The Bootstrap and Jackknife Methods

Session 9

Module 1 Probability & Statistical Inference

The Summer Institutes
DEPARTMENT OF BIOSTATISTICS
SCHOOL OF PUBLIC HEALTH
UNIVERSITY of WASHINGTON

Motivation

In scientific research, we're often interested in estimating some unknown parameter θ, e.g., mean weight of a certain strain of mice, heritability index, a genetic component of variation, a mutation rate, etc.

Two key questions:

1. How do we estimate θ?
2. Given an estimator for θ, how do we estimate its precision/accuracy?

Question 1 can be reasonably answered by the researcher.

Question 2 will be addressed by estimating the standard error of the estimator of θ.

Standard Error

Suppose we want to estimate a parameter θ of a distribution. e.g., the mean/median

- We select a random sample and calculate $\hat{\theta}$, our estimate of θ.
- Any function of our sample is also random.
- So our estimate, $\hat{\theta}$, is random.
- If we collect a new sample, we get a new estimate. Same for another sample, and another.

- Therefore, our estimate has a distribution, called the *sampling distribution*.

The standard deviation of that distribution is the **standard error.**

Motivation for Bootstrap Resampling

Challenges

Estimating the standard error, even for relatively simple estimators (e.g., ratios and other non-linear functions of estimators) can be quite challenging.

Solutions to most estimators are mathematically intractable or too complicated to develop, with or without advanced training in statistical inference.

However, great strides in computing in the last 30 years have made these calculations more feasible.

The bootstrap method allows us to obtain robust estimates of precision.

Limitations of the Central Limit Theorem

Estimating the precision of the sample mean

The central limit theorem gives us the standard error of $\bar{X}$:

$$\widehat{se}\left[\overline{X}\right] = \sqrt{\hat{\sigma}^2 / n}$$

where

$$\hat{\sigma}^2 = \frac{\sum_{i=1}^{n}(X_i - \overline{X})^2}{n-1}$$

However, the CLT only applies to means -- it does not extend to other estimators. The bootstrap is a more general approach that applies to medians, diversity indices, ratios…

Bootstrap Algorithm

Assume the sample dataset accurately reflects the population from which it is drawn

Generate a large number of "bootstrap" samples by resampling (with replacement) from your dataset

Resample with the same structure as used in the original sample

Calculate your estimator $\hat{\theta}$ for each of the bootstrap samples

Calculate the standard deviation of the bootstrapped estimates

Bootstrap Estimation Example: Median

What is the variance of the sample median?

=> Use the bootstrap

	Data	Median
Original sample:	{1, 5, 8, 3, 7}	5
Bootstrap 1:	{1, 7, 1, 3, 7}	3
Bootstrap 2:	{7, 3, 8, 8, 3}	7
Bootstrap 3:	{7, 3, 8, 8, 3}	7
Bootstrap 4:	{3, 5, 5, 1, 5}	5
Bootstrap 5:	{1, 1, 5, 1, 8}	1
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Bootstrap *B* (=1000)	{1, 5, 7, 7, 8}	7

Bootstrap Estimation
Example: Median

Bootstrapped estimates of the standard error for sample median (cont.)

Descriptive statistics for the sample medians from 1000 bootstrap samples

B	1000
Mean	4.964
Standard Deviation	**1.914**
Median	5
Minimum, Maximum	1, 8
25th, 75th percentile	3, 7

We estimate the standard error for the sample median as 1.914

Bootstrap Estimation Example: Relative Risk

Bootstrapped estimates of the standard error for sample relative risk

RR = P[Disease | Exposed]/P[Disease | Not exposed]

Cross-classification of Framingham Men by high systolic blood pressure and heart disease

High Systol BP	Heart Disease: No	Heart Disease: Yes
No	915	48
Yes	322	44

The sample estimate of the relative risk is:

$RR = (44/366)/(48/963) = 2.412$

Bootstrap Estimation Example cont'd

Bootstrapped estimates of the standard error for the relative risk (cont.)

Descriptive statistics for the sample relative risks*:*

B	100000
Bootstrap mean of *RR*	2.464
Bootstrap median of *RR*	2.412
Standard Deviation of *RR*	**0.507**

The bootstrap standard error for the estimated relative risk is 0.507

Bootstrap Summary

Advantages

All-purpose, computer intensive method useful for statistical inference.
Bootstrap estimates of precision do not require knowledge of the theoretical form of an estimator's standard error, no matter how complicated it is.

