

# PLSC 503 – Spring 2024

## Bootstrapping and Missing Data

March 11, 2024

**The population is to the sample as the sample is to the bootstrap sample.**

# Practical (Nonparametric) Bootstrapping

## The General Idea:

- Draw one bootstrap sample of size  $N$  **with replacement** from the original data,
- Estimate the parameter(s)  $\tilde{\theta}_{k \times 1}$ ,
- Repeat steps 1 and 2  $R$  times, to get  $\tilde{\theta}_r$ ,  $r \in \{1, 2, \dots, R\}$ , comprising elements  $\tilde{\theta}_{rk}$ ,
- Examine the empirical characteristics of the resulting distribution(s) of  $\tilde{\theta}_{rk}$ .

# Why Bootstrap?

- **It's intuitive.**
- **It's simple.**
- **It's robust.**

# Bootstrapping: “By Hand”

```
N<-10 # small sample!
reps<-1001

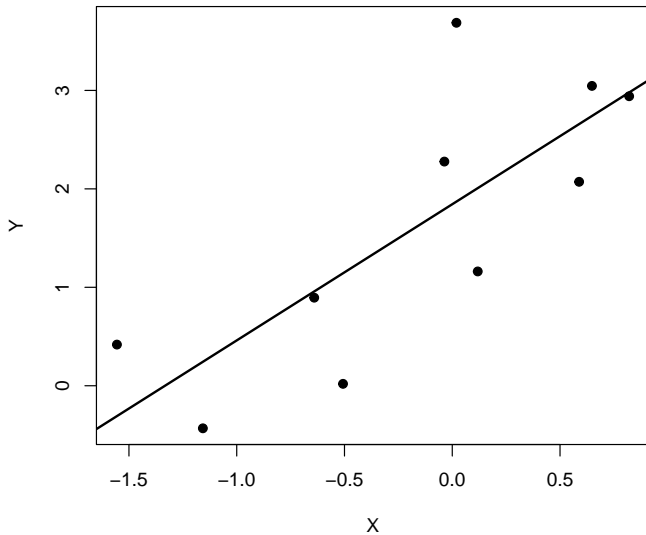
set.seed(1337)
X<-rnorm(N)
Y<-2+2*X+rnorm(N)
data<-data.frame(Y,X)
fitOLS<-lm(Y~X)
CI<-confint(fitOLS)

B0<-numeric(reps)
B1<-numeric(reps)

for (i in 1:reps) {
  temp<-data[sample(1:N,N,replace=TRUE),]
  temp.lm<-lm(Y~X,data=temp)
  B0[i]<-temp.lm$coefficients[1]
  B1[i]<-temp.lm$coefficients[2]
}

ByHandB0<-median(B0)
ByHandB1<-median(B1)
ByHandCI.B0<-quantile(B0,probs=c(0.025,0.975)) # <-- 95% c.i.s
ByHandCI.B1<-quantile(B1,probs=c(0.025,0.975))
```

## Normal Residuals...



# Bootstrapping Via boot

```
library(boot)

Bs<-function(formula, data, indices) { # <- regression function
  dat <- data[indices,]
  fit <- lm(formula, data=dat)
  return(coef(fit))
}

Boot.fit<-boot(data=data, statistic=Bs,
               R=reps, formula=Y~X)

BootB0<-median(Boot.fit$t[,1])
BootB1<-median(Boot.fit$t[,2])
BootCI.B0<-boot.ci(Boot.fit,type="basic",index=1)
BootCI.B1<-boot.ci(Boot.fit,type="basic",index=2)
```

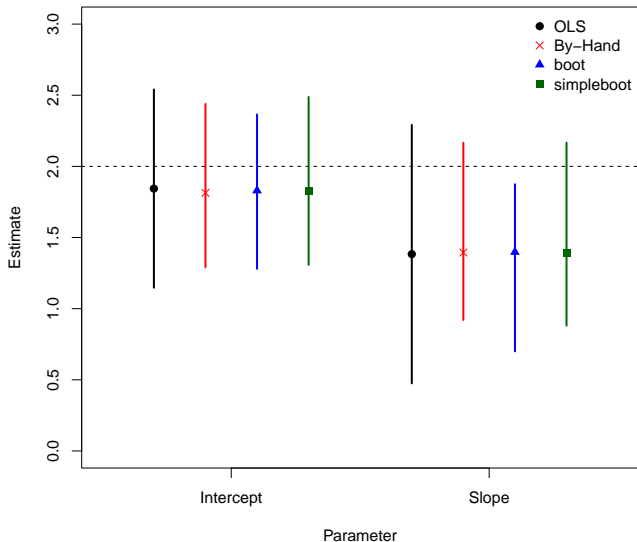
# Bootstrapping Via simpleboot

```
library(simpleboot)

Simple<-lm.boot(fitOLS, reps)
SimpleB0<-perc(Simple, .50)[1]
SimpleB1<-perc(Simple, .50)[2]
Simple.CIs<-perc(Simple, p=c(0.025, 0.975))
```

