### **Introduction to Inferential Statistics**

#### EPIB 607 - FALL 2020

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slides compiled on August 30, 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.

#### **Objectives**

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Where does this course fit in my life?

### Data is the new oil<sup>1</sup>

#### Fuel of the future

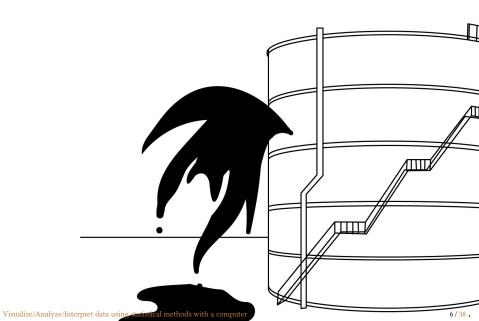
### Data is giving rise to a new economy

How is it shaping up?



<sup>&</sup>lt;sup>1</sup>https://www.economist.com/briefing/2017/05/06/data-is-giving-rise-to-a-new-economy

# $Danger^2 \\$



### Data science<sup>3</sup>



Data Scientist: The Sexiest Job of the 21st Century

by Thomas H. Davenport and D.J. Patil MIGHTHE OCTOBER 2012 ISSUE

<sup>&</sup>lt;sup>3</sup>https://hbr.org/2012/10/data-scientist-the-sexiest-job-of-the-21st-century

# Why R?

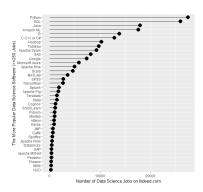


Fig. 1: Data as of May 2019

http://r4stats.com/articles/popularity/

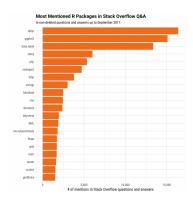


Fig. 2: Popular R packages

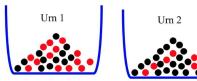
https://stackoverflow.blog/2017/10/10/impressive-growth-r/

### 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.

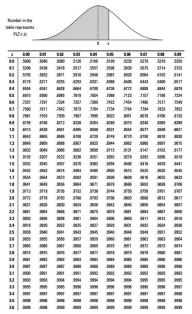
An Urn is selected at random and a ball is selected from that urn.



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

	Sepal.Length <sup>‡</sup>	Sepal.Width <sup>‡</sup>	Petal.Length <sup>‡</sup>	Petal.Width <sup>‡</sup>	Species <sup>‡</sup>
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**

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

### Gather data into analysis ready format

Learn regression

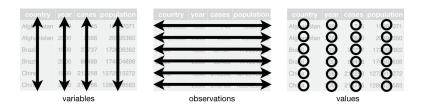
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Where does this course fit in my life?

## 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

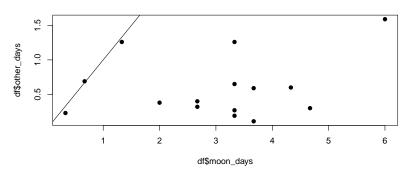
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3	2.67	0.32

**Question: Can I plot the data?** 

patient	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
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Question: Can I fit a <u>meaningful</u> regression model <u>directly</u> to the <u>variables</u> in the <u>data?</u>