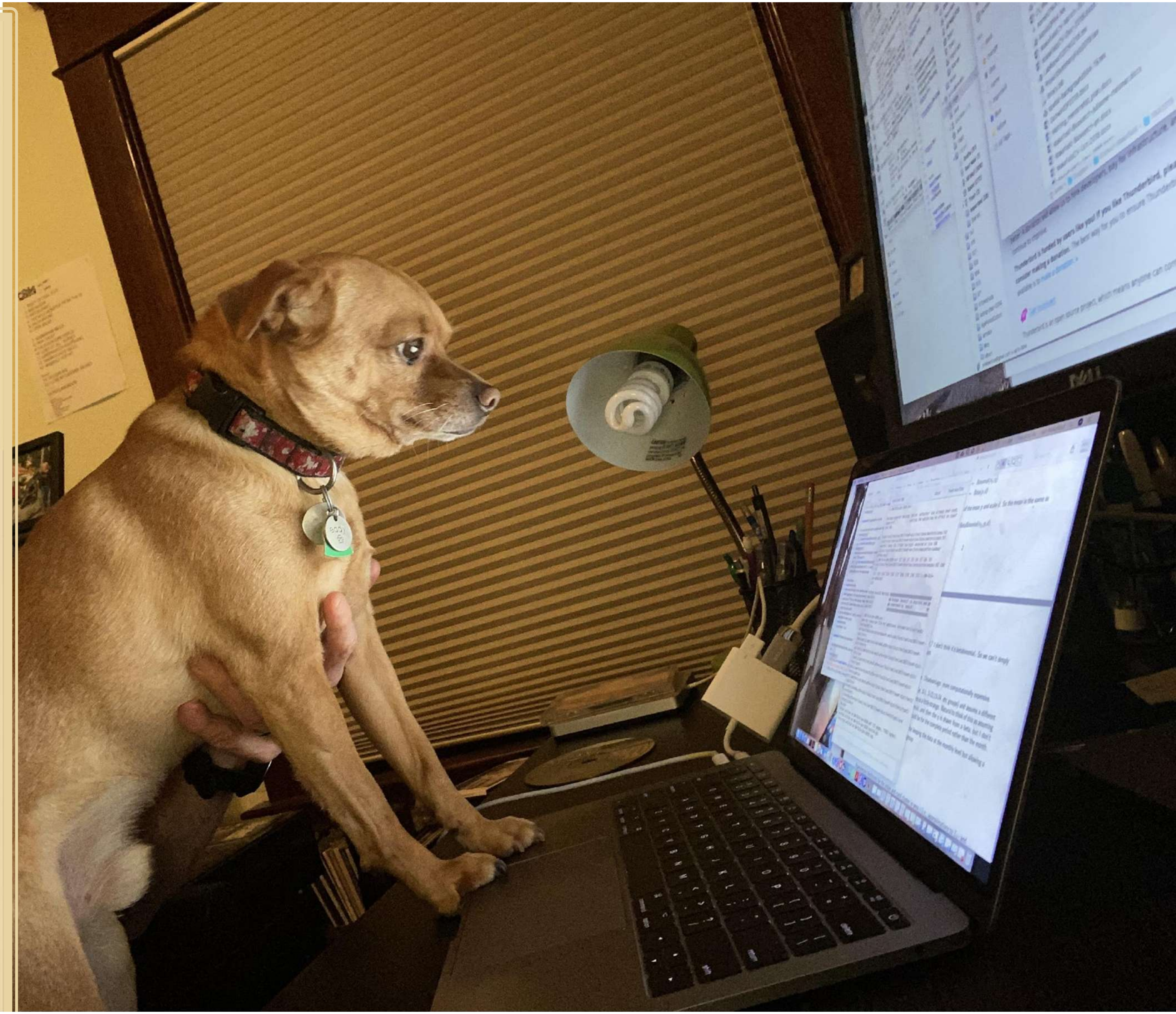
Disadvantages

- Typically not useful for correlated (dependent) data.
- Missing data, censoring, data with outliers are also problematic
- Often used incorrectly

There are many different types of bootstraps: we have only discussed one

Paws-

break time then work on Q1



Jackknife (leave-one-out) estimation

- invented by Quenouille (1949)
- an alternative resampling method to the bootstrap.
- based upon sequentially deleting one observation from the dataset, recomputing the estimator, $\hat{\theta}_{(i)}$, *n* times. So, exactly *n* jackknife estimates for a sample of size *n*.
- like the bootstrap, the jackknife method provides a relatively easy way to estimate the precision of an estimator, θ.
- generally less computationally intensive than the bootstrap

Jackknife Algorithm

Jackknifing

- For a dataset with *n* observations, compute *n* estimates by sequentally omitting each observation from the dataset and estimating $\hat{\theta}$ on the remaining *n* – 1 observations.
- Using the *n* jackknife estimates, $\hat{\theta}_{(1)}, \hat{\theta}_{(2)}, \ldots, \hat{\theta}_{(n)}$,

we estimate the standard error of the estimator as:

$$\widehat{se}_{jack} = \sqrt{\frac{n-1}{n}\sum_{i=1}^{n}(\hat{\theta}_{(i)} - \bar{\hat{\theta}}_{(.)})^2}$$

- Unlike the bootstrap, the jackknife standard error estimate will not change for a given sample

Jackknife Summary

Advantages

- Useful method for estimating and compensating for bias in an estimator.
- Like the bootstrap, the methodology does not require knowledge of the theoretical form of an estimator's standard error.
- Generally less computationally intensive compared to the bootstrap.

Disadvantages

- The jackknife is more conservative than the bootstrap → its estimated standard error tends to be slightly larger.
- Performs poorly when the estimator is not sufficiently smooth, e.g., the median.