

# ABC's of M-estimation



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# Acknowledgements

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**Section 1:** introduction

*Break (15min)*

**Section 2:** applied examples

*Break (15min)*

**Section 3:** extensions, cautions, conclusions

# Overview: Section 2 - Applied Examples

1. Logistic regression
2. Standardization
  - Outcome model based standardization
  - Inverse probability weighting
3. Data fusion

Each section will include time to work with code

As the goal is to illustrate implementation, we will not dwell on identification assumptions of these examples<sup>1</sup>

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<sup>1</sup>Hernan & Robins Causal Inference: What If; Cole et al. *AJE* 2012

$O_i$ : observed data for unit  $i$

- $O_i = (W_i, X_i, Y_i)$
- $O_i = (R_i, W_i, Y_i)$

$$\text{expit}(a) = 1/(1 + \exp(-a))$$

Estimating function

$$\psi(O_i; \theta)$$

Estimating equation

$$\sum_{i=1}^n \psi(O_i; \theta)$$

# Logistic regression

# Example

Data from the Zambia Preterm Birth Prevention Study (ZAPPS)<sup>2</sup>

$O_i = (Y_i, X_i, W_i)$

$Y_i$ , preterm birth

$X_i$ , anemia

$W_i$ , elevated blood pressure

We want to estimate

$$\begin{aligned}\Pr(Y_i = 1) &= \text{expit}(\beta_0 + \beta_1 X_i + \beta_2 W_i) \\ &= \text{expit}(\mathbf{Z}_i \boldsymbol{\beta}^T)\end{aligned}$$

where  $\mathbf{Z}_i = (1, X_i, W_i)$  and  $\boldsymbol{\beta} = (\beta_0, \beta_1, \beta_2)$

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<sup>2</sup>Castillo et al. *Gates Open Research* 2019



$n = 826$  pregnant people

14.2% delivered preterm

14.3% diagnosed with anemia

18.9% diagnosed with elevated blood pressure

# Estimating $\beta$ by MLE

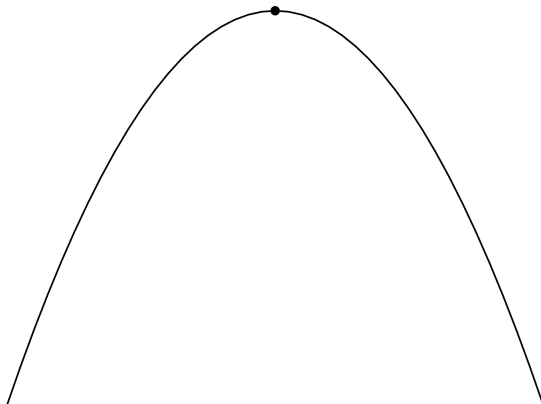
Let  $\tilde{\beta}$  be the estimate of  $\beta$  by MLE

$\tilde{\beta}$  are the values that maximize the log-likelihood

$$\begin{aligned} & \sum_{i=1}^n \ln(L(\mathbf{O}_i, \tilde{\beta})) \\ &= \sum_{i=1}^n Y_i \ln(\text{expit}(\mathbf{Z}_i \tilde{\beta}^T)) + (1 - Y_i) \ln(1 - \text{expit}(\mathbf{Z}_i \tilde{\beta}^T)) \end{aligned}$$

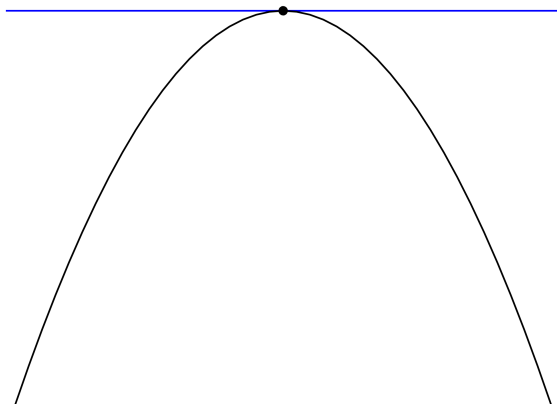
# Estimating $\beta$ by root-finding

Maximum of likelihood is where slope (derivative) is zero ("root")



