

What's the Weight?

Estimating Controlled Outcome Differences in Complex Surveys for Health Disparities Research

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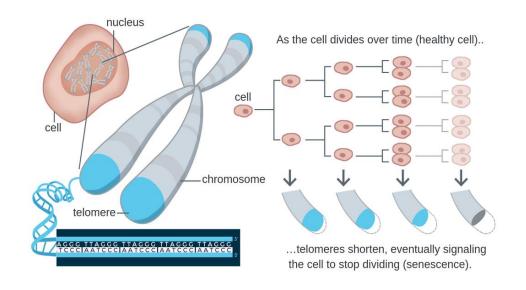


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Paradoxical Relationship Between Telomere Length, Race, and SES

Longer telomeres in Black individuals w/ lower SES, but comparable in similar SES populations

- Regions of DNA that protect against cell death
- Shortening associated w/ cardiometabolic outcomes
- Affected by age, sex, race/ethnicity, genetics,
 SES, environment, psychosocial stressors, ...
- Lower SES often associated w/ shorter telomeres



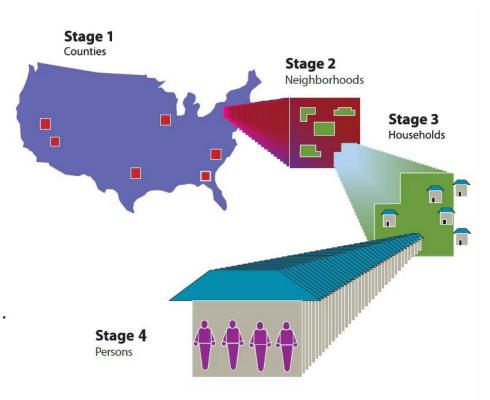
Credit: theory.labster.com/telomere-length

"If we could hypothetically *balance* SES between Black and White individuals in a *nationally representative sample*, would we still see significant Black/White *differences* in telomere length?"

National Health and Nutrition Examination Survey (NHANES)

Nationally Representative Survey by the CDC with a Stratified, Clustered Complex Design

- Primary sampling units (counties)
- Drawn from demographic-specific strata:
 - Oversamples participants living ≤ 130% FPL
 - Oversamples non-Hispanic Black participants
- Rich data from interviews, physical exams, lab tests, ...



Credit: cdc.gov/nchs/nhanes/

Confounding + Selection Bias in a Complex Survey Design

How to weight when selection depends on the group variable under comparison (i.e., race)?

- Within sample confounding, covariate imbalance
- Need to generalizing results to target population (U.S.)
- Statistically challenging because:
 - SES is associated with race
 - Both impact telomere length and selection



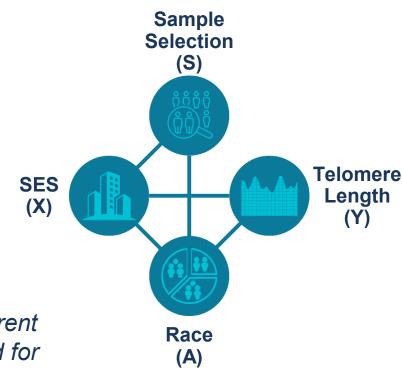
Notation and Target of Inference

Want to Identify a Descriptive Target: Population Average Controlled Difference (ACD)

$$ACD = E_X[E(Y | A = 1, X) - E(Y | A = 0, X)]$$

- A: **Groups** of Interest (race; 1 = Black, 0 = White)
- X: Confounders (SES)
- Y: **Outcome** (log telomere length)
- S: Sample Selection Indicator

Can identify a descriptive or causal target (same estimand, different assumptions), but confounding and selection must be accounted for



G-Formula and Inverse Probability Weighting Approaches

Some Questions: Do you survey weight the propensity model? How to weight the outcome?

