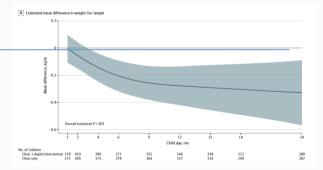
## **Mathematical Statistics II**

STA2212H S LEC9101

Week 11

March 25 2025



Research

#### JAMA | Original Investigation

#### A Digital Health Behavior Intervention to Prevent Childhood Obesity The Greenlight Plus Randomized Clinical Trial

William J. Heerman, MD. MPH: Russell L. Rothman, MD. MPP: Lee M. Sanders, MD. MPH: Jonathan S. Schildcrout, PhD: Kori R. Flower, MD, MS, MPH: Alan M. Delamater, PhD: Melissa C Kay PhD MPH MS RD CLC: Charles T Wood MD MPH: Rachel S Gross MD MS: Aihua Rian MPH: Laura E. Adams, RD. MBA: Evan C. Sommer, BS. BA: H. Shonna Yin, MD. MSc: Eliana M. Perrin, MD. MPH: and the Greenlight Investigators

IMPORTANCE Infant growth predicts long-term obesity and cardiovascular disease. Previous interventions designed to prevent obesity in the first 2 years of life have been largely unsuccessful. Obesity prevalence is high among traditional racial and ethnic minority groups.

OBJECTIVE To compare the effectiveness of adding a digital childhood obesity prevention intervention to health behavior counseling delivered by pediatric primary care clinicians.

DESIGN, SETTING, AND PARTICIPANTS Individually randomized, parallel-group trial conducted at 6 US medical centers and enrolling patients shortly after birth. To be eligible, parents spoke English or Spanish, and children were born after 34 weeks' gestational age. Study enrollment occurred between October 2019 and January 2022, with follow-up through January 2024.

INTERVENTIONS In the clinic-based health behavior counseling (clinic-only) group, pediatric clinicians used health literacy-informed booklets at well-child visits to promote healthy behaviors (n = 451). In the clinic + digital intervention group, families also received health literacy-informed, individually tailored, responsive text messages to support health behavior goals and a web-based dashboard (n = 449).

MAIN OUTCOMES AND MEASURES The primary outcome was child weight-for-length trajectory over 24 months. Secondary outcomes included weight-for-length z score, body mass index (BMI) z score, and the percentage of children with overweight or obesity.

RESULTS Of 900 randomized children, 86.3% had primary outcome data at the 24-month. follow-up time point: 143 (15.9%) were Black, non-Hispanic: 405 (45.0%) were Hispanic: 185 (20.6%) were White, non-Hispanic; and 165 (18.3%) identified as other or multiple races and ethnicities. Children in the clinic + digital intervention group had a lower mean

weight-for-length trajectory, with an estimated reduction of 0.33 kg/m (95% CI, 0.09 to 0.57) at 24 months. There was also an adjusted mean difference of -0.19 (95% CL -0.37 to -0.02) for weight-for-length z score and -0.19 (95% CI, -0.36 to -0.01) for BMI z score. At

Visual Abstract

Multimedia

Supplemental content

### **Today**

- 1. Recap Mar 18 intro to causality
- 2. HW 10 due April 2
- 3. Course evaluation window open
- 4. Theory and methods for missing data
- 5. Project schedule

topics? review?

randomized

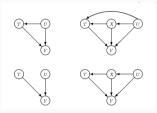
# **Project Schedule April 1**

#### link Start at 10.00

Project Schedule April 1 2025 STA 2212S							
Time	Team Members	Title					
10.00 10.10	Hojung Kim & Markus Kangur Jingxin Wang & Connie Ens	Asymptotics for Lasso type estimators. Testing generalized linear models with					
10.20 10.30	Phyllis Sun & Yufei Liu Abigail McGrory & Aoqi Xie	high-dimensional nuisance parameters.  Longitudinal data analysis using generalized linear models.  Models for exceedances over high thresholds.					
10.40 $10.50$	Lillian Dong & Nevena Ciganovic Regression models and life tables.						
$11.00 \\ 11.10$	Zifan Feng & Shiheng Huang Wenqi Shan & Yunqing Xu	Quantile regression for longitudinal data.  A weakly informative default prior distribution for logistic					
		and other regression models.					
11.20	Break						

