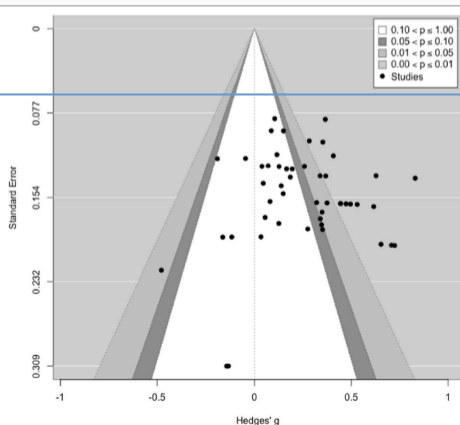


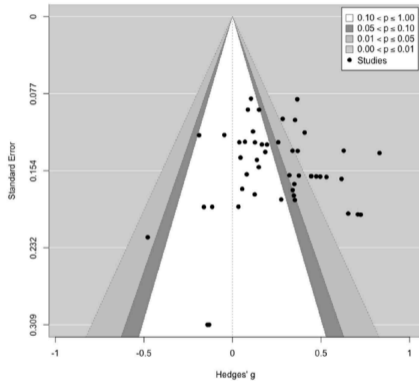
Statistical Theory for Data Science

STA2212H S LEC9101

Week 7

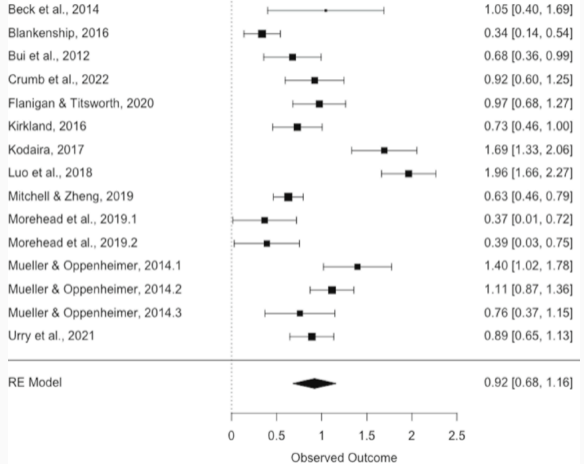
February 24 2026





Note. A positive standardized effect size indicates that students who handwrote their notes had higher achievement than those who typed their notes.

Fig. 1 Funnel plot depicting Hedges' *g* and precision in assessing achievement: handwritten vs. typed note-taking



Today

1. Midterm 2 March 10
2. Recap: Multiple testing, reproducibility
3. Papers re project [Google sheet](#)
4. Inference with missing data
5. Meta-analysis

Recap: Multiple testing

- p -values p_1, \dots, p_m , typically arising from many similar tests on the same data
- e.g. two-sample tests for expressions levels of each of m genes/proteins/test scores/...
- family-wise error rate (fwer) controlled by ensuring

$$\text{pr}(\text{any true null rejected}) \leq \alpha$$

- Bonferroni correction: reject H_{0i} if $p_i \leq \alpha/m$ $\alpha = 0.05$ is conventional

Recap: Multiple testing

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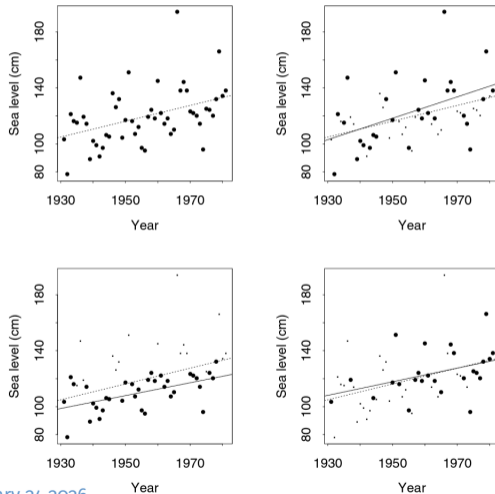
$$\text{pr}(\text{any true null rejected}) \leq \alpha$$

- Bonferroni correction: reject H_{oi} if $p_i \leq \alpha/m$ $\alpha = 0.05$ is conventional
- false discovery rate (fdr) controlled using the BH method Benjamini & Hochberg 1995
- order the p -values $p_{(1)} \leq \dots \leq p_{(m)}$: reject H_{oi} for all i where $p_{(i)} \leq \frac{i}{m}q$ $q = 0.1$ is conventional
- a conservative correction available for dependent p -values

5.5 · Missing Data

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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 y , and data with non-ignorable non-response — missingness depends on both x and y . 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.