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# Estimating $\beta$ by root-finding

Our log-likelihood is

$$\ln(L(\mathbf{O}_i, \beta)) = Y_i \ln(\text{expit}(\mathbf{Z}_i \beta^T)) + (1 - Y_i) \ln(1 - \text{expit}(\mathbf{Z}_i \beta^T))$$

Functions for the slope (derivative<sup>3</sup>)

$$\psi(\mathbf{O}_i, \beta) = \frac{\partial \ln(L(\mathbf{O}_i, \beta))}{\partial \beta} = \begin{bmatrix} (Y_i - \text{expit}(\mathbf{Z}_i \beta^T)) 1 \\ (Y_i - \text{expit}(\mathbf{Z}_i \beta^T)) X_i \\ (Y_i - \text{expit}(\mathbf{Z}_i \beta^T)) W_i \end{bmatrix}$$

---

<sup>3</sup>first-order partial derivatives, called the "score" functions

# Estimating $\beta$ by root-finding

Let  $\hat{\beta}$  be the estimate of  $\beta$  by root finding (M-estimation)

Our estimating equation

$$\sum_{i=1}^n \psi(\mathbf{O}_i, \hat{\beta}) = \sum_{i=1}^n \begin{bmatrix} Y_i - \text{expit}(\mathbf{Z}_i \hat{\beta}^T) \\ (Y_i - \text{expit}(\mathbf{Z}_i \hat{\beta}^T)) X_i \\ (Y_i - \text{expit}(\mathbf{Z}_i \hat{\beta}^T)) W_i \end{bmatrix} = \begin{bmatrix} 0 \\ 0 \\ 0 \end{bmatrix}$$

To implement, we need to define the estimating functions

Under **Defining estimating equation**

- R

```
p <- plogis(beta[1] + beta[2]*dat$anemia + beta[3]*dat$bp)

ef_1 <- (dat$ptb - p)
ef_2 <- (dat$ptb - p)*dat$anemia
ef_3 <- (dat$ptb - p)*dat$bp
```

To implement, we need to define the estimating functions

Under **Defining estimating equation**

- SAS

```
p = 1 / (1 + exp(-(beta[1] + beta[2]*anemia + beta[3]*bp)));  
ef_1 = ptb - p;  
ef_2 = (ptb - p)#anemia;  
ef_3 = (ptb - p)#bp;
```



To implement, we need to define the estimating functions

Under **Defining estimating equation**

- Python

```
yhat = inverse_logit(theta[0]*1 +  
                      theta[1]*d['X'] + theta[2]*d['W'])
```

```
residual = d['Y'] - yhat
```

```
score = [residual*1, residual*d['X'], residual*d['W']]
```

# $\tilde{\beta}$ and $\hat{\beta}$ from R

	$\beta_0$	$\beta_1$	$\beta_2$
$\tilde{\beta}$	-1.89450081814 <b>3</b>	0.1187353484 <b>28</b>	0.3605113262 <b>76</b>
$\hat{\beta}$	-1.89450081814 <b>8</b>	0.1187353484 <b>32</b>	0.3605113262 <b>83</b>

Difference is result of algorithm/tolerance

Let  $\mathbf{V}(\boldsymbol{\beta})$  be the  $3 \times 3$  covariance matrix of  $\boldsymbol{\beta}$

The diagonal vector of  $\mathbf{V}/n$  (i.e.,  $\text{diag}(\mathbf{V}/n)$ ) are the finite sample variances

For Wald-type confidence intervals, we use  $\text{SE}(\boldsymbol{\beta}) = \sqrt{\text{diag}(\mathbf{V}/n)}$

