#### Module 7: Simulations

Siyue Yang

06/03/2022

#### Simulation study

- Simulation: A numerical techniques for conducting experiments on the computer
- Monte Carlo simulation: Computer experiment involving random sampling from probability distributions

## Why simulation?

To establish/validate the properties of statistical methods

- Exact analytical derivations of properties are rarely possible
- Large sample approximations to properties are often possible, but need to evaluate their relevance to (finite) sample sizes likely to be encountered in practice

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Moreover, analytical results may require **assumptions** (e.g., normality)

- But what happens when these assumptions are violated?
- ► Analytical results, even large sample ones, may not be possible

#### Considerations for simulation

- Is an estimator biased in finite samples? Is it still consistent under departures from assumptions? What is its sampling variance?
- How does it compare to competing estimators on the basis of bias, precision, etc.?

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- Is an estimator biased in finite samples? Is it still consistent under departures from assumptions? What is its sampling variance?
- ▶ How does it compare to competing estimators on the basis of bias, precision, etc.?
- Does a procedure for constructing a confidence interval for a parameter achieve the advertised nominal level of coverage?
- Does a hypothesis testing procedure attain the advertised level or size?
- ▶ If it does, what **power** is possible against different alternatives to the null hypothesis? Do different test procedures deliver different power?

#### Monte Carlo simulation

- Generate S independent data sets under the conditions of interest
- Compute the numerical value of the estimator/test statistic T (data) for each data set  $\Rightarrow T_1, \ldots, T_S$
- ► If S is large enough, summary statistics across T<sub>1</sub>,..., T<sub>S</sub> should be good approximations to the true sampling properties of the estimator/test statistic under the conditions of interest

# Simulations for properties of estimators

Example: Compare 3 estimators for the **mean**  $\mu$  of a distribution based on i.i.d. draws  $Y_1, \ldots, Y_n$ 

- ightharpoonup Sample mean  $T^{(1)}$
- $\triangleright$  Sample 20% trimmed mean  $T^{(2)}$
- ▶ Sample median  $T^{(3)}$

# Simulations for properties of estimators (cont'd)

Simulation procedure: For a particular choice of  $\mu$ ,  $\emph{n}$ , and true underlying distribution

- ▶ Generate independent draws  $Y_1, ..., Y_n$  from the distribution
- ► Compute  $T^{(1)}$ ,  $T^{(2)}$ ,  $T^{(3)}$
- ▶ Repeat *S* times  $T_1^{(1)}, \dots, T_S^{(1)}; \quad T_1^{(2)}, \dots, T_S^{(2)}; \quad T_1^{(3)}, \dots, T_S^{(3)}$
- ightharpoonup Compute for k = 1, 2, 3

$$\widehat{\text{mean}} = S^{-1} \sum_{s=1}^{S} T_s^{(k)} = \overline{T}^{(k)}, \ \widehat{\text{bias}} = \overline{T}^{(k)} - \mu$$

$$\widehat{\text{SD}} = \sqrt{(S-1)^{-1} \sum_{s=1}^{S} \left(T_s^{(k)} - \overline{T}^{(k)}\right)^2}$$

$$\widehat{\text{MSE}} = S^{-1} \sum_{s=1}^{S} \left(T_s^{(k)} - \mu\right)^2 \approx \widehat{\text{SD}}^2 + \widehat{\text{bias}}^2$$

## Simulations for properties of estimators (cont'd)

Another important property we care about is the **relative efficiency** (RE).

▶ If the estimators are unbiased,

$$RE = \frac{\operatorname{var}\left(T^{(1)}\right)}{\operatorname{var}\left(T^{(2)}\right)}$$

▶ If the estimators are biased,

$$RE = \frac{\mathsf{MSE}\left(T^{(1)}\right)}{\mathsf{MSE}\left(T^{(2)}\right)}$$

In either case RE < 1 means estimator 1 is preferred (estimator 2 is inefficient relative to estimator 1 in this sense)

## Set up parameters

```
set.seed(3)
S <- 1000
n <- 15
mu <- 1
sigma <- sqrt(5/3)

trimmean <- function(Y) mean(Y, 0.2)</pre>
```

