The Bootstrap and Jackknife

Summer 2017 Summer Institutes 249

Bootstrap & Jackknife Motivation

In scientific research

- Interest often focuses upon the estimation of some unknown parameter, θ . The parameter θ can represent for example, mean weight of a certain strain of mice, heritability index, a genetic component of variation, a mutation rate, etc.
- Two key questions need to be addressed:
 - 1. How do we estimate θ ?
 - 2. Given an estimator for θ , how do we estimate its precision/accuracy?
- We assume Question 1 can be reasonably well specified by the researcher
- Question 2, for our purposes, will be addressed via the estimation of the estimator's standard error

st de erroi = estimate
of std dev. of

Summer 2017

Summer Institutes

an estimates

What is a standard error?

Suppose we want to estimate a parameter theta (eg. the mean/median/squared-log-mode) of a distribution

- Our sample is random, so...
- Any function of our sample is random, so...
- Our estimate, theta-hat, is random! So...
- If we collected a new sample, we'd get a new estimate. Same for another sample, and another... So
- Our estimate has a distribution! It's called a sampling distribution!

The standard deviation of that distribution is the standard error

Summer 2017 Summer Institutes 251

Bootstrap Motivation

Challenges

- Answering Question 2, 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, particularly in the last 25 years, have made computational intensive calculations feasible.
- We will investigate how the bootstrap allows us to obtain robust estimates of precision for our estimator, θ , with a simple example...

Bootstrap Estimation

Estimating the precision of the sample mean

• A dataset of n observation provides more than an estimate of the population mean (denoted here as X), where

$$\bullet \quad \overline{X} = \frac{1}{n} \sum_{i=1}^{n} X_i$$

• It gives an estimate of the precision of \overline{X} , namely

gives an estimate of the precision of
$$\widehat{se}\left[\overline{X}\right] = \sqrt{\widehat{\sigma}^2 / n}$$
where
$$\widehat{\sigma}^2 = \frac{\sum_{i=1}^{n} (X_i - \overline{X})}{n-1}$$

is an estimate of the population variance.

- This standard error estimate is that it does not extend to estimators other than \overline{X}
- •Standard deviation: square root of variance; standard error: estimate of standard deviation

Bootstrap Estimation

Estimating the precision of the sample mean

- From the formulas on the previous page, we can obtain an estimate of precision for \overline{X} by estimating the population variance and "plugging" it into the formula for the standard error estimate.
- **Question**: What IF you did not know the formula for the standard error of the sample mean, BUT you had access to modern PC. How might you obtain an estimate of precision?
- **Answer**: The bootstrap!

Bootstrap Algorithm

Bootstrapping

- Assuming the sample accurately reflects the population from which it is drawn \overline{X}
- Generate a large number of "bootstrap" samples by resampling (with replacement) from the dataset
- Resample with the same structure (dependence, sample sizes) as used in the original sample
- Compute your estimator, θ , (here, $\theta = X$), for each of the bootstrap samples
- Compute the "standard deviation" from the statistics calculated above.

Bootstrap Algorithm

Bootstrap sample

Bootstrap estimates

1:
$$(X_1^{(1)}, X_2^{(1)}, \dots, X_n^{(1)}) \rightarrow \hat{\theta}(X_1^{(1)}, X_2^{(1)}, \dots, X_n^{(1)}) = \overline{X}_{(2)}^{(1)}$$

2:
$$(X_1^{(2)}, X_2^{(2)}, \dots, X_n^{(2)}) \rightarrow \hat{\theta}(X_1^{(2)}, X_2^{(2)}, \dots, X_n^{(2)}) = \overline{X}^{(2)}$$

B:
$$(X_1^{(B)}, X_2^{(B)}, \dots, X_n^{(B)}) \rightarrow \hat{\theta}(X_1^{(B)}, X_2^{(B)}, \dots, X_n^{(B)}) = \overline{X}^{(B)}$$

Compute $\hat{\sigma}_b^2$, where $\hat{\sigma}_b^2 = \frac{\sum_{j=1}^B (\overline{X}^{(j)} - \overline{X}^{(j)})^2}{B-1}$, and

$$\overline{X}^{(.)} = \frac{1}{B} \sum_{j=1}^{B} \overline{X}^{(j)}$$
.

The bootstrap standard error is $\widehat{se_{boot}}\left[\overline{X}\right] = \sqrt{\widehat{\sigma}_b^2}$.

For other estimators, simply replace \overline{X} with the θ of your choice.

What is the variance of the sample median? No idea! => Use the bootstrap!

Bootstrapped estimates of the standard error for sample median

<u>Data</u> Median	
Original sample: {1, 5, 8, 3, 7}	5
Bootstrap 1: {1, 7, 1, 3, 7} Bootstrap 2: {7, 3, 8, 8, 3} Bootstrap 3: {7, 3, 8, 8, 3} Bootstrap 4: {3, 5, 5, 1, 5} Bootstrap 5: {1, 1, 5, 1, 8}	3 7 7 7 5 1
etc. Bootstrap B (=1000)	
1 /	
	B

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

• Descriptive statistics for the sample medians from 1000 bootstrap samples

В	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
- A 95% asymptotic (with n=5?) confidence interval (using the 0.975 quantile of the standard normal distribution) is

$$5 + / -1.96(1.914) = (1.25, 8.75)$$

Bootstrapped estimates of the standard error for sample relative risk

r = P[D|Exposed]/P[D|Not exposed]

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

Heart Disease

High Systol BP	No	Yes
No	915	48
Yes	322	44

The sample estimate of the relative risk is

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

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

• Descriptive statistics for the sample relative risks

100000
2.464
2.412
0.507

- The bootstrap standard error for the estimated relative risk is 0.507
- A 95% bootstrap confidence interval is

$$2.412 + -1.96(0.507) = (1.42, 3.41)$$

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

Jackknife

Jackknife Estimation

- The jackknife (or leave one out) method, invented by Quenouille (1949), is an alternative resampling method to the bootstrap.
- The method is based upon sequentially deleting one observation from the dataset, recomputing the estimator, here, $\hat{\theta}_{(i)}$, n times. That is, there are exactly n jackknife estimates obtained in a sample of size n.
- Like the bootstrap, the jackknife method provides a relatively easy way to estimate the precision of an estimator, θ .
- The jackknife is 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, $\hat{\theta}_{(1)}, \hat{\theta}_{(2)}, \dots, \hat{\theta}_{(n)}$, we estimate the standard error of the estimator as

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

• 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.
- Is generally less computationally intensive compared to the bootstrap method.

Disadvantages

- The jackknife method is more conservative than the bootstrap method, that is, its estimated standard error tends to be slightly larger.
- Performs poorly when the estimator is not sufficiently smooth, i.e., a non-smooth statistic for which the jackknife performs poorly is the median.