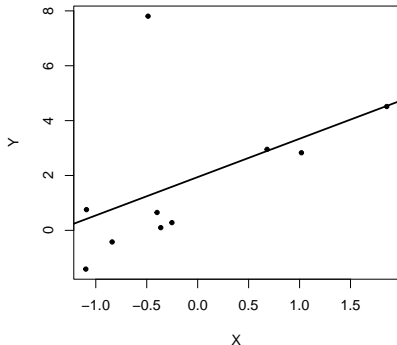


# Bootstrapping Results

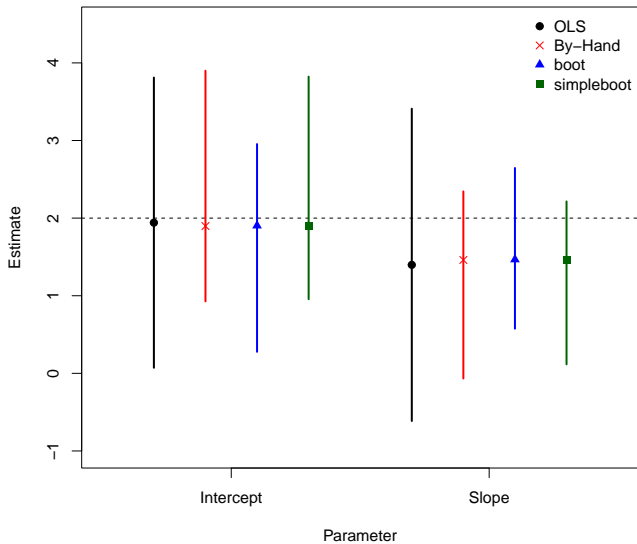


# Bootstrapping: Skewed Residuals

```
N<-10  
reps<-1001  
  
set.seed(1337)  
X<-rnorm(N)  
ustar<-rgamma(N,shape=0.2,scale=1)*6 # <- skewed u.s  
Y<-2+2*X+(ustar-mean(ustar))  
data<-data.frame(Y,X)  
fitOLS<-lm(Y~X)  
CI<-confint(fitOLS)
```



# Skewed Residuals: Results



# When Should I Bootstrap?

## A few canonical applications:

- When  $N$  is small, and the estimator is consistent (but not unbiased / efficient)
- When the estimand(s) is/are complex
- When the distribution *of the estimand(s)* is unknown
- As a robustness check on your findings when data are complex

## R things:

- A [simple introduction](#) at StatMethods
- [Bootstrap in R](#) (at DataCamp)
- Packages: [boot](#), [bootstrap](#), [simpleboot](#), [car::Boot](#), [broom](#) (tidy), many more

## Other Resources:

- Efron's [original \(1979\) paper](#)
- [Chernick and Labudde \(2011\)](#) (a solid R-based bootstrapping book)
- A good little [online intro](#), by James Scott
- Many other books, etc.

# Missing Data

## Why are data missing?

- The observation itself does not exist
- Data don't exist for that observation
- Data exist, but are *impossible* to measure
- Data exist, but were not measured

# Missing Data, Part II: Flavors

Notation:

$$\mathbf{x}_i \in \{\mathbf{w}_i, \mathbf{z}_i\}$$

$\mathbf{w}_i$  have some missing values,  
 $\mathbf{z}_i$  are “complete”

$$R_{ik} = \begin{cases} 1 & \text{if } W_{ik} \text{ is missing,} \\ 0 & \text{otherwise.} \end{cases}$$

$$\pi_{ik} = \Pr(R_{ik} = 1)$$



# Missing Data, Part II: Rubin's Flavors

Missing completely at random ("MCAR"):

$$\mathbf{R} \perp \{\mathbf{Z}, \mathbf{W}\}$$

Missing at random ("MAR"):

$$\mathbf{R} \perp \mathbf{W} | \mathbf{Z}$$

Anything else is "informatively" (or "non-ignorably," or sometimes "MNAR") missing.

# MCAR vs. MAR vs. MNAR, Explained

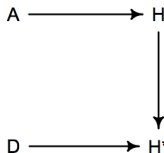
H: Homework

H\*: Homework with missing values

A: Attribute of student

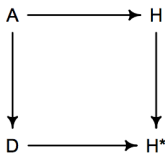
D: Dog (missingness mechanism)

**DOG EATS  
ANY  
HOMEWORK**



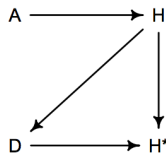
**MISSING COMPLETELY  
AT RANDOM**

**DOG EATS  
STUDENTS'  
HOMEWORK**



**MISSING  
AT RANDOM**

**DOG EATS  
BAD  
HOMEWORK**



**MISSING NOT  
AT RANDOM**

(Source)

# Missing Data: Consequences

In general:

- MCAR:
  - Missing data are a fully random sample of all the data
  - $\rightarrow$  Missingness does not bias  $\hat{\theta}$ , *but*
  - There is some loss of information (and therefore efficiency)
- MAR
  - Missing data are a nonrandom sample of all the data
  - Ignoring missingness can lead to bias in  $\hat{\theta}$ , *but*
  - Conditioning on the variable(s) that drive the missingness can eliminate the bias
- Informative Missingness / MNAR
  - Missing data are a nonrandom sample of all the data
  - Ignoring missingness can lead to bias in  $\hat{\theta}$
  - In general, conditioning cannot eliminate the bias

## Example, Simulated

```
> set.seed(7222009)
> Npop <- 1000
> X<-runif(Npop,0,10)    # NOTE: X, Z are correlated a bit...
> Z<-(0.3*X)+(0.7*runif(Npop,0,10))
> Y<-0+(2*X)+(2*Z)+rnorm(Npop,mean=0,sd=4)
> DF<-data.frame(X=X,Z=Z,Y=Y)
> fit.pop<-lm(Y~X+Z,DF)
> summary(fit.pop)
```

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	0.4051	0.3260	1.24	0.21
X	1.9553	0.0466	41.97	<2e-16 ***
Z	1.9812	0.0617	32.09	<2e-16 ***

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 3.98 on 997 degrees of freedom

