# Significance and Hypothesis testing 

## Martin Popel

ÚFAL (Institute of Formal and Applied Linguistics)
Charles University in Prague
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## Motivation

Reporting significance and confidence intervals is ubiquitous in quantitative research.

Goals of this lecture

- Understand the basic principles (and names).

Understand papers, e.g.
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# Motivation <br> Reporting significance and confidence intervals is ubiquitous in quantitative research. 

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- Understand the basic principles (and names).

Understand papers, e.g.
"significantly better than the baseline ( $p<0.05$ )"
Does it mean "much better'? No!
Don't use "significant" unless you can prove it! So what does it mean?

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## Fisher vs. Neyman \& Pearson

They were rivals, their approaches are not compatible.


## Recap: Statistics

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- standard deviation $(s, \sigma)$, variance $\left(s^{2}, \sigma^{2}\right)$,
- median, Xth quantile,
- for difference tests: difference mean, difference median,...
- BLEU, LAS, $\mathrm{F}_{1}$-score,...


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- one-sample
- two-sample (difference test)
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- paired
correlated samples have lower variance of the difference mean


## P-value

Null hypothesis $\left(H_{0}\right)$ :

- no effect, status quo, what could be expected
- defines a distribution
$P$-value is:
- "the probability of obtaining a test statistic result at least as extreme as the one that was actually observed, assuming that the null hypothesis is true"
- $p=P\left(\right.$ data or more extreme $\left.\mid H_{0}\right)$
- informal measure of evidence against $H_{0}$

P -value is not:

- $P\left(H_{0}\right), P\left(H_{0} \mid\right.$ data $), 1-P\left(H_{A}\right)$ (see Lindley's paradox)
- size or importance of the observed effect
- probability that the measured effect is just a random fluke
- probability of falsely rejecting $H_{0}$, i.e. false positive error rate, i.e. Type I error rate


## Significance level

Fisher's Significance level:

- popular but arbitrary value is 0.05 (or 0.01 in some areas)
- threshold for p -values (reject $H_{0}$ if $p<0.05$ )
- sometimes called $\alpha$, but should not be confused with Neyman\&Pearson's $\alpha=$ Type I error rate.
- should be set before the experiment (prior to data collection)

It is better to report the (rounded) p-value instead of just $p<0.05$.


A p-value (shaded green area) is the probability of an observed (or more extreme) result arising by chance

## Experiment 1: Five heads in a row

- Story: A magician claims to bias a coin toward more heads.
- Experiment: Flip a coin 5 times (i.e. sample size $=5$ ).
- Result: HHHHH (i.e. five heads in a row)
- Analysis: p-value =
- Conclusion:


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- Result: HHHHH (i.e. test statistic $=5$ )
- Analysis: p-value $=P\left(H H H H H\right.$ or more $\left.H_{0}\right)=\left(\frac{1}{2}\right)^{5} \doteq 0.03$ Event HHHHH is significant, p-value $<0.05$.
- Conclusion: Reject $H_{0}$ (on the 0.05 significance level). Either $H_{0}$ is false or a highly unprobable event occured.


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- Test statistic: total number of heads
- Significance level: 0.05 (i.e. confidence level $=95 \%$ )
- One vs. two tails: two-tailed test or alternative hypothesis $H_{A}: p($ head $) \neq 0.5$
- Result: HHHHH (i.e. test statistic $=5$ )
- Analysis: p-value $=P\left(H H H H\right.$ or more $\left.H_{0}\right)=2 \cdot\left(\frac{1}{2}\right)^{5} \doteq 0.06$ Event HHHHH is not significant, p-value $>0.05$.
- Conclusion: Cannot reject $H_{0}$ (on the 0.05 significance level).


## Experiment 1 moral

One tail vs. two tails: It matters.

p-value-two-tailed $=2 \cdot$ p-value-one-tailed (for symmetric $H_{0}$ ) Which one is more strict?

