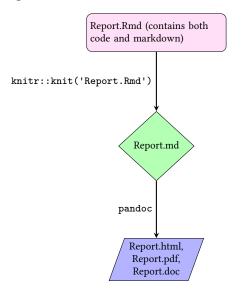
What rmarkdown does

RMarkdown example:



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RMarkdown example:

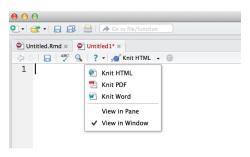


Compiling a . Rmd document

The two steps on previous slide can be executed in one command:

rmarkdown::render()

or in RStudio:



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Topics by level of exposure

Level of Exposure



Mainstream media

- Scientific Article
- Statistical Analysis
- Cleaning Data
- Collecting Data
- Research Ethics Board (REB) Approval
- Study Design
- Research Question
- Statistical Methods Development

First year courses

Mainstream media evel of Exposure Scientific Article Statistical Analysis EPIB 607/621 **EPIB 613** Cleaning Data Collecting Data Research Ethics Board (REB) Approval Study Design FPIB 601/602 Research Question Statistical Methods Development



- Scientific Article
- Statistical Analysis
- Cleaning Data
- Collecting Data
- Research Ethics Board (REB) Approval
- Study Design
- Research Question

EPIB 601/602

EPIB 607/621 EPIB 613

Statistical Methods Development

What I do

Session Info

```
R version 3.6.2 (2019-12-12)
Platform: x86_64-pc-linux-gnu (64-bit)
Running under: Pop! OS 19.10
Matrix products: default
        /usr/lib/x86_64-linux-gnu/openblas/libblas.so.3
BLAS:
LAPACK: /usr/lib/x86_64-linux-gnu/libopenblasp-r0.3.7.so
attached base packages:
[1] tools
                        graphics
                                  grDevices utils
                                                       datasets methods
              stats
[8] base
other attached packages:
 [1] NCStats 0.4.7
                        FSA 0.8.30
                                            forcats 0.5.0
                                                               stringr 1.4.0
 [5] dplyr_1.0.2
                        purrr_0.3.4
                                            readr_1.3.1
                                                               tidyr_1.1.2
 [9] tibble 3.0.3
                        ggplot2_3.3.2.9000 tidyverse_1.3.0
                                                               knitr_1.29
loaded via a namespace (and not attached):
 [1] Rcpp_1.0.4.6
                        lubridate_1.7.4
                                            lattice_0.20-38
                                                               assertthat_0.2.1
 [5] digest_0.6.25
                        R6_2.4.1
                                            cellranger 1.1.0
                                                               plvr 1.8.6
 [9] backports_1.1.9
                        reprex_0.3.0
                                            evaluate 0.14
                                                               httr 1.4.1
[13] highr_0.8
                        pillar_1.4.6
                                            TeachingDemos_2.12 rlang_0.4.7
[17] readxl 1.3.1
                        rstudioapi 0.11
                                            minga 1.2.4
                                                               nloptr 1.2.2.1
[21] Matrix 1.2-18
                        labeling 0.3
                                            splines 3.6.2
                                                               lme4 1.1-21
[25] munsell_0.5.0
                        broom_0.7.0
                                            compiler_3.6.2
                                                               modelr_0.1.5
[29] xfun_0.16
                        pkgconfig_2.0.3
                                            tidyselect_1.1.0
                                                               fansi_0.4.1
[33] crayon_1.3.4
                        dbplyr_1.4.2
                                            withr 2.2.0
                                                               ggpubr_0.2.5
[37] MASS_7.3-51.5
                        grid_3.6.2
                                            nlme_3.1-143
                                                               jsonlite_1.7.0
[41] gtable_0.3.0
                        lifecycle_0.2.0
                                            DBI_1.1.0
                                                               magrittr_1.5
[45] scales 1.1.1
                        cli 2.0.2
                                            stringi 1.4.6
                                                               farver 2.0.3
[49] ggsignif_0.6.0
                        fs_1.3.2
                                            xm12_1.3.0
                                                               ellipsis_0.3.1
[53] generics_0.0.2
                        vctrs_0.3.4
                                            boot_1.3-24
                                                               glue_1.4.2
[57] hms 0.5.3
                        colorspace 1.4-1
                                            rvest 0.3.5
                                                               haven 2.3.1
```