Multiple R-squared: 0.823, Adjusted R-squared: 0.823

F-statistic: 2.32e+03 on 2 and 997 DF, p-value: <2e-16

# Simulated MCAR

```
> pmis<-0.50
> DF$Ymcar<-rbinom(Npop,1,pmis)
> DF$Ymcar<-ifelse(DF$Ymcar==1,NA,DF$Y)
>
> # Regression w/listwise deletion:
>
> fit.s<-lm(Ymcar~X+Z,DF) # <-- looks fine
> summary(fit.s)
```

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	0.4442	0.4653	0.95	0.34
X	1.9661	0.0658	29.87	<2e-16 ***
Z	1.9763	0.0862	22.92	<2e-16 ***

---

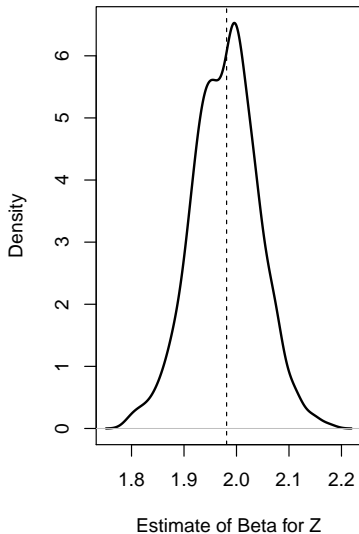
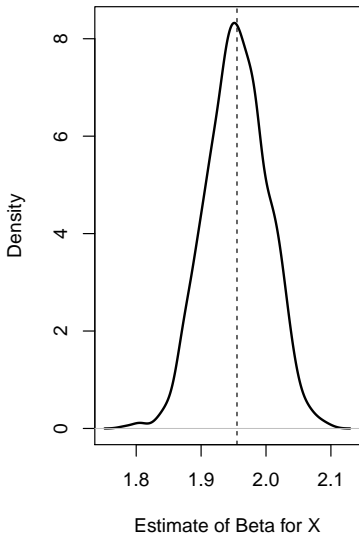
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 4 on 507 degrees of freedom  
(490 observations deleted due to missingness)

Multiple R-squared: 0.822, Adjusted R-squared: 0.821

F-statistic: 1.17e+03 on 2 and 507 DF, p-value: <2e-16

## Do That A Bunch Of Times...



```
> set.seed(7222009)
> DF$Ymar<-rbinom(Npop,1,(DF$Z/10))
> DF$Ymar<-ifelse(DF$Ymar==1,NA,DF$Y)
>
> summary(lm(Ymar~X,DF))
```

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	3.6600	0.3610	10.1	<2e-16 ***
X	2.9923	0.0648	46.2	<2e-16 ***

---

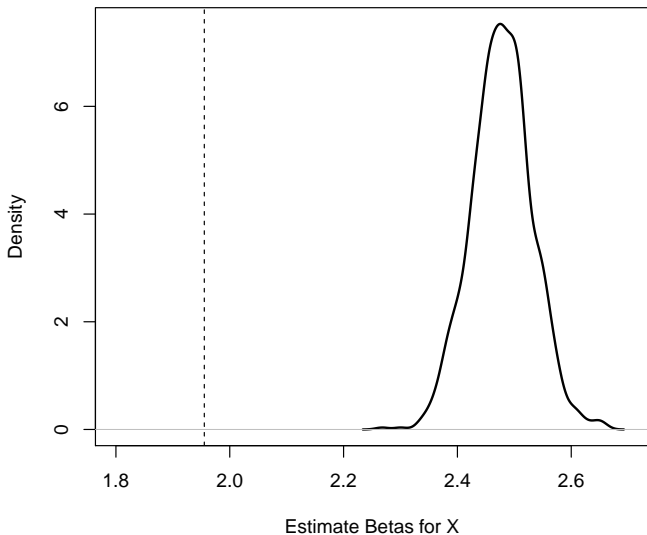
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 4.75 on 547 degrees of freedom  
(451 observations deleted due to missingness)

Multiple R-squared: 0.796, Adjusted R-squared: 0.795

F-statistic: 2.13e+03 on 1 and 547 DF, p-value: <2e-16

# Do That A Bunch Of Times...





## More MAR: Add Z...

```
> summary(lm(Ymar~X+Z,DF))
```

Call:

```
lm(formula = Ymar ~ X + Z, data = DF)
```

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	0.2529	0.4367	0.58	0.56
X	2.0200	0.0663	30.49	<2e-16 ***
Z	1.9499	0.0979	19.91	<2e-16 ***

---

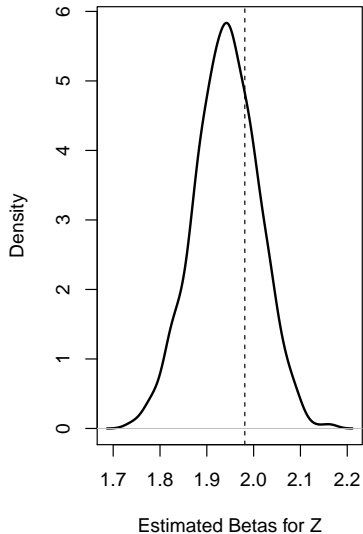
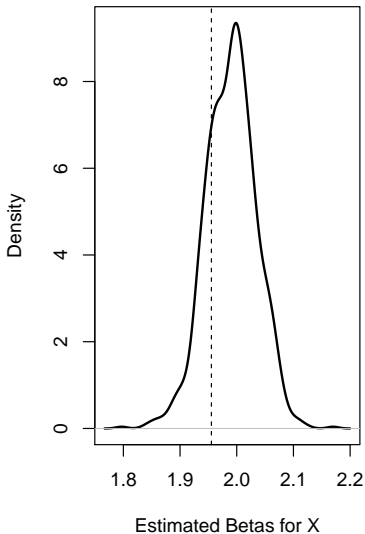
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 4.02 on 499 degrees of freedom  
(498 observations deleted due to missingness)