# Estimating $V(\tilde{\beta})$ by MLE

$$V(\tilde{\beta}) = I(\tilde{\beta})^{-1}$$

where  $I(\tilde{\beta})$  is the observed information matrix at  $\tilde{\beta}$

There are two ways to estimate  $I(\tilde{\beta})$

1. Hessian-based estimator

$$I(\tilde{\beta}) = \frac{1}{n} \sum_{i=1}^n \left[ -\psi'(\mathbf{O}_i, \tilde{\beta}) \right]$$

$$\text{where } \psi' = \frac{\partial \psi(\mathbf{O}_i, \tilde{\beta})}{\partial \tilde{\beta}} = \frac{\partial^2 \ln(L(\mathbf{O}_i, \beta))}{\partial \tilde{\beta}^2}$$

There are two ways to estimate  $I(\tilde{\beta})$

2. Residual-based estimator

$$I(\tilde{\beta}) = \frac{1}{n} \sum_{i=1}^n \left[ \psi(\mathbf{O}_i, \tilde{\beta}) \psi(\mathbf{O}_i, \tilde{\beta})^T \right]$$

# Estimating $V(\tilde{\beta})$

There are two ways to estimate  $I(\tilde{\beta})$

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$$I(\tilde{\beta}) = \frac{1}{n} \sum_{i=1}^n \left[ -\psi'(\mathbf{O}_i, \tilde{\beta}) \right]$$

2. Residual-based estimator

$$I(\tilde{\beta}) = \frac{1}{n} \sum_{i=1}^n \left[ \psi(\mathbf{O}_i, \tilde{\beta}) \psi(\mathbf{O}_i, \tilde{\beta})^T \right]$$

Asymptotically equal when chosen parametric family is correct  
Hessian is more efficient and default (logistic, glm, GLM)

# Estimating $V(\hat{\beta})$ by sandwich variance estimator

$$V(\hat{\beta}) = B(\hat{\beta})^{-1} F(\hat{\beta}) [B(\hat{\beta})^{-1}]^T$$

Recall

- $B(\hat{\beta}) = \frac{1}{n} \sum_i \left[ -\psi'(O_i, \hat{\beta}) \right]$   
This corresponds to the Hessian-based info matrix estimator
- $F(\hat{\beta}) = \frac{1}{n} \sum_i \left[ \psi(O_i, \hat{\beta}) \psi(O_i, \hat{\beta})^T \right]$   
This corresponds to the residual-based info matrix estimator



When chosen parametric family is correct (here, logistic)

$$\begin{aligned} \mathbf{V}(\boldsymbol{\beta}) &= \mathbf{B}(\boldsymbol{\beta})^{-1} \mathbf{F}(\boldsymbol{\beta}) [\mathbf{B}(\boldsymbol{\beta})^{-1}]^T \\ &\sim \mathbf{I}(\boldsymbol{\beta})^{-1} \mathbf{I}(\boldsymbol{\beta}) [\mathbf{I}(\boldsymbol{\beta})^{-1}]^T \\ &= \mathbf{I}(\boldsymbol{\beta})^{-1} \end{aligned}$$

# Estimating $V(\hat{\beta})$

1. Use the same code for the estimating equation
2. Baking the bread
  - Use software to estimate the Hessian
3. Cooking the filling
  - Use transpose and dot product functions
4. Assembling the sandwich
  - Use inverse function

## Standard errors from R

	$\beta_0$	$\beta_1$	$\beta_2$
MLE, Hessian-based	0.01 <b>4</b> 96046	0.077 <b>6</b> 4837	0.056 <b>6</b> 0453
MLE, Residual-based	0.01 <b>5</b> 08328	0.077 <b>6</b> 1366	0.056 <b>7</b> 0628
Sandwich estimator	0.01 <b>4</b> 84043	0.077 <b>7</b> 2035	0.056 <b>5</b> 2969

Time to work with code

# Standardization

Using the same data as the prior example

$$O_i = (Y_i, X_i, W_i)$$

$Y_i$ , preterm birth

$X_i$ , anemia

$W_i$ , elevated blood pressure

We want to estimate the marginal risk difference and risk ratio of anemia on preterm birth, standardized by elevated blood pressure.