- context: independent observations $(y_i, x_i), i = 1, \dots, n$
- model $f(\mathbf{y} \mid \mathbf{x}; \theta)$ or sometimes $f(\mathbf{y}, \mathbf{x}; \theta)$
- some observations on \mathbf{y} may be missing

x_i could be a vector
linear regression; glm; etc

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 - model $f(\mathbf{y} \mid \mathbf{x}; \theta)$ or sometimes $f(\mathbf{y}, \mathbf{x}; \theta)$
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 - 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
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- context: independent observations $(y_i, x_i), i = 1, \dots, n$ x_i could be a vector
- model $f(\mathbf{y} | \mathbf{x}; \theta)$ or sometimes $f(\mathbf{y}, \mathbf{x}; \theta)$ linear regression; glm; etc
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- 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
- contribution to likelihood function from **complete** observation

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

- contribution to likelihood function from **incomplete** observation no θ

$$f(x_i, R_i; \theta) = \int \text{pr}(R_i = 0 | x_i, y_i) f(y_i | x_i; \theta) f(x_i; \theta) dy_i$$

in usual regression settings, $f(x_i)$

- contribution to likelihood function from **incomplete** observation

 $R_i = 0$

$$f(x_i, R_i; \theta) = \int \text{pr}(R_i = 0 \mid x_i, y_i) f(y_i \mid x_i; \theta) f(x_i; \theta) dy_i$$

- missing completely at random:** $\text{pr}(R_i = 0 \mid x_i, y_i) = \text{pr}(R_i = 0)$
- missing at random:** $\text{pr}(R_i = 0 \mid x_i, y_i) = \text{pr}(R_i = 0 \mid x_i)$
- non-ignorable non-response** $\text{pr}(R_i = 0 \mid x_i, y_i)$

MCAR

MAR

no simplification

- contribution to likelihood function from **incomplete** observation

$$R_i = 0$$

$$f(x_i, R_i; \theta) = \int \text{pr}(R_i = 0 \mid x_i, y_i) f(y_i \mid x_i; \theta) f(x_i; \theta) dy_i$$

- missing completely at random:** $\text{pr}(R_i = 0 \mid x_i, y_i) = \text{pr}(R_i = 0)$ MCAR
- missing at random:** $\text{pr}(R_i = 0 \mid x_i, y_i) = \text{pr}(R_i = 0 \mid x_i)$ MAR
- non-ignorable non-response** $\text{pr}(R_i = 0 \mid x_i, y_i)$ no simplification
- likelihood function for sample $(y_i, x_i, R_i), i = 1, \dots, n$

$$L(\theta; \mathbf{R}, \mathbf{x}, \mathbf{y}) =$$

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

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

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

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- under MAR or MCAR ,

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

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

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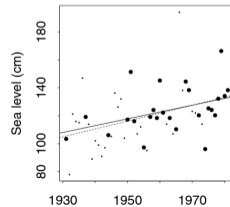
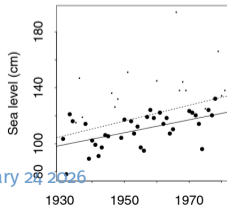
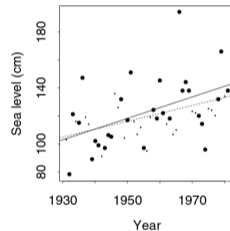
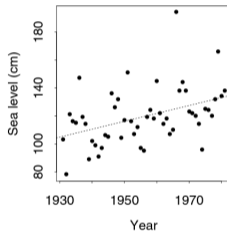
- usually $f(x_i)$ free of θ , so $L(\theta) \propto \prod_{i=1}^n \{f(y_i \mid x_i; \theta)\}^{r_i} = \prod_{\text{complete cases}} f(y_i \mid x_i; \theta)$
- expected information** $I(\theta) = \mathbb{E}_\theta \{-\ell''(\theta)\}$ will depend on $\text{pr}(R_i = 1)$
- use **observed information** $J(\hat{\theta}) = -\ell''(\hat{\theta})$ for estimating standard error of MLE