Answer: Depends on factorization of the joint probability, $Pr(S = 1, A = a \mid X)$

(1)
$$\mathbb{E}_{X}\left[\mathbb{E}[Y\mid A=a,S=1,X]\cdot\frac{\Pr(S=1)}{\Pr(S=1\mid X)}\mid S=1\right]$$
 Estimate via **g-formula** (1) or **inverse probability weighting** (2, 3)
 (2)
$$\mathbb{E}_{X}\left[\frac{AY}{\Pr(A=a\mid X)}\cdot\frac{\Pr(S=1)}{\Pr(S=1\mid A=a,X)}\mid S=1\right]$$
 Either we **weight** our **propensity score** and specifically take **selection given** $A=a$

(2)
$$\mathbb{E}_{X} \left[\frac{AY}{\Pr(A = a \mid X)} \cdot \frac{\Pr(S = 1)}{\Pr(S = 1 \mid A = a, X)} \middle| S = 1 \right]$$

(3)
$$\mathbb{E}_{X} \left[\frac{AY}{\Pr(A=a \mid S=1,X)} \cdot \frac{\Pr(S=1)}{\Pr(S=1 \mid X)} \mid S=1 \right]$$

Or we fit a within-sample propensity score and marginalize A out of the selection probability

Assumptions for Identifying ACD versus PATE

Pop. average treatment effect (PATE) potential outcome means, E[Ya], w/ stronger assumptions

| Assumption | Definition | ACD | PATE |
|--|--|----------|-------------------------|
| Positivity | $Pr(A=a \mid X=x) > 0 \ \forall a \in A, \text{ every } x \text{ s.t. } f_X(x) > 0$ | ✓ | \checkmark |
| Selection Positivity | $Pr(S=1 \mid A=a, X=x) > 0 \text{ for every a, x s.t. } f_{A,X}(a, x) > 0$ | ✓ | \checkmark |
| Weak Selection Exchangeability | $E[Y A = a, X] = E[Y A = a, S = 1, X]$ or $E[Y^a A = a, X] = E[Y^a A = a, S = 1, X]$ | ✓ | $\overline{\checkmark}$ |
| Stable Unit Treatment Value Assumption | $Y_i = Y_i^a \ \forall i, A_i = a \in A$ | | \checkmark |
| Weak Treatment Exchangeability | E[Y ^a X] = E[Y ^a A=a, X] | | ✓ |

Estimation and Inference

Proposed G-formula (OM) estimator and IPW-based estimators (IPW1, IPW2)

• OM is consistent of the outcome model is correctly specified, IPW1/2 are if the propensity model is

$$\hat{\mu}_{\text{OM}}(a) = \frac{1}{n} \sum_{i=1}^{n} \hat{g}_{a}(X_{i}) \frac{\Pr(S_{i} = 1)}{\Pr(S_{i} = 1 \mid X_{i})} \qquad \hat{\mu}_{\text{IPW1}}(a) = \frac{1}{n} \sum_{i=1}^{n} \frac{I(A_{i} = a)Y_{i}Pr(S_{i} = 1)}{\hat{e}_{a}^{w}(X_{i})Pr(S_{i} = 1 \mid A_{i} = a, X_{i})} \qquad \hat{\mu}_{\text{IPW2}}(a) = \frac{1}{n} \sum_{i=1}^{n} \frac{I(A_{i} = a)Y_{i}Pr(S_{i} = 1)}{\hat{e}_{a}(X_{i})Pr(S_{i} = 1 \mid X_{i})}$$

Inference for ith ind. in jth clust. in kth strat. via empirical sandwich estimator with influence func., φ:

$$\hat{V}(\hat{\theta}) = \sum_{k=1}^{K} \frac{J_k}{J_k - 1} \sum_{i=1}^{J_k} \left\{ \phi_{\cdot jk}(Y, \hat{\theta}, \mathcal{P}) - \bar{\phi}_{\cdot \cdot k}(Y, \hat{\theta}, \mathcal{P}) \right\} \left\{ \phi_{\cdot jk} \left(Y, \hat{\theta}, \mathcal{P} \right) - \bar{\phi}_{\cdot \cdot k}(Y, \hat{\theta}, \mathcal{P}) \right\}^{\top}$$