We will illustrate two standardization approaches

- Outcome model based standardization (i.e., g-computation)
- Inverse probability weighting

# Outcome model based standardization

Following implementation illustrated in Snowden et al. AJE 2011

1. Estimate outcome model (from prior example)

$$\Pr(Y_i = 1) = \text{expit}(\beta_0 + \beta_1 X_i + \beta_2 W_i) = \text{expit}(\mathbf{Z}_i \boldsymbol{\beta}^T)$$

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2. Use  $\hat{\boldsymbol{\beta}}$  to predict outcome when  $X_i = 1$ . Take the mean.

$$\hat{\mu}_1 = \frac{1}{n} \sum_{i=1}^n \text{expit}(\hat{\beta}_0 + \hat{\beta}_1 \times 1 + \hat{\beta}_2 W_i)$$



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3. Use  $\hat{\boldsymbol{\beta}}$  to predict outcome when  $X_i = 0$ . Take the mean.

$$\hat{\mu}_2 = \frac{1}{n} \sum_{i=1}^n \text{expit}(\hat{\beta}_0 + \hat{\beta}_1 \times 0 + \hat{\beta}_2 W_i)$$

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$$\hat{\mu}_2 = \frac{1}{n} \sum_{i=1}^n \text{expit}(\hat{\beta}_0 + \hat{\beta}_1 \times 0 + \hat{\beta}_2 W_i)$$

4. Calculate the risk difference  $\hat{\delta}_1 = \hat{\mu}_1 - \hat{\mu}_2$  & log risk ratio  $\hat{\delta}_2 = \ln(\hat{\mu}_1 / \hat{\mu}_2)$

# Outcome model based standardization

Our estimating equation

$$\sum_{i=1}^n \psi(\mathbf{O}_i, \hat{\boldsymbol{\theta}}) = \sum_{i=1}^n \begin{bmatrix} Y_i - \text{expit}(\mathbf{Z}_i \hat{\boldsymbol{\beta}}^T) \\ (Y_i - \text{expit}(\mathbf{Z}_i \hat{\boldsymbol{\beta}}^T)) X_i \\ (Y_i - \text{expit}(\mathbf{Z}_i \hat{\boldsymbol{\beta}}^T)) W_i \\ \text{expit}(\hat{\beta}_0 + \hat{\beta}_1 \times 1 + \hat{\beta}_2 W_i) - \hat{\mu}_1 \\ \text{expit}(\hat{\beta}_0 + \hat{\beta}_1 \times 0 + \hat{\beta}_2 W_i) - \hat{\mu}_2 \\ (\hat{\mu}_1 - \hat{\mu}_2) - \hat{\delta}_1 \\ \ln(\hat{\mu}_1 / \hat{\mu}_2) - \hat{\delta}_2 \end{bmatrix} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \\ 0 \\ 0 \\ 0 \end{bmatrix}$$

where  $\boldsymbol{\theta} = (\boldsymbol{\beta}, \boldsymbol{\mu}, \boldsymbol{\delta})$

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where  $\boldsymbol{\theta} = (\boldsymbol{\beta}, \boldsymbol{\mu}, \boldsymbol{\delta})$

# Estimating functions: Code

- R

```
p <- plogis(beta[1] + beta[2]*dat$anemia + beta[3]*dat$bp)
ef_1 <- (dat$ptb - p)
ef_2 <- (dat$ptb - p)*dat$anemia
ef_3 <- (dat$ptb - p)*dat$bp

ef_r1 <- plogis(beta[1] + beta[2]*1 + beta[3]*dat$bp) - mu[1]
ef_r0 <- plogis(beta[1] + beta[2]*0 + beta[3]*dat$bp) - mu[2]

ef_rd <- (mu[1] - mu[2]) - delta[1]
ef_lnr <- log(mu[1]/mu[2]) - delta[2]
```