Annual maximum sea-level in Venice, 1931 – 1981

5.5 · Missing Data

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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 y , and data with non-ignorable non-response — missingness depends on both x and y . 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.



```
> faraway::summary(venice.lm)
              Estimate Std. Error t value Pr(>|t|)
(Intercept)    119.60784    2.60729  45.8744 < 2.2e-16
I(year - mean(year))  0.56697    0.17713   3.2009  0.002406
```

n = 51, p = 2, Residual SE = 18.61977, R-Squared = 0.17

simulate 1000 samples from linear model with $\beta_0 = 120$, $\beta_1 = 0.5$, $\sigma = 20$

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$n = 51$, $p = 2$, Residual SE = 18.61977, R-Squared = 0.17

simulate 1000 samples from linear model with $\beta_0 = 120$, $\beta_1 = 0.5$, $\sigma = 20$

generate missing data indicators as

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

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| | Truth | Average estimate (average standard error) | | | |
|-----------|-------|---|-------------|-------------|-------------|
| | | Full | MCAR | MAR | NIN |
| β_0 | 120 | 120 (2.79) | 120 (4.02) | 120 (4.73) | 132 (3.67) |
| β_1 | 0.50 | 0.49 (0.19) | 0.48 (0.28) | 0.50 (0.32) | 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 and with the same covariate x . We then computed maximum likelihood estimates for the full data and for those observations that remain after applying the non-response mechanisms

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 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 0.09 and 0.002.

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5 · Models

| Trial | Magnesium r/m | Control r/m | n | $\hat{\mu}$ | $(v/n)^{1/2}$ |
|---------------|--------------------|------------------|-------|-------------|---------------|
| 1 | 1/25 | 3/23 | 48 | 1.18 | 1.05 |
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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 $\hat{\mu}$ 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.

- 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 $Y \rightarrow \hat{\mu}, X \rightarrow n$, both pot. missing

$$\hat{\mu} = \mu + \sigma n^{-1/2} U_1, \quad Z = \gamma_0 + \gamma_1 n^{1/2} + U_2, \quad \text{cor}(U_1, U_2) = \rho \geq 0$$

$$U_1 =$$

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- non-ignorable non-response unless $\rho = 0$

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- non-ignorable non-response unless $\rho = 0$
- estimate of μ is biased: small γ_1

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

$\zeta = \phi / \Phi$

- estimate of μ is biased:

small γ_1

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- 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
-

$$f(\hat{\mu}_j \mid R_j = 1; \theta) = \frac{f(\hat{\mu}_j; \theta) \text{pr}(Z_j > 0 \mid \hat{\mu}_j; \theta)}{\text{pr}(Z_j > 0)}$$

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- log-likelihood function

 $\theta =$

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

- log-likelihood function

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

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

- if we set $\rho = 0$, $\hat{\mu} = \frac{\sum n_j \hat{\mu}_j}{\sum n_j} \sim N\left(0, \frac{\sigma^2}{\sum n_j}\right)$

no publication bias

208 5 · Models

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

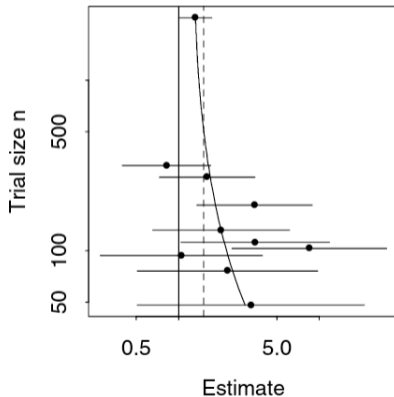
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$$\hat{\mu}_j = \log(r_{2j}/m_{2j}) - \log(r_{1j}/m_{1j}), \quad \sigma^2 \doteq 41.24$$

5.5 · Missing Data

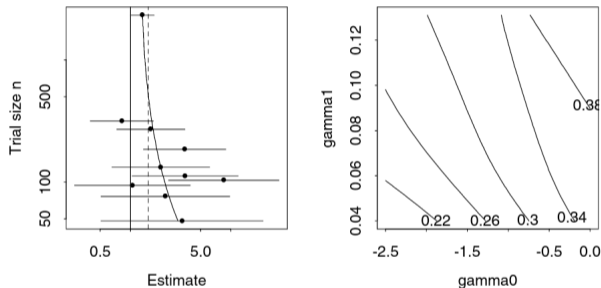
Figure 5.13 Likelihood analysis of magnesium data. Left: funnel plot showing variation of $\hat{\mu}$ with trial size n , with 95% confidence interval for μ based on each trial. The vertical dotted line is the combined estimate of μ from the ten small trials, ignoring the possibility of publication bias; the vertical solid line shows no treatment effect. The solid line is the estimated conditional mean (5.33). Right: contours of $\hat{\mu}$ as a function of γ_0 and γ_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