## **Recap causality**

- · correlation or association is different than causality
- randomized assignment of treatment to units increases the strength of a causal claim



observational studies can support a causal claim under some assumptions

not testable

- consistency
- no unmeasured confounding

## ... Recap causality

• Potential outcomes Y(a), a = 0 and 1

or continuous

- Observed outcomes  $Y \mid A = a$ ; a = o or 1
- Causal treatment effect  $E\{Y(1) Y(0)\}$

ATE, ACE

- Estimable effect  $E(Y \mid A = 1) E(Y \mid A = 0)$
- blue = red if  $(Y(0), Y(1)) \perp A$

tmt assignment independent of potential outcomes

- in observational studies we rely on adjusting for potential confounders X
- Causal treatment effect  $E_X E\{Y(1) Y(0) \mid X\}$
- Estimable effect  $\int E(Y \mid A = 1, X = x) f_X(x) dx E(Y \mid A = 0, X = x) f_X(x) dx$
- Estimate

$$\frac{1}{n}\sum \hat{r}(1,X_i)-\frac{1}{n}\sum \hat{r}(0,X_i)$$

some fitted model

$$\frac{1}{n}\sum \hat{\mathbf{E}}(Y\mid A=1,X_i)-\frac{1}{n}\sum \hat{\mathbf{E}}(Y\mid A=0,X_i)$$

## ... Recap causality

Estimate

$$\frac{1}{n}\sum_{i=1}^{n}\hat{r}(1,X_i)-\frac{1}{n}\sum_{i=1}^{n}\hat{r}(0,X_i)$$

some fitted model

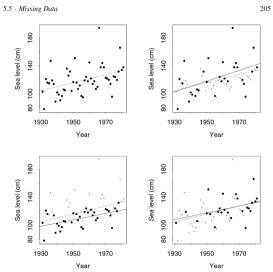
$$\frac{1}{n}\sum \hat{\mathbf{E}}(\mathbf{Y}\mid \mathbf{A}=\mathbf{1},X_i)-\frac{1}{n}\sum \hat{\mathbf{E}}(\mathbf{Y}\mid \mathbf{A}=\mathbf{0},X_i)$$

· A different estimate

$$\frac{1}{n}\sum_{i=1}^n\frac{A_iY_i}{\widehat{\operatorname{pr}}(A=1\mid X_i)}-\frac{1}{n}\sum_{i=1}^n\frac{(1-A_i)Y_i}{\widehat{\operatorname{pr}}(A=0\mid X_i)}-$$

combine these to get a so-called "doubly robust estimator"





• context: independent observations  $(y_i, x_i), i = 1, ..., n$ 

 $x_i$  could be a vector

• model  $f(y \mid x; \theta)$  or sometimes  $f(y, x; \theta)$ 

linear regression; glm; etc

- some observations on  ${\it y}$  may be missing
- e.g. clinical trial,  $x_i$  covariate(s) measured at baseline,  $y_i$  response after treatment, or after some time has elapsed
- observation on subject i becomes  $(y_i, x_i, R_i)$ ,  $R_i = 1$  for complete observation  $R_i = 0$  for incomplete observation