- SAS

```
p = 1/(1+exp(-(theta[1] + theta[2]*anemia + theta[3]*bp)));  
ef_1 = ptb - p;  
ef_2 = (ptb - p)#anemia;  
ef_3 = (ptb - p)#bp;  
  
ef_r1 = 1/(1+exp(-(theta[1] + theta[2]*1 + theta[3]*bp)))  
                                     - theta[4];  
  
ef_r0 = 1/(1+exp(-(theta[1] + theta[2]*0 + theta[3]*bp)))  
                                     - theta[5];  
  
ef_rd = j(n,1,(theta[4] - theta[5]) - theta[6]);  
ef_lnrr = j(n,1,log(theta[4]/theta[5]) - theta[7]);
```

# Estimating functions: Code

- Python

```
ee_logit = ee_regression(theta=beta,
                        y=d['Y'],
                        X=d[['intercept', 'X', 'W']],
                        model='logistic')
```

```
y1_hat = inverse_logit(np.dot(d1[['intercept', 'X', 'W']], beta))
y0_hat = inverse_logit(np.dot(d0[['intercept', 'X', 'W']], beta))
```

```
ee_r1 = y1_hat - mu1
ee_r0 = y0_hat - mu0
```

```
ee_rd = np.ones(d.shape[0])*((mu1 - mu0) - delta1)
ee_rr = np.ones(d.shape[0])*(np.log(mu1 / mu0) - delta2)
```

# Inverse probability weighting

1. Estimate propensity score model

$$\Pr(X_i = 1) = \text{expit}(\hat{\alpha}_0 + \hat{\alpha}_1 W_i)$$

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$$\Pr(X_i = 1) = \text{expit}(\hat{\alpha}_0 + \hat{\alpha}_1 W_i)$$

2. Among those with  $X_i = 1$ , take a mean of the outcome, weighted by the inverse probability of  $X_i = 1$ ,

$$\hat{\mu}_1 = \frac{1}{n} \sum_{i=1}^n \left\{ \frac{X_i Y_i}{\text{expit}(\hat{\alpha}_0 + \hat{\alpha}_1 W_i)} \right\}$$

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3. Among those with  $X_i = 0$ , take a mean of the outcome, weighted by the inverse probability of  $X_i = 0$ ,

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where  $\theta = (\alpha, \mu, \delta)$

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- R

```
pscore <- plogis(alpha[1] + alpha[2]*bp)
ef_1 <- (anemia - pscore)
ef_2 <- (anemia - pscore)*bp

wt <- anemia/pscore + (1-anemia)/(1-pscore)
ef_r1 <- anemia*wt*ptb - mu[1]
ef_r0 <- (1 - anemia)*wt*ptb - mu[2]

ef_rd <- (mu[1] - mu[2]) - delta[1]
ef_lnr <- log(mu[1]/mu[2]) - delta[2]
```

- SAS

```
pscore = 1/(1 + exp(-(theta[1] + theta[2]*bp)));  
ef_1 = anemia - pscore;  
ef_2 = (anemia - pscore)#bp;  
  
wt = anemia/pscore + (1-anemia)/(1-pscore);  
ef_r1 = anemia#wt#ptb - theta[3];  
ef_r0 = (1-anemia)#wt#ptb - theta[4];  
  
ef_rd = j(n,1,(theta[3] - theta[4]) - theta[5]);  
ef_lnr = j(n,1,log(theta[3]/theta[4]) - theta[6]);
```

# Estimating functions: Code

- Python

```
ee_logit = ee_regression(theta=alpha ,  
                        y=d['X'],  
                        X=d[['intercept', 'W']],  
                        model='logistic')  
  
pscore = inverse_logit(np.dot(d1[['intercept', 'W']], alpha))  
wt = d['X']/pscore + (1-d['X'])/(1-pscore)  
  
ee_r1 = d['X']*d['Y']*wt - mu1  
ee_r0 = (1-d['X'])*d['Y']*wt - mu0  
  
ee_rd = np.ones(d.shape[0])*((mu1 - mu0) - delta1)  
ee_rr = np.ones(d.shape[0])*(np.log(mu1 / mu0) - delta2)
```