#### Generate data

Note: for this very simple data generation, we can get the data in one step, no looping. In more complex statistical models, looping is often required.

```
generate.normal <- function(S, n, mu, sigma){
  dat <- matrix(rnorm(n*S, mu, sigma), ncol=n, byrow=T)
  out <- list(dat=dat)
  return(out)
}</pre>
```

```
out <- generate.normal(S, n, mu, sigma)
out_mean <- apply(out$dat, 1, mean)
out_trimmean <- apply(out$dat, 1, trimmean)
out_median <- apply(out$dat, 1, median)</pre>
```

#### View the simulated data

```
## mean trim median
## 1 0.753935 0.7131731 1.0388898
## 2 0.643902 0.4580396 0.3745711
## 3 1.555288 1.6710299 1.9394763
## 4 0.517147 0.4826527 0.4118927
## 5 1.360281 1.4620501 1.3451583
## 6 1.359185 1.3955097 1.4949135
```

#### View the estimator properties

```
simsum <- function(dat, trueval){
   S <- nrow(dat)
   MCmean <- apply(dat,2,mean)</pre>
   MCbias <- MCmean-trueval
   MCrelbias <- MCbias/trueval
   MCstddev <- sqrt(apply(dat,2,var))</pre>
   MCMSE <- apply((dat-trueval)^2,2,mean)</pre>
# MCMSE <- MCbias^2 + MCstddev^2 # alternative lazy calculation
   MCRE <- MCMSE[1]/MCMSE
   sumdat <- rbind(rep(trueval,3), S, MCmean, MCbias,</pre>
                    MCrelbias, MCstddev, MCMSE, MCRE)
   names <- c("true value", "# sims", "MC mean", "MC bias", "MC relative bias",
              "MC standard deviation", "MC MSE", "MC relative efficiency")
   ests <- c("Sample mean", "Trimmed mean", "Median")
   dimnames(sumdat) <- list(names,ests)</pre>
   round(sumdat.5)
```

## View the estimator properties (cont'd)

```
results <- simsum(summary.sim, mu)
results
```

```
Sample mean Trimmed mean
                                                     Median
##
## true value
                             1.00000
                                         1.00000
                                                    1.00000
## # sims
                          1000.00000
                                       1000.00000 1000.00000
## MC mean
                             0.98515
                                         0.98690
                                                    0.99173
## MC bias
                            -0.01485
                                      -0.01310
                                                  -0.00827
## MC relative bias
                            -0.01485
                                        -0.01310
                                                  -0.00827
## MC standard deviation
                            0.33088
                                         0.34800
                                                  0.39763
## MC MSE
                            0.10959
                                         0.12116
                                                  0.15802
## MC relative efficiency
                           1.00000
                                         0.90456
                                                  0.69356
```

# Performance of estimates of uncertainty

How well do estimated standard errors represent the true sampling variation?

- Compare the average of the estimated standard errors to MC standard deviation.
- For sample mean  $\bar{Y}$ ,  $SE(\bar{Y}) = \frac{s}{\sqrt{n}}, \quad s^2 = (n-1)^{-1} \sum_{j=1}^n \left(Y_j \bar{Y}\right)^2$

```
results["MC standard deviation", "Sample mean"]
```

```
mean_se <- sqrt(apply(out$dat, 1, var)/n)
ave_mean_se <- mean(mean_se)
round(ave_mean_se, 3)</pre>
```

```
## [1] 0.329
```

## [1] 0.33088

#### Confidence interval

Based on the sample mean,

$$\left[\bar{Y}-t_{1-\alpha/2,n-1}\frac{s}{\sqrt{n}},\bar{Y}+t_{1-\alpha/2,n-1}\frac{s}{\sqrt{n}}\right]$$

Does the interval achieve the nominal level of coverage  $1-\alpha$  ?

```
## [1] 0.949
```

# Simulations for properties of hypothesis testing

Example: Size and power of the usual *t*-test for the mean

$$H_0: \mu = \mu_0$$
 vs.  $H_1: \mu \neq \mu_0$ 

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To evaluate whether size/level of test achieves advertised  $\alpha$ 

- ightharpoonup Approximates the true probability of rejecting  $H_0$  when it is true
- Generate data under  $H_0: \mu = \mu_0$
- lacktriangle Calculate proportion of rejections of  $H_0$ , should pprox lpha