• context: independent observations  $(y_i, x_i), i = 1, ..., n$ 

 $x_i$  could be a vector

• model  $f(y \mid x; \theta)$  or sometimes  $f(y, x; \theta)$ 

linear regression; glm; etc

- some observations on y may be missing
- e.g. clinical trial, x<sub>i</sub> covariate(s) measured at baseline,
   y<sub>i</sub> response after treatment, or after some time has elapsed
- observation on subject i becomes  $(y_i, x_i, R_i)$ ,  $R_i = 1$  for complete observation  $R_i = 0$  for incomplete observation
- · contribution to likelihood function from complete observation

$$f(y_i, x_i, R_i; \theta) = \operatorname{pr}(R_i = 1 \mid x_i, y_i) f(y_i \mid x_i; \theta) f(x_i; \theta)$$

contribution to likelihood function from incomplete observation

no  $\theta$ 

$$f(x_i, R_i; \theta) = \int \operatorname{pr}(R_i = O \mid x_i, y_i) f(y_i \mid x_i; \theta) f(x_i; \theta) dy_i$$

in usual regression settings,  $f(x_i)$ 

• contribution to likelihood function from incomplete observation

$$R_i = 0$$

$$f(x_i, R_i; \theta) = \int \operatorname{pr}(R_i = O \mid x_i, y_i) f(y_i \mid x_i; \theta) f(x_i; \theta) dy_i$$

- missing completely at random:  $pr(R_i = o \mid x_i, y_i) = pr(R_i = o)$
- missing at random:  $pr(R_i = o \mid x_i, y_i) = pr(R_i = o \mid x_i)$
- non-ignorable non-response  $pr(R_i = o \mid x_i, y_i)$

MCAR MAR

no simplification

no simplification

• contribution to likelihood function from incomplete observation

$$R_i = 0$$

$$f(x_i, R_i; \theta) = \int \operatorname{pr}(R_i = O \mid x_i, y_i) f(y_i \mid x_i; \theta) f(x_i; \theta) dy_i$$

 $\operatorname{pr}(R_i = O \mid X_i, V_i) = \operatorname{pr}(R_i = O \mid X_i)$ 

• missing completely at random:  $pr(R_i = o \mid x_i, y_i) = pr(R_i = o)$ 

MAR

**MCAR** 

- missing at random:
- non-ignorable non-response  $pr(R_i = o \mid x_i, y_i)$

no simplification

• likelihood function for sample  $(y_i, x_i, R_i)$ , i = 1, ..., n

$$L(\theta; \mathbf{R}, \mathbf{x}, \mathbf{y}) = \prod_{i \in \mathcal{M}} \int \operatorname{pr}(R_i = 0 \mid x_i, y_i) f(y_i \mid x_i; \theta) f(x_i; \theta) dy_i \times \prod_{i \notin \mathcal{M}} \operatorname{pr}(R_i = 1 \mid x_i, y_i) f(y_i \mid x_i; \theta) f(x_i; \theta)$$

10

• likelihood function for sample  $(y_i, x_i, R_i), i = 1, \dots, n$ 

$$L(\theta; \boldsymbol{R}, \boldsymbol{x}, \boldsymbol{y}) = \prod_{i \in \mathcal{M}} \int \operatorname{pr}(R_i = 0 \mid x_i, y_i) f(y_i \mid x_i; \theta) f(x_i; \theta) dy_i \times \prod_{i \notin \mathcal{M}} \operatorname{pr}(R_i = 1 \mid x_i, y_i) f(y_i \mid x_i; \theta) f(x_i; \theta)$$

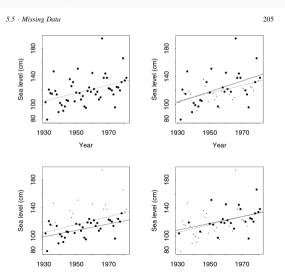
• under MAR or MCAR,

$$L(\theta; \mathbf{R}, \mathbf{x}, \mathbf{y}) \propto \prod_{i=1}^{n} f(y_i \mid x_i; \theta) f(x_i; \theta)$$