## Aside: IPW GEE trick for conservative SE/intervals

The propensity score parameters are excluded from the stack of estimating functions, i.e. treats  $\alpha$  parameters as known

$$\sum_{i=1}^n \psi(\mathbf{O}_i, \hat{\boldsymbol{\theta}}) = \sum_{i=1}^n \begin{bmatrix} \cancel{X_i - \text{expit}(\hat{\alpha}_0 + \hat{\alpha}_1 W_i)} \\ \cancel{(X_i - \text{expit}(\hat{\alpha}_0 + \hat{\alpha}_1 W_i)) W_i} \\ \frac{X_i Y_i}{\text{expit}(\hat{\alpha}_0 + \hat{\alpha}_1 W_i)} - \hat{\mu}_1 \\ \frac{(1-X_i) Y_i}{1 - \text{expit}(\hat{\alpha}_0 + \hat{\alpha}_1 W_i)} - \hat{\mu}_2 \\ (\hat{\mu}_1 - \hat{\mu}_2) - \hat{\delta}_1 \\ \ln(\hat{\mu}_1 / \hat{\mu}_2) - \hat{\delta}_2 \end{bmatrix}$$

Then use sandwich variance estimator

Time to work with code

# Data Fusion



"Fusion" study designs combine data from multiple sources<sup>4</sup>

## Examples

- Using external validation data to address measurement error
- Transporting a parameter to an external target population
- Bridged treatment comparisons (combine trials)

---

<sup>4</sup>Cole et al. AJE 2022

## Example: Misclassification

We want to estimate the point prevalence of HIV treatment for 950 HIV positive adults in MACS/WIHS cohort study ( $R = 1$ ) in 1995<sup>5</sup>

$$\Pr(Y = 1 | R = 1)$$

In MACS/WIHS, HIV treatment is measured by self-report which may be inaccurate. Let  $W$  be self-reported treatment.

In a separate data source ( $R = 0$ ), we have data on both self-reported treatment  $W$  and treatment documented in medical and pharmacy records (the gold-standard,  $Y$ ) for 331 individuals.

$$O_i = (R_i, Y_i(1 - R_i), W_i)$$

---

<sup>5</sup>Cole et al. AJE 2010;171(1)

Study cohort: 680 reported HIV treatment

$$\hat{\Pr}(W = 1|R = 1) = 0.716$$

Validation data

		Medical Records	
		+	-
Reported	+	204	18
	-	38	71
		242	89

Sensitivity:  $\hat{\Pr}(W = 1|Y = 1) = 0.843$

Specificity:  $\hat{\Pr}(W = 0|Y = 0) = 0.798$

Let  $Se = \Pr(W = 1|Y = 1)$  (sensitivity)

Let  $Sp = \Pr(W = 0|Y = 0)$  (specificity)

$$\Pr(Y = 1) = \frac{\Pr(W = 1) - (1 - Sp)}{Se - (1 - Sp)}$$

Thus, we can calculate the true outcome risk using the mismeasured outcome risk, sensitivity, and specificity

We will apply this Rogan Gladen equation in our study sample

$$\Pr(Y = 1|R = 1) = \frac{\Pr(W = 1|R = 1) - (1 - \text{Sp})}{\text{Se} - (1 - \text{Sp})}$$

leveraging our validation sample to estimate Se and Sp

Four steps to estimate 4 parameters

1. Estimate  $\theta_1 = \Pr(W = 1|R = 1)$
2. Estimate  $\theta_2 = \Pr(W = 1|Y = 1, R = 0)$
3. Estimate  $\theta_3 = \Pr(W = 0|Y = 0, R = 0)$
4. Estimate  $\theta_4 = \Pr(Y = 1|R = 1) = \frac{\theta_1 - (1 - \theta_3)}{\theta_2 - (1 - \theta_3)}$