5.5 · Missing Data

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

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

Large RCT (ISIS-4) found no benefit

February 24 2026

preview

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

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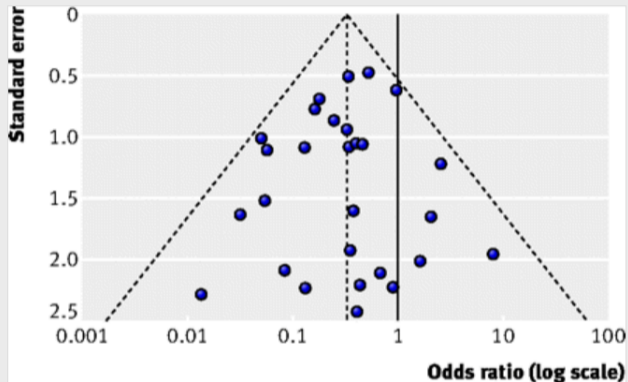
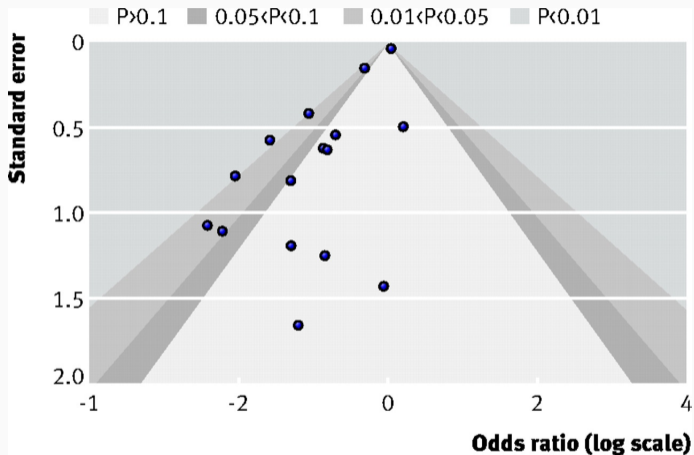
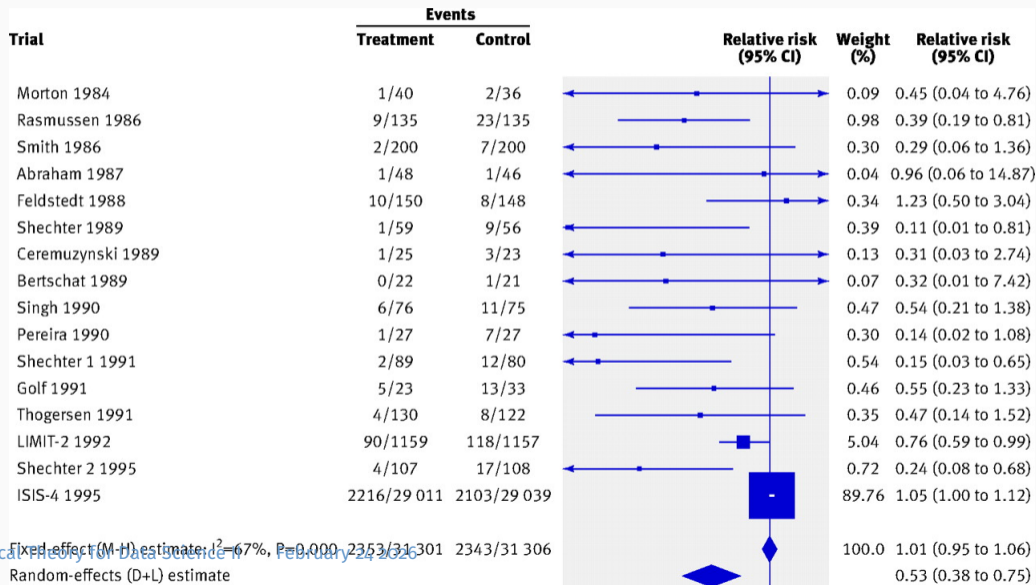