- and very often  $f(x_i)$  free of  $\theta$ , so  $L(\theta) \propto \prod_{i=1}^n f(y_i \mid x_i; \theta)$  as usual
- expected information  $I(\theta) = \mathbb{E}_{\theta}\{-\ell''(\theta)\}$  will depend on  $\operatorname{pr}(R_i = 1)$
- use observed information  $J(\hat{ heta}) = -\ell''(\hat{ heta})$  for estimating standard error of MLE

#### Annual maximum sea-level in Venice, 1931 - 1981

Figure 5.12 Missing data in straight-line regression for Venice sea-level data. Clockwise from top left: original data, data with values missing completely at random, data with values missing at random missingness depends on x but not on v, and data with non-ignorable non-response missingness depends on both x and v. Missing values are represented by a small dot. The dotted line is the fit from the full data, the solid lines those from the non-missing data



simulate 1000 samples from linear model with  $\beta_0=$  120,  $\beta_1=$  0.5,  $\sigma=$  20 generate missing data indicators as

$$pr(R = 1 \mid x, y) = \begin{cases} 0.5, \\ \Phi\{0.05(x - \bar{x})\}, \\ \Phi[0.05(x - \bar{x}) + \{y - \beta_0 - \beta_1(x - \bar{x})\}/\sigma] \end{cases}$$

...Simulations SM 5.3

$$pr(R = 1 \mid x, y) = \begin{cases} 0.5, \\ \Phi\{0.05(x - \bar{x})\}, \\ \Phi[0.05(x - \bar{x}) + \{y - \beta_0 - \beta_1(x - \bar{x})\}/\sigma] \end{cases}$$

		Avera	ge estimate (av	(average standard error)				
	Truth	Full	MCAR	MAR	NIN			
$eta_0 \ eta_1$	120 0.50	120 (2.79) 0.49 (0.19)	120 (4.02) 0.48 (0.28)	120 (4.73) 0.50 (0.32)	132 (3.67) 0.20 (0.25)			

To assess the extent of this bias, we generated 1000 samples from a model with parameters  $\beta_0 = 120$ ,  $\beta_1 = 0.5$  and  $\sigma = 20$ , close to the estimates for the Venice data

random (MAR) and with non-ignorable non-response (NIN). 1000 samples were taken. Standard errors for the averages for  $\hat{\beta}_0$  and  $\hat{\beta}_1$  are at most 0.16 and 0.01; those for their standard

errors are at most 0.013 and

0.002.

Table 5.8 Average estimates and standard errors for missing value simulation based on Venice data, for full dataset, with data missing completely at random (MCAR), missing at

and with the same covariate x. We then computed maximum likelihood estimates for Mathematike full data and for those observations that remain after applying the non-response

208

٠	Models

	Magnesium	Control			
Trial	r/m	r/m	n	$\widehat{\mu}$	$(v/n)^{1/2}$
1	1/25	3/23	48	1.18	1.05
2	1/40	2/36	76	0.80	0.83
3	2/48	2/46	94	0.04	0.75
4	1/50	9/53	103	2.14	0.72
5	4/56	14/56	112	1.25	0.69
6	3/66	6/66	132	0.69	0.63
7	2/92	7/93	185	1.24	0.53
8	27/135	43/135	270	0.47	0.44
9	10/160	8/156	316	-0.20	0.41
10	90/1159	118/1157	2316	0.27	0.15
Meta-analysis			3652	0.41	0.11
ISIS-4	2216/29011	2103/29039	58050	-0.05	0.03

Table 5.9 Data from 11 clinical trials to compare magnesium treatment for heart attacks with control. with n patients randomly allocated to treatment and control: there are r deaths out of m patients in each group (Copas, 1999). The estimated log treatment effect  $\widehat{\mu}$  will be positive if treatment is effective:  $(v/n)^{1/2}$  is its standard error. The huge ISIS-4 trial is not included in the meta-analysis.