Our estimating equation is

$$\sum_{i=1}^n \psi(\mathbf{O}_i, \hat{\boldsymbol{\theta}}) = \sum_{i=1}^n \begin{bmatrix} R_i(W_i - \hat{\theta}_1) \\ (1 - R_i)Y_i(W_i - \hat{\theta}_2) \\ (1 - R_i)(1 - Y_i)((1 - W_i) - \hat{\theta}_3) \\ \hat{\phi}(\hat{\theta}_2 - (1 - \hat{\theta}_3)) - (\hat{\theta}_1 - (1 - \hat{\theta}_4)) \end{bmatrix} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \end{bmatrix}$$

Our estimating equation is

$$\sum_{i=1}^n \psi(\mathbf{O}_i, \hat{\boldsymbol{\theta}}) = \sum_{i=1}^n \begin{bmatrix} R_i(W_i - \hat{\theta}_1) \\ (1 - R_i)Y_i(W_i - \hat{\theta}_2) \\ (1 - R_i)(1 - Y_i)((1 - W_i) - \hat{\theta}_3) \\ \hat{\theta}_4(\hat{\theta}_2 - (1 - \hat{\theta}_3)) - (\hat{\theta}_1 - (1 - \hat{\theta}_3)) \end{bmatrix} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \end{bmatrix}$$



# Deriving our estimating functions

$$\text{Step 1. } \theta_1 = \Pr(W = 1|R = 1) = \frac{\Pr(W=1,R=1)}{\Pr(R=1)}$$

---

$$\hat{\theta}_1 = \frac{\sum_{i=1}^n W_i R_i}{\sum_{i=1}^n R_i}$$

$$\hat{\theta}_1 \sum_{i=1}^n R_i = \sum_{i=1}^n W_i R_i$$

$$0 = \sum_{i=1}^n W_i R_i - \sum_{i=1}^n \hat{\theta}_1 R_i$$

$$0 = \sum_{i=1}^n \{W_i R_i - \hat{\theta}_1 R_i\}$$

---

Therefore, our estimating function is  $W_i R_i - \hat{\theta}_1 R_i = R_i(W_i - \hat{\theta}_1)$

Our estimating equation is

$$\sum_{i=1}^n \psi(\mathbf{O}_i, \hat{\boldsymbol{\theta}}) = \sum_{i=1}^n \begin{bmatrix} R_i(W_i - \hat{\theta}_1) \\ (1 - R_i)Y_i(W_i - \hat{\theta}_2) \\ (1 - R_i)(1 - Y_i)((1 - W_i) - \hat{\theta}_3) \\ \hat{\theta}_4(\hat{\theta}_2 - (1 - \hat{\theta}_3)) - (\hat{\theta}_1 - (1 - \hat{\theta}_3)) \end{bmatrix} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \end{bmatrix}$$

Our estimating equation is

$$\sum_{i=1}^n \psi(\mathbf{O}_i, \hat{\boldsymbol{\theta}}) = \sum_{i=1}^n \begin{bmatrix} R_i(W_i - \hat{\theta}_1) \\ (1 - R_i)Y_i(W_i - \hat{\theta}_2) \\ (1 - R_i)(1 - Y_i)((1 - W_i) - \hat{\theta}_3) \\ \hat{\theta}_4(\hat{\theta}_2 - (1 - \hat{\theta}_3)) - (\hat{\theta}_1 - (1 - \hat{\theta}_3)) \end{bmatrix} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \end{bmatrix}$$

# Deriving our estimating functions

$$\text{Step 2. } \theta_2 = \Pr(W = 1|Y = 1, R = 0) = \frac{\Pr(W=1, Y=1, R=0)}{\Pr(Y=1, R=0)}$$