Fig 1 Example of symmetrical funnel plot. The outer dashed lines indicate the triangular region within which 95% of studies are expected to lie in the absence of both biases and heterogeneity (fixed effect summary log odds ratio $\pm 1.96 \times$ standard error of summary log odds ratio). The solid vertical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

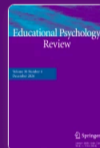


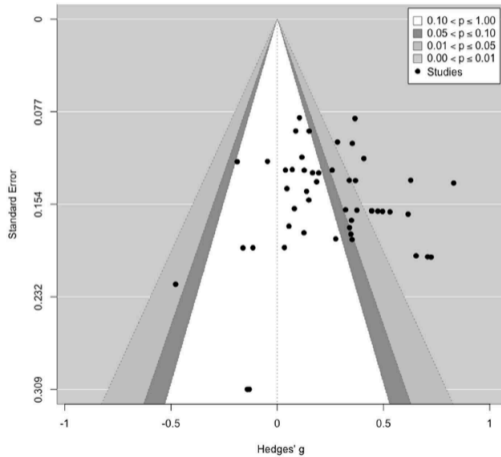


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Typed Versus Handwritten Lecture Notes and College Student Achievement: A Meta-Analysis

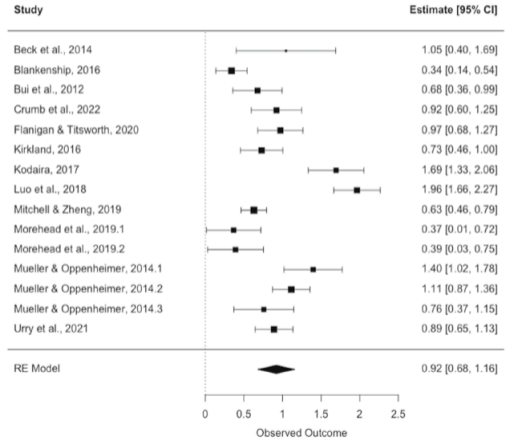
META-ANALYSIS | Published: 12 July 2024

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Note. A positive standardized effect size indicates that students who handwrote their notes had higher achievement than those who typed their notes.

Fig. 1 Funnel plot depicting Hedges' g and precision in assessing achievement: handwritten vs. typed



Note. A positive standardized effect size indicates that students who typed their notes recorded a larger number of words compared to those who handwrote their notes.

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
- adjustments will depend on the missing-ness model being correct
- there is a large literature on re-weighting standard estimators to accommodate missing-ness

pub bias

... Inference with missing data

- 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 |

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 (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

- + 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. $X_i \sim \text{Uniform from } \{1, \dots, B\}$
 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, $0 \leq \theta_j \leq 1$
- $\xi = (\xi_1, \dots, \xi_B)$ known, $0 < \delta \leq \xi_j \leq 1 - \delta < 1$

- data $(X_1, R_1, Y_1), \dots, (X_n, R_n, Y_n)$ i.i.d.
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- $\theta = (\theta_1, \dots, \theta_B)$ unknown, $0 \leq \theta_j \leq 1$
- $\xi = (\xi_1, \dots, \xi_B)$ known, $0 < \delta \leq \xi_j \leq 1 - \delta < 1$
- **parameter of interest** $\psi = \text{pr}(Y_i = 1) = \sum_{j=1}^B \text{pr}(Y_i = 1 \mid X_i = j) \text{pr}(X_i = j) = \frac{1}{B} \sum_j \theta_j$

- An unbiased estimator of ψ :

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

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

IPW estimator

- data $(X_1, R_1, Y_1), \dots, (X_n, R_n, Y_n)$ i.i.d.
 1. $X_i \sim \text{Uniform from } \{1, \dots, B\}$
 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:

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

- data $(X_1, R_1, Y_1), \dots, (X_n, R_n, Y_n)$ i.i.d.
 1. $X_i \sim \text{Uniform from } \{1, \dots, B\}$
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- likelihood function: $L(\boldsymbol{\theta}) \propto \prod_{i=1}^n \theta_{X_i}^{Y_i R_i} (1 - \theta_{X_i})^{(1-Y_i)R_i} = \prod_{j=1}^B \theta_j^{n_j} (1 - \theta_j)^{m_j}$
- $n_j = \#\{i : Y_i = 1, R_i = 1, X_i = j\}$, $m_j = \#\{i : Y_i = 1, R_i = 0, X_i = j\}$
- most $n_j, m_j = 0$ (B very large) \implies mle of θ_j doesn't exist for many j
 $\implies \pi(\boldsymbol{\theta} | \text{data}) \propto \pi(\boldsymbol{\theta})$

$$f(\mathbf{y}; \vartheta) = \sum_{r=1}^p f(\mathbf{y}_r; \vartheta) \pi_r, \quad \mathbf{0} \leq \pi_r \leq \mathbf{1}, \Sigma \pi_r = \mathbf{1}$$