Multiple R-squared: 0.801, Adjusted R-squared: 0.8

F-statistic: 1e+03 on 2 and 499 DF, p-value: <2e-16

## Do That A Bunch Of Times...



# Informative Missingness / "MNAR"

```
> set.seed(7222009)
> DF$Yim<-rbinom(Npop,1,rescale(DF$Z-(4*DF$Y)))
> DF$Yim<-ifelse(DF$Yim==1,NA,DF$Y)
>
> summary(lm(Yim~X+Z,DF))
```

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	2.0518	0.5463	3.76	0.00019 ***
X	1.8420	0.0671	27.45	< 2e-16 ***
Z	1.9171	0.0859	22.32	< 2e-16 ***

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

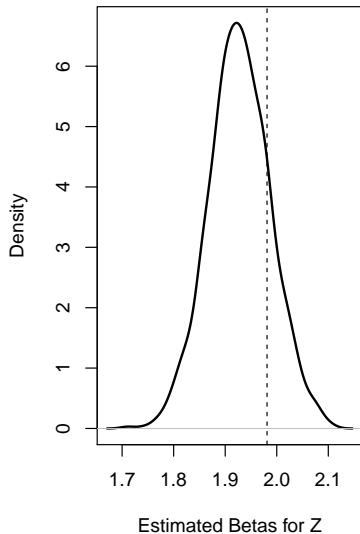
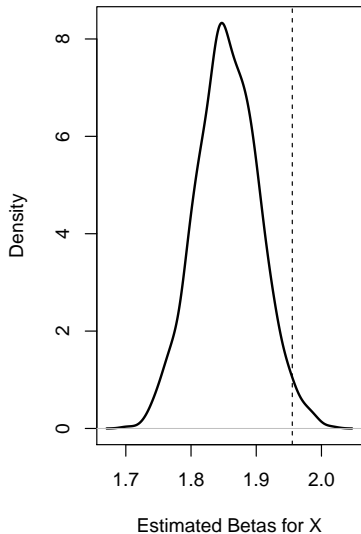
Residual standard error: 3.85 on 465 degrees of freedom

(532 observations deleted due to missingness)

Multiple R-squared: 0.797, Adjusted R-squared: 0.796

F-statistic: 911 on 2 and 465 DF, p-value: <2e-16

## Do That A Bunch Of Times...



# A Real-Data Examples: 2020 ANES

Model is:

$$\begin{aligned}\text{Biden Thermometer}_i &= \beta_0 + \beta_1 \text{R's Conservatism}_i + \\ &= \beta_2 \text{R Labor Union}_i + \beta_3 \text{Female}_i + \\ &= \beta_4 \text{Latino}_i + \beta_5 \text{Age} / 10_i + \\ &= \beta_6 \text{Education}_i + u_i\end{aligned}$$

Data: ANES 2016-2020 Panel data, 2020 pre-election survey ( $N = 2839$ ).

Three models:

- All data ( $N = 2291$ )
- 67% MCAR (via simulation) ( $N = 709$ )
- (MNAR) Data *only* on individuals who stated that they “strongly approved” of how then-President Trump was doing his job ( $N = 743$ )

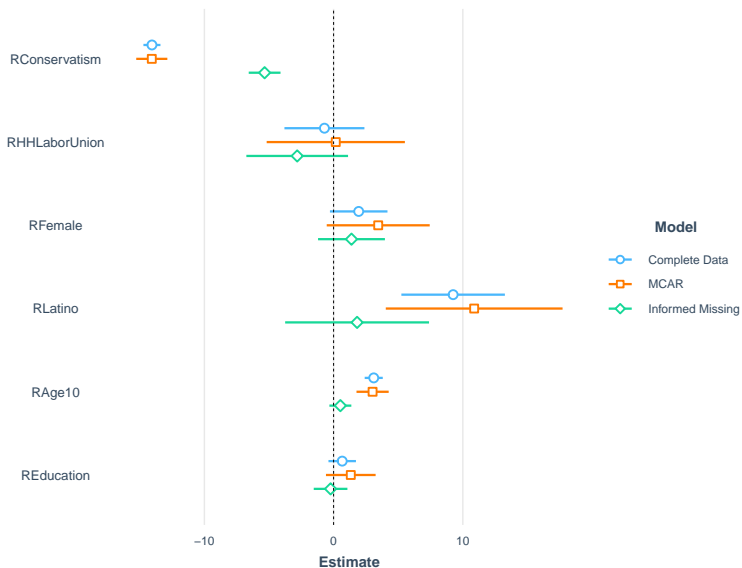
# Biden Thermometer Models

	<i>Dependent variable:</i>		
	Biden Thermometer Score		
	Complete	2/3 MCAR data	MNAR (Trump Supporters)
R's Conservatism	−14.060*** (0.336)	−14.070*** (0.613)	−5.340*** (0.627)
R in Labor Union	−0.710 (1.578)	0.168 (2.723)	−2.817 (2.004)
R is Female	1.943* (1.135)	3.454* (2.029)	1.384 (1.317)
R is Latino	9.251*** (2.042)	10.880*** (3.483)	1.818 (2.835)
R's Age(/10)	3.106*** (0.357)	3.018*** (0.634)	0.524 (0.432)
R's Education	0.666 (0.542)	1.330 (0.978)	−0.234 (0.663)
Constant	86.870*** (3.306)	82.890*** (5.915)	40.440*** (4.750)
Observations	1,942	583	621
R <sup>2</sup>	0.497	0.511	0.114
Adjusted R <sup>2</sup>	0.495	0.506	0.105
Residual Std. Error	24.770 (df = 1935)	24.240 (df = 576)	16.350 (df = 614)
F Statistic	318.300*** (df = 6; 1935)	100.500*** (df = 6; 576)	13.150*** (df = 6; 614)