 $cor(U_1, U_2) = \rho$ 

- study with *n* individuals leads to estimate  $\hat{\mu} \sim N(\mu, \sigma^2/n)$
- study is published (R = 1), if Z > 0 some measure of randomness in publication
- Suppose  $\hat{\mu}$  and Z are related according to the model

$$\hat{\mu} = \mu + \sigma n^{-1/2} U_1, \quad Z = \gamma_0 + \gamma_1 n^{1/2} U_2$$

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$$\hat{\mu} = \mu + \sigma n^{-1/2} U_1, \quad Z = \gamma_0 + \gamma_1 n^{1/2} U_2$$

• 
$$pr(R = 1) = pr(Z > 0) = \Phi(\gamma_0 + \gamma_1 n^{1/2})$$

• 
$$\operatorname{pr}(R = 1 \mid \hat{\mu}) = \operatorname{pr}(Z > 0 \mid \hat{\mu}) = \Phi\left\{\frac{\gamma_0 + \gamma_1 n^{1/2} + \rho n^{1/2}(\hat{\mu} - \mu)/\sigma)}{(1 - \rho^2)^{1/2}}\right\}$$

non-ignorable non-response

unless  $\rho = 0$ 

 $cor(U_1, U_2) = \rho$ 

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non-ignorable non-response

unless  $\rho = 0$ 

 $cor(U_1, U_2) = \rho$ 

• estimate of  $\mu$  is biased:

small  $\gamma_1$ 

$$E(\hat{\mu} \mid R = 1) = \mu + \rho \sigma n^{-1/2} \zeta(\gamma_0 + \gamma_1 n^{1/2})$$
$$= \mu + \rho \sigma \gamma_1 \zeta'(\gamma_0) + \rho \sigma \zeta(\gamma_0) n^{-1/2}$$

• estimate of  $\mu$  is biased:

small 
$$\gamma_1$$

$$E(\hat{\mu} \mid R = 1) = \mu + \rho \sigma n^{-1/2} \zeta(\gamma_0 + \gamma_1 n^{1/2})$$
  
$$\doteq \mu + \rho \sigma \gamma_1 \zeta'(\gamma_0) + \rho \sigma \zeta(\gamma_0) n^{-1/2}$$

- Suppose now we have k published studies of the same treatment  $\hat{\mu}_1, \dots, \hat{\mu}_k$ ,
- assume  $\hat{\mu}_j \sim N(\mu, \sigma^2/n_j)$

same mean, variance depends on study size

 $f(\hat{\mu}_j \mid R_j = 1; \mu, \sigma^2, \rho) = \frac{f(\hat{\mu}_j; \mu, \sigma^2) \operatorname{pr}(R_j = 1 \mid \hat{\mu}_j)}{\operatorname{pr}(R_j = 1)}$ 

log-likelihood function

$$\ell(\theta; \hat{\mu}) = -\sum_{j=1}^{k} \left\{ \frac{1}{2} \log \sigma^2 + \frac{n_j}{2\sigma^2} (\hat{\mu} - \mu)^2 + \log \Phi(a_j) - \log \Phi(b_j) \right\}$$

$$a_j = \gamma_0 + \gamma^1 n_j^{1/2}, b_j = \{a_j + \rho n_j^{1/2} (\hat{\mu}_j - \mu)/\sigma\} (1 - \rho^2)^{-1/2}$$

log-likelihood function

$$\ell(\theta; \hat{\mu}) = -\sum_{i=1}^k \left\{ \frac{1}{2} \log \sigma^2 + \frac{n_j}{2\sigma^2} (\hat{\mu} - \mu)^2 + \log \Phi(a_j) - \log \Phi(b_j) \right\}$$