- missing data: $U_1, \dots, U_p; U_r \sim \text{Bernoulli}(\pi_r)$ indexes sub-model
- complete-data log-likelihood function:

$$\log f(\mathbf{y}, \mathbf{u}; \theta) = \sum_{r=1}^p \mathbf{1}(U = r) \{ \log \pi_r + \log f_r(\mathbf{y}; \theta) \} \quad = \log f(\mathbf{y}; \theta) + \log f(\mathbf{u} \mid \mathbf{y}; \theta)$$

$$f(\mathbf{y}; \vartheta) = \sum_{r=1}^p f(\mathbf{y}_r; \vartheta) \pi_r, \quad \mathbf{0} \leq \pi_r \leq \mathbf{1}, \Sigma \pi_r = \mathbf{1}$$

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- maximize $Q(\theta, \theta') = \log f(\mathbf{y}; \theta) + E_{\theta'} \log f(\mathbf{u} | \mathbf{y}; \theta)$
- conditional distribution

$$\text{pr}(U = r | Y = \mathbf{y}; \theta') = \frac{\pi_r' f_r(\mathbf{y}; \theta')}{\sum_{s=1}^p \pi_s' f_s(\mathbf{y}; \theta')} = w_r(\mathbf{y}; \theta')$$

The EM Algorithm

(0) Pick a starting value θ^0 . Now for $j = 1, 2, \dots$, repeat steps 1 and 2 below:

(1) (The E-step): Calculate

$$J(\theta|\theta^j) = \mathbb{E}_{\theta^j} \left(\log \frac{f(Y^n, Z^n; \theta)}{f(Y^n, Z^n; \theta^j)} \mid Y^n = y^n \right).$$

The expectation is over the missing data Z^n treating θ^j and the observed data Y^n as fixed.

(2) Find θ^{j+1} to maximize $J(\theta|\theta^j)$.

Maximizing this function over the five parameters is hard. Imagining that we were given extra information telling us which of the two normals every observation came from. These “complete” data are of the form $(Y_1, Z_1), \dots, (Y_n, Z_n)$, where $Z_i = 0$ represents the first normal and $Z_i = 1$ represents the second. Note that $\mathbb{P}(Z_i = 1) = p$. We shall soon see that the likelihood for the complete data $(Y_1, Z_1), \dots, (Y_n, Z_n)$ is much simpler than the likelihood for the observed data Y_1, \dots, Y_n . ■

Now we describe the EM algorithm.

The EM Algorithm

(0) Pick a starting value θ^0 . Now for $j = 1, 2, \dots$, repeat steps 1 and 2 below:

(1) (The E-step): Calculate

$$J(\theta|\theta^j) = \mathbb{E}_{\theta^j} \left(\log \frac{f(Y^n, Z^n; \theta)}{f(Y^n, Z^n; \theta^j)} \mid Y^n = y^n \right).$$

The expectation is over the missing data Z^n treating θ^i and the observed data Y^n as fixed.

(2) Find θ^{j+1} to maximize $J(\theta|\theta^j)$.

We now show that the EM algorithm always increases the likelihood, that is, $\mathcal{L}(\theta^{j+1}) \geq \mathcal{L}(\theta^j)$. Note that

$$\begin{aligned} J(\theta^{j+1}|\theta^j) &= \mathbb{E}_{\theta^j} \left(\log \frac{f(Y^n, Z^n; \theta^{j+1})}{f(Y^n, Z^n; \theta^j)} \mid Y^n = y^n \right) \\ &= \log \frac{f(y^n; \theta^{j+1})}{f(y^n; \theta^j)} + \mathbb{E}_{\theta^j} \left(\log \frac{f(Z^n|Y^n; \theta^{j+1})}{f(Z^n|Y^n; \theta^j)} \mid Y^n = y^n \right) \end{aligned}$$