## Experiment 2: Sample size

Test statistic ( $x$ ): proportion of heads

- HHHHH (5 heads out of 5 flips): $x=1$

$$
p_{\text {two-tailed }}=\frac{1}{16} \doteq 0.06
$$

- HHHHHHHHHH (10 heads out of 10 flips): $x=1$ $p_{\text {two-tailed }}=$
- HHHHHHTHHH (9 heads out of 10 flips): $x=0.9$
$p_{\text {two-tailed }}=$


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- HHHHHHHHHH (10 heads out of 10 flips): $x=1$

$$
p_{\text {two-tailed }}=2 \cdot \frac{1}{2^{10}}=\frac{1}{512} \doteq 0.002
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- HHHHHHTHHH ( 9 heads out of 10 flips): $x=0.9$

$$
p_{\text {two-tailed }}=2 \cdot \frac{1+10}{2^{10}}=\frac{11}{512} \doteq 0.02
$$

Experiment 2 morals:

- Sample size matters.
- P-value conflates effect size and our confidence.


## Experiment 3: Alternating coin flips

Null hypothesis: fair coin

Test statistic: number of heads

- HTHTHTHTHT:
$p_{\text {two-tailed }}=$

Test statistic $(x)$ : number of "alternations" ("HT" or "TH")

- HTHTHTHTHT:
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Experiment 3 morals:

- Test statistic matters.


## Confidence Interval

Always report confidence interval for a statistic! E.g. $\mathrm{BLEU}=12.1$ ([10.6; 12.5])

What influences the size of a confidence interval?

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What influences the size of a confidence interval?

- level of confidence (e.g. 95\% confidence interval)


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Always report confidence interval for a statistic!
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What influences the size of a confidence interval?

- level of confidence (e.g. 95\% confidence interval)
- population variance


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- population variance
- sample size


## How to compute confidence interval?

There are three ways

- informal
- traditional normal-based formula
- bootstrapping


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- informal Median $\pm 1.5 \cdot \frac{I Q R}{\sqrt{n}}$ $I Q R=$ Inter-Quartile Range $=Q_{3}-Q_{1}$ $\sim 99 \%$ confidence interval

- traditional normal-based formula
- bootstrapping



## Normal-based CI

traditional normal-based formula $\bar{x} \pm t \cdot s t d . e r r$

- standard error $=\frac{s}{\sqrt{n}}=\frac{\text { sample standard deviation }}{\sqrt{\text { sample size }}}$ $\mathrm{t}=\mathrm{t}$-statistic $=$ function (confidence level, df ) $\mathrm{df}=\mathrm{n}-1=$ degrees of freedom
- from scipy.stats import t; print t.ppf(0.975, 99)
- Excel, Calc: $\operatorname{TINV}(0.05,99)$
- https://www.wolframalpha.com/input/?i=t-interval

For example: $n=100, s=1, \bar{x}=10$ the $95 \%$ interval is
$95 \%$ of (population) values lie within this interval. True or false?

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For example: $n=100, s=1, \bar{x}=10$ the $95 \%$ interval is $10 \pm 0.198$
$95 \%$ of (population) values lie within this interval. True or false? False. We are $95 \%$ sure that the population mean lies within this interval.

## Bootstrap

- popular since 90 's thanks to faster computers
- distribution-independent
- All the information about the population we have is the sample.
- Resampling produces a similar distribution to repeated sampling from the population.
- The new samples (called "resamples" or "bootstrap samples") must have the same size as the original sample.
- We must sample with replacement. Otherwise all resamples would be identical.
- Sort resamples based on the statistic (mean, BLEU,...).
- Take central $95 \%$ of resamples.


## Conclusion

Sources and further reading

- http://statslc.com/ youtube videos
- http://en.wikipedia.org/wiki/P-value etc.
- http://vassarstats.net/ can compute test statistic (JS)
- http://www.statisticsdonewrong.com