Note:

\* p<0.1; \*\* p<0.05; \*\*\* p<0.01

# Biden Thermometer Models (II)



# How Much Missing Data Is A Problem?

*"It is often supposed that there exists something like a critical missing rate up to which missing values are not too dangerous. The belief in such a global missing rate is rather stupid."*

*– Vach (1994, 113)*



# What to Do About Missing Data?

- Listwise Deletion...
- Mean Substitution / Imputation
- “Nearest Neighbor” methods
- “Hot Deck” Imputation
- **Multiple Imputation**
- **Model-Based Solutions**

For MAR data:

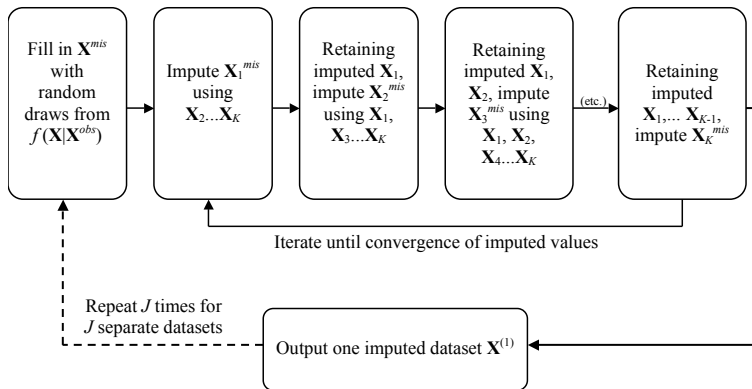
$$\mathbf{R} \perp \mathbf{W} | \mathbf{Z}$$

so  $\mathbf{W}$  and  $\mathbf{Z}$  factorize independently.

Sources of variation we need to consider:

1. Prediction
2. Predictive variation
3. Parameter variation / uncertainty

# MAR: Multiple Imputation



# Multiple Imputation (continued)

Original Data  $\mathbf{X}$  With Missing Data

$i$	$X_1$	$X_2$	$X_3$	$X_4$	...	$X_K$
1	$X_{11}$	$X_{21}$	$X_{31}$	$X_{41}$	...	$X_{K1}$
2	•	$X_{22}$	$X_{32}$	•	...	$X_{K2}$
3	$X_{13}$	$X_{23}$	•	$X_{43}$	...	$X_{K3}$
4	$X_{14}$	•	$X_{34}$	$X_{44}$	...	$X_{K4}$
5	•	$X_{25}$	$X_{35}$	•	...	•
6	$X_{16}$	$X_{26}$	$X_{36}$	$X_{46}$	...	$X_{K6}$
⋮	⋮	⋮	⋮	⋮	⋮	⋮
$N$	$X_{1N}$	$X_{2N}$	$X_{3N}$	$X_{4N}$	...	$X_{KN}$

# Multiple Imputation (continued)

## Iteration One:

Step One: "Fill In" Missing Values of  $\mathbf{X}$

$i$	$X_1$	$X_2$	$X_3$	$X_4$	...	$X_K$
1	$X_{11}$	$X_{21}$	$X_{31}$	$X_{41}$	...	$X_{K1}$
2	$R_{12}$	$X_{22}$	$X_{32}$	$R_{42}$	...	$X_{K2}$
3	$X_{13}$	$X_{23}$	$R_{33}$	$X_{43}$	...	$X_{K3}$
4	$X_{14}$	$R_{24}$	$X_{34}$	$X_{44}$	...	$X_{K4}$
5	$R_{15}$	$X_{25}$	$X_{35}$	$R_{45}$	...	$R_{K5}$
6	$X_{16}$	$X_{26}$	$X_{36}$	$X_{46}$	...	$X_{K6}$
$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$
$N$	$X_{1N}$	$X_{2N}$	$X_{3N}$	$X_{4N}$	...	$X_{KN}$

# Multiple Imputation (continued)

Step Two: Use  $\{X_2, X_3, \dots, X_K\}$  To Impute  $X_1^{\text{mis}}$

$i$	$X_1$	$X_2$	$X_3$	$X_4$	...	$X_K$
1	$X_{11}$	$X_{21}$	$X_{31}$	$X_{41}$	...	$X_{K1}$
2	$\hat{X}_{12}^{(1)}$	$X_{22}$	$X_{32}$	$R_{42}$	...	$X_{K2}$
3	$X_{13}$	$X_{23}$	$R_{33}$	$X_{43}$	...	$X_{K3}$
4	$X_{14}$	$R_{24}$	$X_{34}$	$X_{44}$	...	$X_{K4}$
5	$\hat{X}_{15}^{(1)}$	$X_{25}$	$X_{35}$	$R_{45}$	...	$R_{K5}$
6	$X_{16}$	$X_{26}$	$X_{36}$	$X_{46}$	...	$X_{K6}$
$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$
$N$	$X_{1N}$	$X_{2N}$	$X_{3N}$	$X_{4N}$	...	$X_{KN}$