Augmented Inverse Probability Weighting

Can combine the g-formula (1) and IPW formula (3) to form a doubly robust estimator

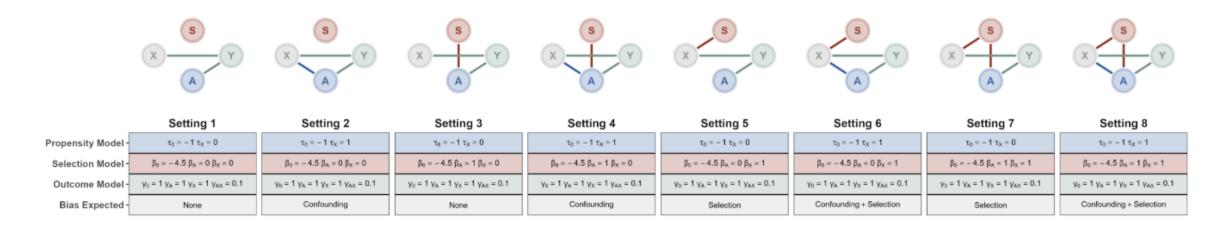
- G-formula most efficient if correctly specified, but
- AIPW Consistent if either the outcome model, g(), or the propensity model, e(), is correctly specified

$$\hat{\mu}_{DR}(a) = \frac{1}{n} \sum_{i=1}^{n} \frac{\Pr(S=1)}{\Pr(S=1 \mid X)} \left\{ \hat{g}_{a}(X_{i}) + \frac{1(A_{i}=a)}{\hat{e}_{a}(X_{i})} \left(Y_{i} - \hat{g}_{a}(X_{i}) \right) \right\}$$

Simulation Study Overview

Compared existing methods to proposed over a range of possible dependence settings

- Generated stratified, clustered population-level confounders, X
- Specified propensity model (A | X), selection model (S | A, X), and outcome model (Y | A, X)
- Systematically varied relationships between X, A, S, and Y:



Simulation Results – No Confounding or Selection Bias

Facilitates a fair comparison – All perform comparably w.r.t. bias, MSE, SE, coverage

| Method | ACD | Est. ACD | Rel. Bias | ASE | MCSE | MSE | Cov. |
|--|-------|----------|-----------|-------|-------|-------|-------|
| Setting 1: No Bias Expected ($\tau_X = 0$, $\beta_A = 0$, $\beta_X = 0$) | | | | | | | |
| Oracle Estimator | 1.003 | 1.000 | -0.003 | 0.109 | 0.107 | 0.012 | 0.964 |
| Simple Reg. | 1.003 | 1.004 | 0.002 | 0.204 | 0.206 | 0.042 | 0.948 |
| Multiple Reg. | 1.003 | 1.001 | -0.002 | 0.109 | 0.107 | 0.012 | 0.958 |
| IPTW Estimator | 1.003 | 1.008 | 0.005 | 0.110 | 0.263 | 0.069 | 0.586 |
| Survey-Weighted Multiple Reg. | 1.003 | 1.001 | -0.002 | 0.109 | 0.108 | 0.012 | 0.956 |
| IPTW Multiple Reg. | 1.003 | 1.001 | -0.002 | 0.109 | 0.108 | 0.012 | 0.958 |
| IPTW + Survey-Weighted Multiple Reg. | 1.003 | 1.001 | -0.002 | 0.109 | 0.108 | 0.012 | 0.956 |
| Weighted IPTW + Survey-Weighted Multiple Reg. | 1.003 | 1.001 | -0.002 | 0.109 | 0.108 | 0.012 | 0.956 |
| Outcome Modeling and Direct Standardization | 1.003 | 1.001 | -0.002 | 0.109 | 0.107 | 0.012 | 0.962 |
| Inverse Probability Weighting 1 | 1.003 | 1.001 | -0.002 | 0.110 | 0.108 | 0.012 | 0.956 |
| Inverse Probability Weighting 2 | 1.003 | 1.002 | -0.001 | 0.113 | 0.110 | 0.012 | 0.960 |
| Augmented Inverse Probability Weighting | 1.003 | 1.001 | -0.002 | 0.110 | 0.108 | 0.012 | 0.958 |