• if we set 
$$ho = \mathbf{o}$$
,  $\hat{\mu} = \frac{\sum n_j \hat{\mu}_j}{\sum n_j} \sim \mathbf{N}\left(\mathbf{o}, \frac{\sigma^2}{\sum n_j}\right)$ 

no publication bias

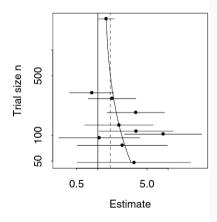
						_	Table 5.9 Data from 11
Trial	Magnesium r/m	Control r/m	п	$\hat{\mu}$	$(v/n)^{1/2}$		clinical trials to compare magnesium treatment for heart uttacks with control,
1	1/25	3/23	48	1.18	1.05		with n patients randomly allocated to treatment and
2	1/40	2/36	76	0.80	0.83		control; there are r deaths
3	2/48	2/46	94	0.04	0.75		out of m patients in each group (Copas, 1999). The
4	1/50	9/53	103	2.14	0.72		estimated log treatment
5	4/56	14/56	112	1.25	0.69		effect $\widehat{\mu}$ will be positive if
6	3/66	6/66	132	0.69	0.63		treatment is effective;
7	2/92	7/93	185	1.24	0.53		$(v/n)^{1/2}$ is its standard
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9	10/160	8/156	316	-0.20	0.41		trial is not included in the meta-analysis.
10	90/1159	118/1157	2316	0.27	0.15		invarannysis.
Meta-analysis			3652	0.41	0.11		
ISIS-4	2216/29011	2103/29039	58050	-0.05	0.03		

 $\exp(\hat{\mu}) = 1.51$ , 95% CI (1.22, 1.86)

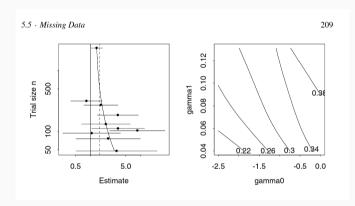
Funnel plot SM 5.5

5.5 · Missing Data

Figure 5.13 Likelihood analysis of magnesium data. Left: funnel plot showing variation of  $\widehat{\mu}$ with trial size n with 95% confidence interval for  $\mu$ based on each trial. The vertical dotted line is the combined estimate of u from the ten small trials. ignoring the possibility of publication bias; the vertical solid line shows no treatment effect. The solid line is the estimated conditional mean (5.33). Right: contours of  $\widehat{\mu}$  as a function of  $\nu_0$  and  $\nu_1$ .



- smaller studies have wider confidence intervals
- seem to be missing small, negative, studies
- simple weighted average is positive (dashed line)
- estimate of average conditional on publication favours smaller studies



back-of the envelope calculation suggests  $\hat{
ho} =$  0.5 and  $\hat{\mu} =$  0.27  $\pm$  0.12

$$\exp(\hat{\mu}) = 1.31, 95\% \text{ CI } (1.03, 1.66)$$

Large RCT (ISIS-4) found no benefit



BMJ 2011:342:d4002 doi: 10.1136/bmi.d4002

Page 1 of 8

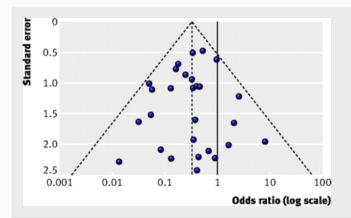
#### **RESEARCH METHODS & REPORTING**

# Recommendations for examining and interpreting funnel plot asymmetry in meta-analyses of randomised controlled trials

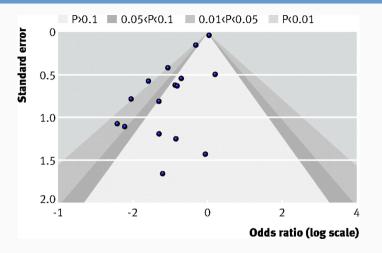
Funnel plots, and tests for funnel plot asymmetry, have been widely used to examine bias in the results of meta-analyses. Funnel plot asymmetry should not be equated with publication bias, because it has a number of other possible causes. This article describes how to interpret funnel plot asymmetry, recommends appropriate tests, and explains the implications for choice of meta-analysis model