# Multiple Imputation (continued)

Step Three: Use The Imputed  $X_1$ , Along With  $\{X_3, X_4, \dots, X_K\}$  To Impute  $X_2^{\text{mis}}$

$i$	$X_1$	$X_2$	$X_3$	$X_4$	...	$X_K$
1	$X_{11}$	$X_{21}$	$X_{31}$	$X_{41}$	...	$X_{K1}$
2	$I_{12}^{(1)}$	$X_{22}$	$X_{32}$	$R_{42}$	...	$X_{K2}$
3	$X_{13}$	$X_{23}$	$R_{33}$	$X_{43}$	...	$X_{K3}$
4	$X_{14}$	$I_{24}^{(1)}$	$X_{34}$	$X_{44}$	...	$X_{K4}$
5	$I_{15}^{(1)}$	$X_{25}$	$X_{35}$	$R_{45}$	...	$R_{K5}$
6	$X_{16}$	$X_{26}$	$X_{36}$	$X_{46}$	...	$X_{K6}$
$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$
$N$	$X_{1N}$	$X_{2N}$	$X_{3N}$	$X_{4N}$	...	$X_{KN}$

# Multiple Imputation (continued)

Step Four: Use The Imputed  $X_1$  and  $X_2$ , Along With  $\{X_4, \dots, X_K\}$  To Impute  $X_3^{\text{mis}}$

$i$	$X_1$	$X_2$	$X_3$	$X_4$	...	$X_K$
1	$X_{11}$	$X_{21}$	$X_{31}$	$X_{41}$	...	$X_{K1}$
2	$I_{12}^{(1)}$	$X_{22}$	$X_{32}$	$R_{42}$	...	$X_{K2}$
3	$X_{13}$	$X_{23}$	$I_{33}^{(1)}$	$X_{43}$	...	$X_{K3}$
4	$X_{14}$	$I_{24}^{(1)}$	$X_{34}$	$X_{44}$	...	$X_{K4}$
5	$I_{15}^{(1)}$	$X_{25}$	$X_{35}$	$R_{45}$	...	$R_{K5}$
6	$X_{16}$	$X_{26}$	$X_{36}$	$X_{46}$	...	$X_{K6}$
$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$
$N$	$X_{1N}$	$X_{2N}$	$X_{3N}$	$X_{4N}$	...	$X_{KN}$



(etc.)

# Multiple Imputation (continued)

Step  $K + 1$ : Use The Imputed  $X_1, X_2, \dots, X_{K-1}$  To Impute  $X_K^{\text{mis}}$

$i$	$X_1$	$X_2$	$X_3$	$X_4$	...	$X_K$
1	$X_{11}$	$X_{21}$	$X_{31}$	$X_{41}$	...	$X_{K1}$
2	$I_{12}^{(1)}$	$X_{22}$	$X_{32}$	$I_{42}^{(1)}$	...	$X_{K2}$
3	$X_{13}$	$X_{23}$	$I_{33}^{(1)}$	$X_{43}$	...	$X_{K3}$
4	$X_{14}$	$I_{24}^{(1)}$	$X_{34}$	$X_{44}$	...	$X_{K4}$
5	$I_{15}^{(1)}$	$X_{25}$	$X_{35}$	$I_{45}^{(1)}$	...	$I_{K5}^{(1)}$
6	$X_{16}$	$X_{26}$	$X_{36}$	$X_{46}$	...	$X_{K6}$
$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$
$N$	$X_{1N}$	$X_{2N}$	$X_{3N}$	$X_{4N}$	...	$X_{KN}$

# Multiple Imputation (continued)

## Iteration Two:

Step One: Use The Imputed  $X_2, X_3, \dots, X_K$  To Impute  $X_1^{\text{mis}}$

$i$	$X_1$	$X_2$	$X_3$	$X_4$	...	$X_K$
1	$X_{11}$	$X_{21}$	$X_{31}$	$X_{41}$	...	$X_{K1}$
2	$I_{12}^{(2)}$	$X_{22}$	$X_{32}$	$I_{42}^{(1)}$	...	$X_{K2}$
3	$X_{13}$	$X_{23}$	$I_{33}^{(1)}$	$X_{43}$	...	$X_{K3}$
4	$X_{14}$	$I_{24}^{(1)}$	$X_{34}$	$X_{44}$	...	$X_{K4}$
5	$I_{15}^{(2)}$	$X_{25}$	$X_{35}$	$I_{45}^{(1)}$	...	$I_{K5}^{(1)}$
6	$X_{16}$	$X_{26}$	$X_{36}$	$X_{46}$	...	$X_{K6}$
$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$
$N$	$X_{1N}$	$X_{2N}$	$X_{3N}$	$X_{4N}$	...	$X_{KN}$

# Multiple Imputation (continued)

Step Two: Use The Imputed  $X_1, X_3, \dots, X_K$  To Impute  $X_2^{\text{mis}}$

$i$	$X_1$	$X_2$	$X_3$	$X_4$	...	$X_K$
1	$X_{11}$	$X_{21}$	$X_{31}$	$X_{41}$	...	$X_{K1}$
2	$I_{12}^{(2)}$	$X_{22}$	$X_{32}$	$I_{42}^{(1)}$	...	$X_{K2}$
3	$X_{13}$	$X_{23}$	$I_{33}^{(1)}$	$X_{43}$	...	$X_{K3}$
4	$X_{14}$	$I_{24}^{(2)}$	$X_{34}$	$X_{44}$	...	$X_{K4}$
5	$I_{15}^{(2)}$	$X_{25}$	$X_{35}$	$I_{45}^{(1)}$	...	$I_{K5}^{(1)}$
6	$X_{16}$	$X_{26}$	$X_{36}$	$X_{46}$	...	$X_{K6}$
$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$
$N$	$X_{1N}$	$X_{2N}$	$X_{3N}$	$X_{4N}$	...	$X_{KN}$

(etc.)