patient	moon_days	other_days
1	3.33	0.27
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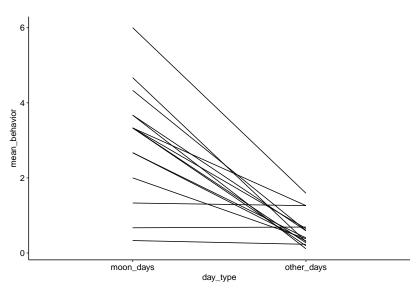
# Question: Can I fit a meaningful regression model directly to the variables in the data?

```
## Call: lm(formula = moon\_days \sim 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 tidv
##
## 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
##
## 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
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### Not tidy vs. tidy data

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Not tidy

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tidy

# 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				
		1			kěy	value
				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

moon\_days

other\_days

moon\_days other\_days

moon\_days other\_days

tidyr::pivot\_longer(data = df, cols = -patient, names\_to = "day\_type", values\_to = "mean\_behavior")

2.67

0.32

0.19

1.26

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Where does this course fit in my life?

Learn regression 22/38.

#### Traditional stats textbook

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- 7.2 General Concepts / 211
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- Mineral Density in Middle-Aged Women / 265 7.12 Derivation of Selected Formulas / 265
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23/38. Learn regression

### This course

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### Statistical concepts

**RESULTS** The total populations were 462 445 in the lowa border 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% CI, -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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<sup>&</sup>lt;sup>4</sup>https://jamanetwork.com/journals/jamanetworkopen/fullarticle/2766229

### 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<sup>a</sup>

Period	Difference in COVID-19 cases per 10 000 residents <sup>b</sup>	Heteroskedasticity robust SE (95% CI) <sup>c</sup>	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.

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<sup>&</sup>lt;sup>a</sup> 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.

<sup>&</sup>lt;sup>b</sup> 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).

<sup>&</sup>lt;sup>c</sup> Heteroskedasticity robust SEs were estimated because homoscedasticity is rejected for all post-period regressions.

<sup>&</sup>lt;sup>5</sup>https://jamanetwork.com/journals/jamanetworkopen/fullarticle/2766229

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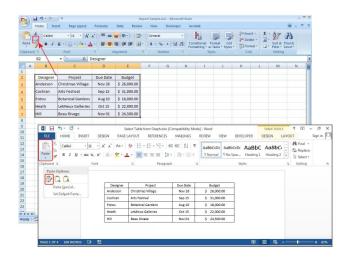
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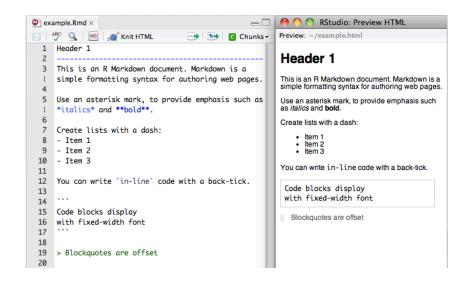
Learn the tools for creating reproducible analyses

Where does this course fit in my life?

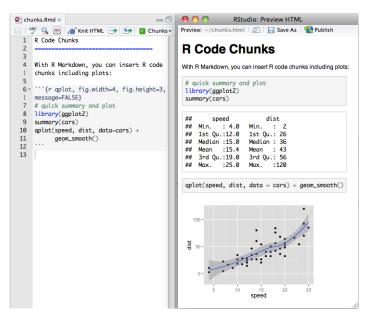
### Copy paste ad nauseam



### Markdown: HTML without knowing HTML

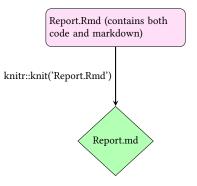


### R + Markdown = RMarkdown



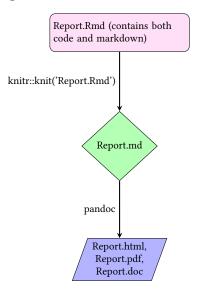
### What rmarkdown does

### RMarkdown example:



### What rmarkdown does

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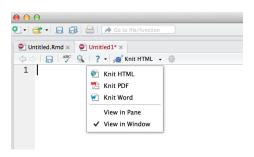


## Compiling a .Rmd document

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 BLAS: /usr/lib/x86\_64-linux-gnu/openblas/libblas.so.3 LAPACK: /usr/lib/x86 64-linux-gnu/libopenblasp-r0.3.7.so attached base packages: [1] stats graphics grDevices utils datasets methods 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 readr 1.3.1 purrr 0.3.4 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 plyr\_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 minqa\_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 pkgconfig\_2.0.3 tidyselect\_1.1.0 fansi 0.4.1 [29] xfun 0.16 [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 xml2 1.3.0 ellipsis 0.3.1 [53] generics\_0.0.2 vctrs 0.3.4 boot 1.3-24 tools 3.6.2 [57] glue\_1.4.2 hms 0.5.3 colorspace 1.4-1 rvest 0.3.5 [61] haven 2.3.1