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- ightharpoonup Approximates the true probability of rejecting  $H_0$  when it is true
- Generate data under  $H_0: \mu = \mu_0$
- ▶ Calculate proportion of rejections of  $H_0$ , should  $\approx \alpha$

#### To evaluate the power

- Approximates the true probability of rejecting  $H_0$  when the alternative is true (power)
- Generate data under some alternative  $H_1: \mu \neq \mu_0$
- Calculate proportion of rejections of H<sub>0</sub>

## Parameters set up

```
set.seed(3)
S <- 1000
n <- 15
sigma <- sqrt(5/3)</pre>
```

## Size/level of test

```
m_{11}O < -1
m11 < -1
out <- generate.normal(S, n, mu, sigma)
samp_mean <- apply(out$dat, 1, mean)</pre>
mean_se <- sqrt(apply(out$dat, 1, var)/n)</pre>
ttests <- (samp_mean - mu0)/mean_se
t05 \leftarrow qt(0.975, n-1)
sum(abs(ttests) > t05)/S
## [1] 0.051
```

#### Power of test

```
m_{11}O < -1
m_{11} < -1.75
out <- generate.normal(S, n, mu, sigma)
samp_mean <- apply(out$dat, 1, mean)</pre>
mean_se <- sqrt(apply(out$dat, 1, var)/n)</pre>
ttests <- (samp_mean - mu0)/mean_se
t05 \leftarrow qt(0.975, n-1)
sum(abs(ttests) > t05)/S
## [1] 0.512
```

#### Simulation studies principles

How well do the Monte Carlo quantities approximate properties of the true sampling distribution of the estimators/test statistics?

- Principle 1: Carefully choose S
- ▶ Principle 2: Save everything
- Principle 3: Keep S small at first
- Principle 4: Set a different seed for each run and keep records
- Principle 5: Document your code

## Principle 1: Carefully choose *S*

Is S=1000 large enough to get a feel for the true sampling properties? How "believable" are the results?

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Estimator for  $\theta$  (true value  $\theta_0$ ) e.g. mean of sampling distribution

$$\sqrt{\operatorname{var}\left(\overline{T} - \theta_0\right)} = \sqrt{\operatorname{var}(\overline{T})} = \sqrt{\operatorname{var}\left(S\sum_{s=1}^{S} T_s\right)} = \frac{\operatorname{SD}\left(T_s\right)}{\sqrt{S}} = d$$

where d is the acceptable error

$$\Rightarrow S = \frac{\{\mathrm{SD}(T_s)\}^2}{d^2}$$

# Principle 1: Carefully choose *S* (cont'd)

Coverage probabilities, size, power e.g. for a hypothesis testing

$$Z = \# \text{ rejections } \sim \text{binomial}(S, p) \Rightarrow \sqrt{\text{var}\left(\frac{Z}{S}\right)} = \sqrt{\frac{p(1-p)}{S}}$$

- ▶ Worst case is at  $p = 1/2 \Rightarrow 1/\sqrt{4S}$
- ▶ d acceptable error  $\Rightarrow$   $S=1/\left(4d^2\right)$ ; e.g., d=0.01 yields S=2500
- For coverage, size, p = 0.05

## Principle 2: Save everything

- ► Save individual estimates in a file then analyze
- ▶ Useful when simulation takes a long time to run

```
# Save txt file.
file name <- paste0("ssl binary",
                     " lab", n,
                     " beta", b,
                     " prev", p,
                     " setting", 1,
                     "_reps", n_sim,
                     ".txt")
write.table(result, file = out_file,
            sep = "\t", row.names = FALSE)
```

```
# Save .Rdata file save(result)
```

## Principle 3: Keep *S* small at first

Test and refine the code until everything is working correctly before trying out final production runs

- ▶ If *S* is large, say, 1000
- ▶ Try S = 20 first
- As it takes less time to run and easy for debugging
- Keep track of how long it will take

# Principle 4: Set a different seed for each run and keep records

- Ensure simulation runs are independent
- Runs may be replicated if necessary

e.g.

```
data_generation <- function(S) {</pre>
  for (i in c(1:S)) {
    set.seed(1234+i)
    X <- ...
   Υ <- ...
  data.frame(X = X, Y = Y)
```

#### Contributions

This module closely follows Marie Davidian's STA810A Preparation for Statistical Research handout of simulation studies in statistics. See links here.