# Multiple Imputation (continued)

Step  $K$ : Use The Imputed  $X_1, X_2, \dots, X_{K-1}$  To Impute  $X_K^{\text{mis}}$

$i$	$X_1$	$X_2$	$X_3$	$X_4$	...	$X_K$
1	$X_{11}$	$X_{21}$	$X_{31}$	$X_{41}$	...	$X_{K1}$
2	$I_{12}^{(2)}$	$X_{22}$	$X_{32}$	$I_{42}^{(2)}$	...	$X_{K2}$
3	$X_{13}$	$X_{23}$	$I_{33}^{(2)}$	$X_{43}$	...	$X_{K3}$
4	$X_{14}$	$I_{24}^{(2)}$	$X_{34}$	$X_{44}$	...	$X_{K4}$
5	$I_{15}^{(2)}$	$X_{25}$	$X_{35}$	$I_{45}^{(2)}$	...	$I_{K5}^{(2)}$
6	$X_{16}$	$X_{26}$	$X_{36}$	$X_{46}$	...	$X_{K6}$
$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$
$N$	$X_{1N}$	$X_{2N}$	$X_{3N}$	$X_{4N}$	...	$X_{KN}$

# Multiple Imputation: Summary

Basically:

- Repeat this process for  $J \approx 10$  iterations until convergence of the  $I_{ki}^{(j)}$ s.
- Output the resulting imputed data  $\mathbf{X}^{(1)}$ .
- Repeat this entire process  $M$  times to create  $M$  imputed datasets  $\{\mathbf{X}^{(1)}, \mathbf{X}^{(2)}, \dots, \mathbf{X}^{(M)}\}$ .
- Rule of thumb: “Set  $M \geq$  the percentage of cases in your data with missingness.”
- Estimate models and conduct inference using multiple analysis and model averaging (see e.g. Schafer 1997, Ch. 4).

For MNAR data:

$$\Pr(\mathbf{R}) = f(\mathbf{W}, \mathbf{Z})$$

i.e., missingness is *nonignorable*.

Common causes / situations:

- Omitted variables ( $\rightarrow$  can't condition on all elements of  $\mathbf{Z}$ )
- Differential response due to unmeasured factors
- Truncation / censoring



# MNAR and Model-Based Solutions

For MNAR data, we must model the joint distribution  $\Pr(\mathbf{X}, \mathbf{R})$ ...

## Approaches:

- *Selection* model:  $\Pr(\mathbf{X}, \mathbf{R}) = \Pr(\mathbf{X}) \Pr(\mathbf{R}|\mathbf{X})$ 
  - E.g., Heckman (1976, 1979, etc.)
  - Specifies a (usually, regression) model for  $\Pr(\mathbf{R} | \mathbf{X})$
- *Pattern-Mixture* model:  $\Pr(\mathbf{X}, \mathbf{R}) = \Pr(\mathbf{X}|\mathbf{R}) \Pr(\mathbf{R})$   
 $= \Pr(\mathbf{X}|\mathbf{R} = 0) \Pr(\mathbf{R} = 0) +$   
 $\Pr(\mathbf{X}|\mathbf{R} = 1) \Pr(\mathbf{R} = 1)$ 
  - E.g., Glynn, Laird, and Rubin (1986)
  - Mixture-type model across “responders” and “non-responders”
- Others... [see, e.g., Little and Rubin (2002)]

# Multiple Imputation Example: ANES

Earlier, we created a data frame with  $\approx 75\%$  MCAR missingness on the `BidenThermometer` variable:

```
> describe(MCAR.ANES)
      vars    n  mean    sd median trimmed   mad min max range  skew kurtosis   se
MCARBidenTherm  1  583  47.85  34.50   50.0   47.66  51.89  0.0  100  100.0 -0.12   -1.43  1.43
RConservatism   2 1942   4.11   1.75    4.0    4.11   2.97  1.0   7    6.0 -0.05   -1.14  0.04
RHHLaborUnion   3 1942   0.15   0.36    0.0    0.06   0.00  0.0   1    1.0  1.95    1.80  0.01
RFemale          4 1942   0.52   0.50    1.0    0.53   0.00  0.0   1    1.0 -0.10   -1.99  0.01
RLatino          5 1942   0.09   0.28    0.0    0.00   0.00  0.0   1    1.0  2.95    6.71  0.01
RAge10           6 1942   5.27   1.62    5.4    5.29   2.08  1.9   8    6.1 -0.08   -1.13  0.04
REducation       7 1942   3.57   1.07    4.0    3.62   1.48  1.0   5    4.0 -0.26   -0.67  0.02
```

We can multiply impute values for `MCARBidenTherm` using (e.g.) `mice`:

```
> mice.mcar<-mice(MCAR.ANES,m=75,seed=7222009) # MICE object

iter imp variable
1    1  MCARBidenTherm
1    2  MCARBidenTherm
1    3  MCARBidenTherm
.
.
.
5   74  MCARBidenTherm
5   75  MCARBidenTherm
```