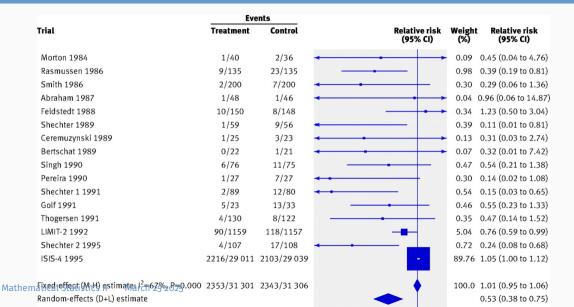
Jonathan A C Steme professor", Alex J Sutton professor and Jonathan A C Steme professor and director\*, Norma Terrin associate professor\*, David R Jones professor\*, Joseph Lau professor and director\*, Carpenter reader\*, Gerta Rücker research assistant\*, Roger M Harbord research associate', Christopher H Schmid professor\*, Jennifer Tetzlaff research coordinator\*, Jonathan J Deeks professor\*, Jaime Peters research fellow\*, Petra Macaskilli associate professor\*, Guido Schwarzer research assistant\*, Sue Duval assistant professor\*1, Douglas G Altman professor\*2, David Moher senior scientifa\*, Julian P T Hiolonis senior statistician\*3

"School of Social and Community Medicine, University of Bristo, Bristo ISBA 975, UK" Department of Health Sciences, University of Lecketter, Locketter, LUSA (See Legal Conference), Technique of Lecketter, Locketter, Lucketter, School of Hydrigen, Statheter, Co. U.School, School of Hydrigen and Topical Medicine, and Health Policy Studies, Taths Medical Context, Obstacl, M.A. U.Sch. "Medical Statistics Unit, London School of Hydrigen and Topical Medicine, University Medical Context, Context, Obstacled Topical Medicine, Context, Context, Obstacled Topical Medicine, University Medical Context, Context, Obstacled Topical Medicine, Context, Obstacled Topical Medicine, University of Medicine, University of Statistics, Obstacled Topical Medicine, University of Statistics, University of Statistic



**Fig 1** Example of symmetrical funnel plot. The outer dashed lines indicate the triangular region within which 95% of studies are expected to lie in the absence of both biases and heterogeneity (fixed effect summary log odds ratio±1.96×standard error of summary log odds ratio). The solid vertical line corresponds to no intervention effect





## Inference with missing data

- if MAR or MCAR, can use usual likelihood-based inference with observed information to estimate variance
- if not, but the missing-ness pattern can be modelled, may be able to adjust estimates accordingly

pub bias

- adjustments will depend on the missing-ness model being correct
- there is a large literature on re-weighting standard estimators to accommodate missing-ness
- the potential outcomes model can be viewed as a type of missing data —
   we see either Y(1) or Y(0) but never both

- · what about missing values of covariates?
- use only complete cases may result in substantial reduction in sample size
- imputation of missing values is a popular choice
- based on prediction of missing covariate value, given observed values of other units

MICE Example									
ID	Age_Original	Income_Original	Age_Imp1	Income_Imp1	Age_Imp2	Income_Imp2			
1	25	50000	25	50000	25	50000			
2		55000	25	55000	50	55000			
3	35		35	65000	35	55000			
4	40	70000	40	70000	40	70000			
5		65000	25	65000	50	65000			
6	50		50	75000	50	65000			
7	45	80000	45	80000	45	80000			
8		90000	35	90000	29	90000			
9	38		38	75000	38	70000			
10	29	75000	29	75000	29	75000			

#### ... Inference with missing data

Research

#### JAMA | Original Investigation

#### A Digital Health Behavior Intervention to Prevent Childhood Obesity The Greenlight Plus Randomized Clinical Trial

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IMPORTANCE Infant growth predicts long-term obesity and cardiovascular disease. Previous interventions designed to prevent obesity in the first 2 years of life have been largely unsuccessful. Obesity prevalence is high among traditional racial and ethnic minority groups.

