Reproducibility and Experimental Design

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Overview

- ► What is reproducibility?
- Source of bias
- Experimental Design
- ► Replication (n = ?)
- Recommendations

Source: https://github.com/tschauer

Reference Book



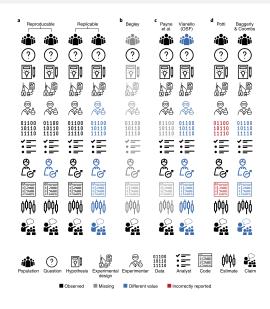
Lazic, 2016

Reproducibility vs. replicablity



Patil, Peng and Leek, 2019

Reproducibility vs. replicablity

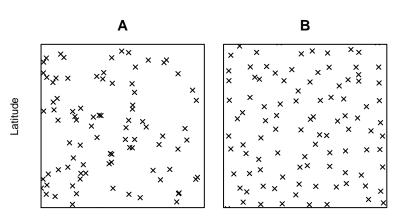


Patil, Peng and Leek, 2019

Reproducibility by Lazic

- ▶ analytical: original data and analysis (code!)
- direct: same conditions, materials, methods
- systematic: different conditions (e.g. cell line, KD vs drug)
- conceptual: general under diverse conditions (paradigm)

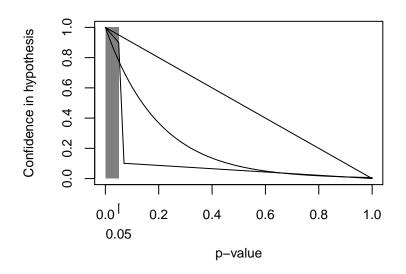
Strategy: which location was bombed randomly?



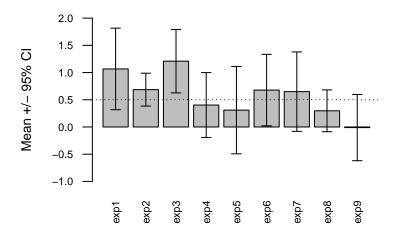
Longitude

- seeing pattern in randomness
- not wanting to miss anything (what else can we get out?)
- ▶ if a hypothesis is derived from the data, then the ability of the data to support that hypothesis is diminished
- exploratory vs. confirmatory research

ightharpoonup psychological cliff at p = 0.05



neglect of sampling variability



ightharpoonup mean = 0.5, standard deviation = 1, n=10 each

- ► lack of independence
 - repeated measures
 - observations are close together in space or time
 - same animal, litter, cell culture dish, fly vial
 - correlated variables
 - different measures of a single underlying effect
 - co-regulated genes, proteins, metabolites
 - disease severity

- confirmation bias
 - ▶ Pubmed search: disease + gene name
 - what about studies which do not find the association?
 - neglecting negative results
 - data transformation until it "gets" significant
 - selecting data to tell the story (data that do not fit excluded)

- expectancy effects (measurements are influenced)
- hindsight bias ('I knew it all along')
- herding effect (scientific inbreeding)

Common problems

- Experimental Design
 - confounding (conditions ~ biological, technical effects)
 - experimental unit (replicates)
 - lack of randomization
 - low statistical power
- Conducting experiments
 - lack of blinding
 - lack of randomization
 - optional stopping

Common problems

- Analysis
 - experimental unit (inflated sample size)
 - ▶ inapproriate model (normal distribution)
 - incorrect interpretation
 - selective reporting

Experimental Goal

	Exploratory	Confirmatory
Question	General	Specific
Hypothesis	Generating	Before
Order	Before	After (independent data)
Analysis	Data dependent	Data independent
Minimize	False Negatives	False Positives
P-value	No Diagnostic Value	Diagnostic Value
Power Analysis	Rarely	Yes

Experimental Goal

Subjects Heterogeneous Homogeneous Environment Varied Standardised Treatments Many Few Levels Many Few Time points Many Few Outcome Many Few Controls Few Many Blinding Possibly Yes			
Environment Varied Standardised Treatments Many Few Levels Many Few Time points Many Few Outcome Many Few Controls Few Many Blinding Possibly Yes		Exploratory	Confirmatory
Treatments Many Few Levels Many Few Time points Many Few Outcome Many Few Controls Few Many Blinding Possibly Yes	Subjects	Heterogeneous	Homogeneous
Levels Many Few Time points Many Few Outcome Many Few Controls Few Many Blinding Possibly Yes	Environment	Varied	Standardised
Time points Many Few Outcome Many Few Controls Few Many Blinding Possibly Yes	Treatments	Many	Few
Outcome Many Few Controls Few Many Blinding Possibly Yes	Levels	Many	Few
Controls Few Many Blinding Possibly Yes	Time points	Many	Few
Blinding Possibly Yes	Outcome	Many	Few
zg	Controls	Few	Many
D 1 : .:	Blinding	Possibly	Yes
Randomization Yes Yes	Randomization	Yes	Yes
Blocking Yes Yes	Blocking	Yes	Yes

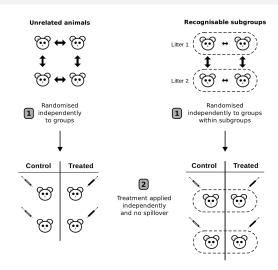
Hypothesis testing?

- p-value
 - ➤ A: "Given these data what is the probability that the null hypothesis is true"
 - ▶ B: "Given that the null hypothesis is true, what is the probability of these (or more extreme) data"
- "The Earth is round (p < .05)" Cohen, 1994
- ► Solution? Pre-registration: https://cos.io/prereg/

Experimental Design Equation

Outcome =	${\sf Treatment} \ +$	$Biological \; + \;$	$Technical\ +$	Error
Gene exp. Protein Cell counts	Environment Compound Inhibitor siRNA Dose Time	Sex Age Weight Litter Genotype Cell line	Person Batch Flask Cage Day Incubator	Experimental Treatment Sampling Measurement

Randomization



Completely randomized vs. randomized blocked design

Lazic, 2018

2-factor design

crossed

	Control	Treated
Wild type	****	****
Mutant	****	****

nested

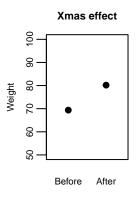
	Control	Treated
Cage 1	****	
Cage 2	****	
Cage 3		****
Cage 4		****

2-factor design

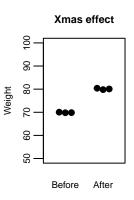
confounded

	Control	Treated
Day 1	******	
Day 2		******

▶ How does Christmas affect human body weight?

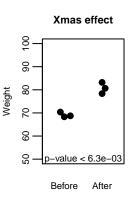


measuerment error (3x within minutes)



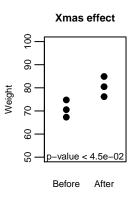
- qPCR well-replicates
- sequencing the same library

Different days same person



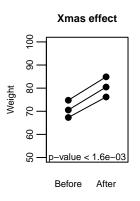
- Christmas signficantly increases human body weight ???
- e.g. cell culture experiments (generalizable?)

Different years same person



year can be used as grouping factor

▶ Different years same person

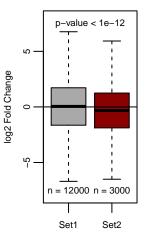


- ▶ it is still just a single person
- ▶ different years not always applicable

Batch effects

What is n?

within sample testing (comparing set of genes)



- ► True difference: -0.25
- ▶ large number of genes inflates p-value (no diagnostic value)