Simulation Results – Confounding + Selection Bias

Assumed relationship in our data – Proposed estimators outperform all current approaches

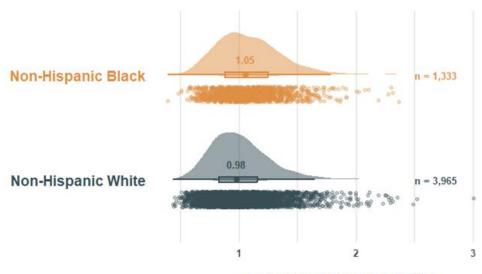
| Method | ACD | Est. ACD | Rel. Bias | ASE | MCSE | MSE | Cov. |
|--|-------|----------|-----------|-------|-------|-------|-------|
| Setting 8: Both Confounding and Selection Bias Expected ($\tau_X = 1, \beta_A = 1, \beta_X = 1$) | | | | | | | |
| Oracle Estimator | 1.001 | 0.991 | -0.010 | 0.097 | 0.093 | 0.009 | 0.956 |
| Simple Reg. | 1.001 | 2.210 | 1.207 | 0.117 | 0.110 | 1.474 | 0.000 |
| Multiple Reg. | 1.001 | 1.232 | 0.230 | 0.073 | 0.079 | 0.059 | 0.126 |
| IPTW Estimator | 1.001 | 3.877 | 2.872 | 0.173 | 0.172 | 8.301 | 0.000 |
| Survey-Weighted Multiple Reg. | 1.001 | 0.994 | -0.007 | 0.118 | 0.134 | 0.018 | 0.920 |
| IPTW Multiple Reg. | 1.001 | 1.233 | 0.231 | 0.074 | 0.081 | 0.060 | 0.144 |
| IPTW + Survey-Weighted Multiple Reg. | 1.001 | 1.016 | 0.015 | 0.109 | 0.123 | 0.015 | 0.912 |
| Weighted IPTW + Survey-Weighted Multiple Reg. | 1.001 | 0.998 | -0.004 | 0.122 | 0.145 | 0.021 | 0.912 |
| Outcome Modeling and Direct Standardization | 1.001 | 0.993 | -0.008 | 0.114 | 0.095 | 0.009 | 0.982 |
| Inverse Probability Weighting 1 | 1.001 | 0.996 | -0.005 | 0.157 | 0.155 | 0.024 | 0.954 |
| Inverse Probability Weighting 2 | 1.001 | 1.017 | 0.016 | 0.226 | 0.264 | 0.070 | 0.968 |
| Augmented Inverse Probability Weighting | 1.001 | 0.992 | -0.010 | 0.154 | 0.158 | 0.025 | 0.946 |

Race + Telomere Length Data from NHANES

5,298 non-Hispanic Black/White adults from 1999 to 2002 with measured telomere length

12 socio-demographic indicators:

Education, Marital Status, Household Size, Household Income, Home Ownership, Home Type, Poverty-Income Ratio, Employment Status, Occupational Category, Insurance Status, Food Security Status, WIC Utilization



8 adjustment covariates:

Age, Sex, White Blood Cell Count, 5-Part Differential

Descriptive Statistics

Univariate tests suggest Black/White differences across all socioeconomic indicators

| Characteristic | Overall, $N = 5,298^{I}$ | Non-Hispanic White, N = 3,965 ¹ | Non-Hispanic Black, N = 1,333 ¹ | p-value |
|---------------------------------|---------------------------------|--|---|---------|
| Telomere Length, Mean T/S Ratio | 1.00 (0.84, 1.18) | 0.98 (0.83, 1.16) | 1.05 (0.88, 1.25) | < 0.001 |
| Age, Years | 50 (35, 67) | 52 (35, 70) | 45 (34, 62) | < 0.001 |
| Sex | | | | 0.4 |
| Male | 2,574 (49%) | 1,939 (49%) | 635 (48%) | |
| Female | 2,724 (51%) | 2,026 (51%) | 698 (52%) | |
| Education | 100 X 10 B 10 10 B 10 C 10 B 10 | Control of the Contro | NOTE: POSSESSE SEA | |
| High School or GED | 2,621 (49%) | 1,790 (45%) | 831 (62%) | |
| Some College | 1,448 (27%) | 1,102 (28%) | 346 (26%) | |
| College Graduate | 1,222 (23%) | 1,068 (27%) | 154 (12%) | |
| Refused/Unknown | 7 (0.1%) | 5 (0.1%) | 2 (0.2%) | |
| Marital Status | | | | < 0.001 |
| Never Married | 754 (14%) | 436 (11%) | 318 (24%) | |
| Widowed/Divorced/Separated | 1,157 (22%) | 775 (20%) | 382 (29%) | |
| Married/Living with Partner | 3,144 (59%) | 2,574 (65%) | 570 (43%) | |
| Refused/Unknown | 243 (4.6%) | 180 (4.5%) | 63 (4.7%) | |
| Household Size | | | 70 07 | < 0.001 |
| • • • | | | | |

Results From Our Study

ACD attenuates as we make appropriate adjustments for confounding + selection bias

- SES *imbalance* across race and *oversampling* not appropriately accounted for by other methods
- Methods which properly incorporate design have more conservative SEs

| Method | ACD Estimate | 95% Confidence Interval |
|--|--------------|-------------------------|
| Multiple Regression | 0.0265 | 0.0106, 0.0424 |
| IPTW Estimator | 0.0262 | 0.0079, 0.0445 |
| Survey-Weighted Multiple Regression | 0.0298 | -0.0010, 0.0606 |
| IPTW Multiple Regression | 0.0183 | 0.0053, 0.0312 |
| IPTW + Survey-Weighted Multiple Regression | 0.0220 | -0.0088, 0.0527 |
| Weighted IPTW + Survey-Weighted Multiple Regression | 0.0186 | -0.0126, 0.0498 |
| Proposed Outcome Modeling and Direct Standardization | 0.0176 | -0.0029, 0.0381 |
| Proposed Inverse Probability Weighting 1 | 0.0141 | -0.0134, 0.0416 |
| Proposed Inverse Probability Weighting 2 | 0.0131 | -0.0081, 0.0342 |
| Proposed Augmented Inverse Probability Weighting | 0.0122 | -0.0077, 0.0321 |

Conclusions

Some thoughts on the approach and our results

- Approach for estimating controlled outcome differences when the group variable of interest and its confounders both influence sample selection
- Proposal minimizes bias and achieves correct inference compared to standard analysis methods
- Context of studying racial disparities presents these challengess in such a way that should be rigorously studied for best practice recommendations

Conclusions

Some thoughts on future work

- Focus on *complex surveys*, but concepts readily extend to other *observational settings*
- Areas of interest for future work include:
 - *Electronic health record* data with unknown sampling probabilities
 - Expanded *relationship diagrams*
 - Extending this framework to two-stage sampling or sequential designs
 - Time-to-event outcomes, AI/ML assisted surveys

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Thank You!

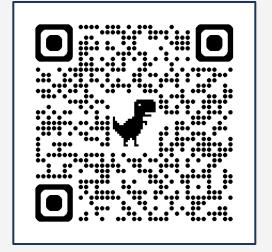








Paper



R Package