OBJECTIVE To compare the effectiveness of adding a digital childhood obesity prevention intervention to health behavior counseling delivered by pediatric primary care clinicians.

DESIGN, SETTING, AND PARTICIPANTS Individually randomized, parallel-group trial conducted at 6 US medical centers and enrolling patients shortly after birth. To be eligible, parents spoke English or Spanish, and children were born after 34 weeks' gestational age. Study enrollment occurred between October 2019 and January 2022, with follow-up through January 2024.

INTERVENTIONS In the clinic-based health behavior counseling (clinic-only) group, pediatric clinicians used health literacy-informed booklets at well-child visits to promote healthy behaviors (n = 451). In the clinic + digital intervention group, families also received health literacy-informed, individually tailored, responsive text messages to support health behavior goals and a web-based dashboard (n = 449).

MAIN OUTCOMES AND MEASURES The primary outcome was child weight-for-length trajectory over 24 months. Secondary outcomes included weight-for-length z score, body mass index Mathematical Statis (BMI):z store, and the perdentage of children with overweight or obesity. 

Visual Abstract

Multimedia

Supplemental content

"Missing baseline variables were imputed 1000 times with chained equations" (p.4)

- data  $(X_1, R_1, Y_1), \dots, (X_n, R_n, Y_n)$  i.i.d.
  - 1.  $X_i \sim \text{Uniform from } \{1, \ldots, B\}$
  - 2.  $R_i \sim \text{Bernoulli}(\xi_{X_i})$
  - 3. If  $R_i = 1$ ,  $Y_i \sim \text{Bernoulli}(\theta_{X_i})$
- $\theta = (\theta_1, \dots, \theta_B)$  unknown,  $0 \le \theta_i \le 1$
- $\xi = (\xi_1, \dots, \xi_B)$  known,  $0 < \delta \le \xi_i \le 1 \delta < 1$
- parameter of interest  $\psi = \operatorname{pr}(Y_i = 1) = \sum_{j=1}^B \operatorname{pr}(Y_i = 1 \mid X_i = j) \operatorname{pr}(X_i = j) = \frac{1}{B} \sum_j \theta_j$
- An unbiased estimator of  $\psi$ :

$$\hat{\psi} = \frac{1}{n} \sum_{i=1}^{n} \frac{R_i Y_i}{\xi_{X_i}}$$

- · observed values are averaged, but weighted by probability of being observed
- Horvitz-Thompson estimator

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  - 1.  $X_i \sim \text{Uniform from } \{1, \dots, B\}$
  - 2.  $R_i \sim \text{Bernoulli}(\xi_{X_i})$
  - 3. If  $R_i = 1$ ,  $Y_i \sim \text{Bernoulli}(\theta_{X_i})$
- one term in likelihood function:

$$f(X_i)f(R_i \mid X_i)f(Y_i \mid X_i)^{R_i} = \frac{1}{B}\xi_{X_i}^{R_i}(1-\xi_{X_i})^{1-R_i}\theta_{X_i}^{Y_iR_i}(1-\theta_{X_i})^{(1-Y_i)R_i}$$

- likelihood function:  $L(\theta) \propto \prod_{i=1}^n \theta_{X_i}^{Y_i R_i} (1 \theta_{X_i})^{(1-Y_i)R_i} = \prod_{j=1}^B \theta_j^{n_j} (1 \theta_j)^{m_j}$
- $n_j = \#\{i : Y_i = 1, R_i = 1, X_i = j\},$   $m_j = \#\{i : Y_i = 1, R_i = 0, X_i = j\}$
- most  $n_j, m_j = o$  (B very large)  $\implies$  mle of  $\theta_j$  doesn't exist for many j  $\implies \pi(\theta \mid \mathsf{data}) \propto \pi(\theta)$