# Multiple Imputation Example: ANES (continued)

Re-run the regression on the multiply-imputed data:

```
> fit.imputed.mcar<-with(mice.mcar,lm(MCARBidenTherm~RConservatism+
+                                     RHHLaborUnion+RFemale+RLatino+RAge10+
+                                     REducation))

> summary(pool(fit.imputed.mcar))
```

	term	estimate	std.error	statistic	df	p.value
1	(Intercept)	84.5889	5.1475	16.4330	152.9	1.336e-35
2	RConservatism	-14.0159	0.5088	-27.5478	163.4	2.676e-63
3	RHHLaborUnion	0.4795	2.3107	0.2075	179.2	8.358e-01
4	RFemale	3.5353	1.9616	1.8022	123.1	7.396e-02
5	RLatino	12.0907	3.2318	3.7412	147.3	2.617e-04
6	RAge10	2.8790	0.5532	5.2045	154.3	6.114e-07
7	REducation	0.8884	0.9066	0.9800	131.2	3.289e-01

Compare to the “complete” data::

```
> summary(fit.all)
```

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	86.868	3.306	26.28	< 2e-16 ***
RConservatism	-14.060	0.336	-41.88	< 2e-16 ***
RHHLaborUnion	-0.710	1.578	-0.45	0.653
RFemale	1.943	1.135	1.71	0.087 .
RLatino	9.251	2.042	4.53	0.0000063 ***
RAge10	3.106	0.357	8.71	< 2e-16 ***
REducation	0.666	0.542	1.23	0.219

---  
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 24.8 on 1935 degrees of freedom  
Multiple R-squared: 0.497, Adjusted R-squared: 0.495  
F-statistic: 318 on 6 and 1935 DF, p-value: <2e-16

# Does this work for MNAR data?

## MNAR ANES data:

```
> describe(MNAR.ANES)
      vars   n  mean   sd median trimmed  mad min max range  skew kurtosis   se
MNARBidenTherm  1  621 12.40 17.28    0.0   9.14 0.00 0.0  90  90.0   1.53    2.24 0.69
RConservatism   2 1942  4.11  1.75    4.0   4.11 2.97 1.0   7   6.0  -0.05   -1.14 0.04
RHHLaborUnion   3 1942  0.15  0.36    0.0   0.06 0.00 0.0   1   1.0   1.95    1.80 0.01
RFemale         4 1942  0.52  0.50    1.0   0.53 0.00 0.0   1   1.0  -0.10   -1.99 0.01
RLatino         5 1942  0.09  0.28    0.0   0.00 0.00 0.0   1   1.0   2.95    6.71 0.01
RAge10          6 1942  5.27  1.62    5.4   5.29 2.08 1.9   8   6.1  -0.08   -1.13 0.04
REducation      7 1942  3.57  1.07    4.0   3.62 1.48 1.0   5   4.0  -0.26   -0.67 0.02

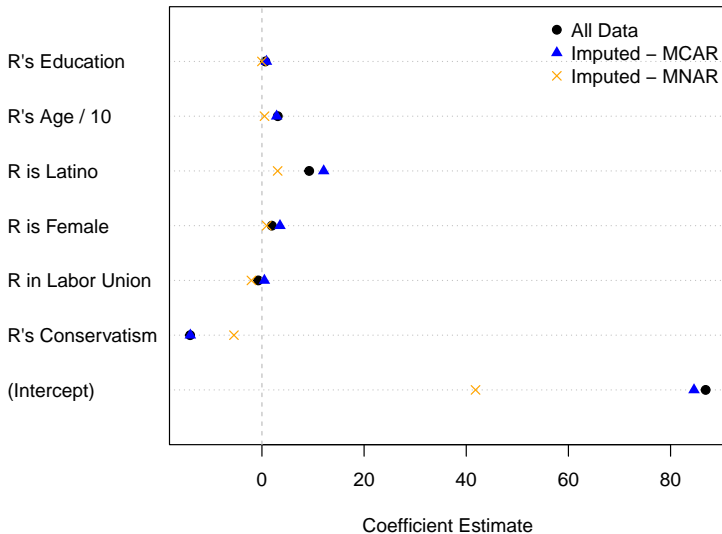
> mice.mnar<-mice(MNAR.ANES,m=75,seed=7222009) # MICE object

iter imp variable
  1   1 MNARBidenTherm
  1   2 MNARBidenTherm
.
.
.

> fit.imputed.mnar<-with(mice.mnar,lm(MNARBidenTherm~RConservatism+RHHLaborUnion+RFemale+RLatino+
+                                     RAge10+REducation))

> summary(pool(fit.imputed.mnar))
      term estimate std.error statistic    df p.value
1 (Intercept) 41.816443   4.9672   8.418599 153.9 2.472e-14
2 RConservatism -5.478921   0.5402  -10.141586 132.6 3.051e-18
3 RHHLaborUnion -2.058441   2.5739   -0.799749 129.1 4.253e-01
4 RFemale      0.936825   1.8626   0.502967 127.6 6.159e-01
5 RLatino      3.073983   3.5359   0.869358 115.8 3.864e-01
6 RAge10       0.508512   0.5681   0.895163 135.3 3.723e-01
7 REducation  -0.004455   0.7982  -0.005581 162.0 9.956e-01
```

# Imputed Thermometer Model Estimated $\hat{\beta}$ s



# Missing Data Resources, R and Otherwise

Check out:

- The [Missing Data CRAN Task View](#)
- Packages:
  - [Amelia](#)
  - [mi](#), [mice](#), and [miceFast](#)
  - [miceMNAR](#) (MNAR imputation using a Heckman-style selection model)
  - [naniar](#) (tidy-cult, but enables [cool visualizations](#))
  - [VIM](#) (joint visualization and imputation of missing data; also used to have a GUI)
  - [Many](#) others...
- van Buuren's [Flexible Imputation of Missing Data 2e](#) e-book