---

$$\hat{\theta}_2 = \frac{\sum_{i=1}^n W_i(1 - R_i)Y_i}{\sum_{i=1}^n (1 - R_i)Y_i}$$

$$\hat{\theta}_2 \sum_{i=1}^n (1 - R_i)Y_i = \sum_{i=1}^n W_i(1 - R_i)Y_i$$

$$0 = \sum_{i=1}^n W_i(1 - R_i)Y_i - \sum_{i=1}^n \hat{\theta}_2(1 - R_i)Y_i$$

$$0 = \sum_{i=1}^n \{W_i(1 - R_i)Y_i - \hat{\theta}_2(1 - R_i)Y_i\}$$

---

Therefore, our estimating function is  $(1 - R_i)Y_i(W_i - \hat{\theta}_2)$

Our estimating equation is

$$\sum_{i=1}^n \psi(\mathbf{O}_i, \hat{\boldsymbol{\theta}}) = \sum_{i=1}^n \begin{bmatrix} R_i(W_i - \hat{\theta}_1) \\ (1 - R_i)Y_i(W_i - \hat{\theta}_2) \\ (1 - R_i)(1 - Y_i)((1 - W_i) - \hat{\theta}_3) \\ \hat{\theta}_4(\hat{\theta}_2 - (1 - \hat{\theta}_3)) - (\hat{\theta}_1 - (1 - \hat{\theta}_3)) \end{bmatrix} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \end{bmatrix}$$

Our estimating equation is

$$\sum_{i=1}^n \psi(\mathbf{O}_i, \hat{\boldsymbol{\theta}}) = \sum_{i=1}^n \begin{bmatrix} R_i(W_i - \hat{\theta}_1) \\ (1 - R_i)Y_i(W_i - \hat{\theta}_2) \\ (1 - R_i)(1 - Y_i)((1 - W_i) - \hat{\theta}_3) \\ \hat{\theta}_4(\hat{\theta}_2 - (1 - \hat{\theta}_3)) - (\hat{\theta}_1 - (1 - \hat{\theta}_3)) \end{bmatrix} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \end{bmatrix}$$

Our estimating equation is

$$\sum_{i=1}^n \psi(\mathbf{O}_i, \hat{\boldsymbol{\theta}}) = \sum_{i=1}^n \begin{bmatrix} R_i(W_i - \hat{\theta}_1) \\ (1 - R_i)Y_i(W_i - \hat{\theta}_2) \\ (1 - R_i)(1 - Y_i)((1 - W_i) - \hat{\theta}_3) \\ \hat{\theta}_4(\hat{\theta}_2 - (1 - \hat{\theta}_3)) - (\hat{\theta}_1 - (1 - \hat{\theta}_3)) \end{bmatrix} = \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \end{bmatrix}$$

# Estimating functions: Code

- R

```
ef_1 <- r*(w - theta[1])  
ef_2 <- (1 - r)*y*(w - theta[2])  
ef_3 <- (1 - r)*(1 - y)*((1 - w) - theta[3])  
ef_4 <- theta[4]*(theta[2] - (1 - theta[3]))  
          - (theta[1] - (1 - theta[3]))
```



- SAS

```
ef_1 = r#(w - theta[1]);  
ef_2 = (1 - r)#y#(w - theta[2]);  
ef_3 = (1 - r)#(1 - y)#((1 - w) - theta[3]);  
ef_4 = j(n,1,  
        (theta[1] - (1 - theta[3]))/(theta[2]  
        - (1 - theta[3])) - theta[4]);
```

# Estimating functions: Code

- Python

```
ee_1 = r*(w - nu)
ee_2 = (1-r) * y * (w - gamma)
ee_3 = (1-r) * (1-y) * ((1-w) - eta)
ee_4 = np.ones(y.shape[0])*phi*(gamma + eta - 1) - (nu + eta - 1)
```

Time to work with code

**Section 1:** introduction

*Break* (15min)

**Section 2:** applied examples

*Break* (15min)

**Section 3:** extensions, cautions, conclusions

Extra slides