

A Sampling of IMPACT Research: Methods for Analysis with Dropout and Identifying Optimal Treatment Regimes

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Outline

- Overview: Projects 2 and 5
- Methods for primary and longitudinal analyses in the presence of dropout
- Identifying optimal treatment regimes from a restricted, feasible set
- Computational Resource and Dissemination Core

Overview

IMPACT:

- *P01 Program Project* grant from NCI
- Five *research projects*
- Three *cores*

Focus here: Research being carried out in *two* of the projects

- *Project 2*: Methods for Missing and Auxiliary Covariates in Clinical Trials
- *Project 5*: Methods for Discovery and Analysis of Dynamic Treatment Regimes
- *Of necessity*: Simplest cases

Project 2

Specific aims:

1. Improving efficiency of inferences in randomized clinical trials using auxiliary covariates
2. *Methods for primary and longitudinal analyses in the presence of drop-out*
3. Diagnostic measures for longitudinal and joint models in the presence of missing data
4. Inference for sensitivity analyses of missing data

Doubly robust methods in the presence of dropout

Motivation:

- Subject drop-out is *commonplace* in clinical trials
- Particularly problematic in studies of *longitudinal markers*, e.g., QOL measures, biomarkers
- *Monotone* pattern of missingness

Missing at random (MAR): Probability of drop-out depends only on information *observed prior to drop-out*

- *Likelihood methods*: Do not require specification of drop-out mechanism but do require *correct* full data model
- *Inverse weighted methods*: Do not require full data model but do require *correct* drop-out model
- *Doubly robust methods*: Require both, but *only one* need be correct

Doubly robust methods in the presence of dropout

Doubly robust methods:

- *Obvious* appeal
- But “*usual*” doubly robust methods can exhibit *disastrous performance* under “*slight*” model misspecification (Kang and Schafer, 2007)

Goal: Can *alternative* doubly robust methods be developed that do not suffer this shortcoming?

The simplest setting

Clinical trial:

- *Outcome* Y , interested in $\mu = E(Y)$
- *Full data*: (Y_i, X_i) , $i = 1, \dots, n$, iid, $X_i =$ *baseline covariates* for subject i
- But Y_i is *missing* for some i (e.g., due to *drop-out*)
- *Observed data*: $(R_i, R_i Y_i, X_i)$, $i = 1, \dots, n$, iid, $R_i = I(Y_i \text{ observed})$

MAR assumption: $R_i \perp\!\!\!\perp Y_i \mid X_i$, implies

$$\mu = E(Y) = E\{E(Y|X)\} = E\{E(Y|R = 1, X)\} \quad (1)$$

Estimators for μ

Outcome regression estimator: MAR (1) suggests *positing* a model $m(X, \beta)$ for $E(Y|X)$

$$\hat{\mu}^{OR} = n^{-1} \sum_{i=1}^n m(X_i, \hat{\beta}) \quad \text{for some } \hat{\beta}$$

- By MAR (1), can use *complete cases* with $R_i = 1$; e.g. *least squares*

$$\sum_{i=1}^n R_i \{Y_i - m(X_i, \beta)\} m_{\beta}(X_i, \beta) = 0, \quad m_{\beta}(X, \beta) = \frac{\partial m(X_i, \beta)}{\partial \beta}$$

- $\hat{\mu}^{OR}$ *consistent* for μ if $m(X, \beta)$ is *correct*

Estimators for μ

Inverse propensity score weighted estimator: *Propensity score*

$P(R = 1|X)$

- If $\pi(X)$ is the *true* propensity score, by MAR

$$n^{-1} \sum_{i=1}^n \frac{R_i Y_i}{\pi(X_i)} \xrightarrow{p} \mu$$

- *Posit* a model $\pi(X, \gamma)$, estimate γ by ML on (R_i, X_i) , $i = 1, \dots, n$

$$\hat{\mu}^{IPW} = n^{-1} \sum \frac{R_i Y_i}{\pi(X_i, \hat{\gamma})}$$

- $\hat{\mu}^{IPW}$ *consistent* for μ if $\pi(X, \gamma)$ is *correct*

Semiparametric theory

Robins et al. (1994): If the *propensity model* is *correct*, with *no additional assumptions* on the distribution of the data

- All *consistent and asymptotically normal* estimators are asymptotically equivalent to estimators of the form

$$n^{-1} \sum \left\{ \frac{R_i Y_i}{\pi(X_i, \hat{\gamma})} + \frac{R_i - \pi(X_i, \hat{\gamma})}{\pi(X_i, \hat{\gamma})} h(X_i) \right\} \quad \text{for some function } h(X)$$

- *Optimal* $h(X)$ leading to *smallest variance (asymptotically)* is

$$h(X) = -E(Y|X)$$

- Suggests *modeling* $E(Y|X)$ by $m(X, \beta)$, estimating β , and estimating μ by

$$n^{-1} \sum \left\{ \frac{R_i Y_i}{\pi(X_i, \hat{\gamma})} - \frac{R_i - \pi(X_i, \hat{\gamma})}{\pi(X_i, \hat{\gamma})} m(X_i, \hat{\beta}) \right\} \quad (2)$$

New perspective

Double robustness: DR

- Such estimators are *consistent* for μ if *either* model is *correct*
- *Kang and Schafer (2007)*: *Simulation scenario* where the “*usual*” DR estimator of form (2) with β *estimated by least squares* is *severely biased and inefficient* when $m(X, \beta)$ and $\pi(X, \gamma)$ are only “*slightly*” misspecified *or* some $\pi(X_i, \hat{\gamma})$ are *close to 0*
- $\hat{\mu}^{OR}$ performed *much better*, even under *misspecification* of $m(X, \beta)$

Key finding: With DR estimators, the method for estimating β *matters*

- The method that is best for estimating β *is not best* for estimating μ
- *Instead*: Find an estimator for β that *minimizes* the (large sample) *variance* of DR estimators of form (2)...

New perspective

Idea: Assume $\pi(X)$ fixed (no unknown γ) and consider estimators

$$n^{-1} \sum \left\{ \frac{R_i Y_i}{\pi(X_i)} - \frac{R_i - \pi(X_i)}{\pi(X_i)} m(X_i, \beta) \right\} \text{ indexed by } \beta \quad (3)$$

- If $\pi(X)$ is *correct* but $m(X, \beta)$ *may not be*, all estimators of form (3) are *consistent* with *asymptotic variance*

$$\text{var}(Y) + E \left[\left\{ \frac{1 - \pi_0(X)}{\pi_0(X)} \right\} \{Y - m(X, \beta)\}^2 \right] \quad (4)$$

- *Minimize* (4) in $\beta \implies \beta^{opt}$ satisfies

$$E \left[\left\{ \frac{1 - \pi_0(X)}{\pi_0(X)} \right\} \{Y - m(X, \beta^{opt})\} m_{\beta}(X, \beta^{opt}) \right] = 0 \quad (5)$$

- Find an estimator $\hat{\beta} \xrightarrow{p} \beta^{opt}$ under these conditions *and* $\hat{\beta} \xrightarrow{p} \text{true } \beta_0$ if $m(X, \beta)$ is *correct* even if $\pi(X)$ is *not*

New perspective

Result: Instead of estimating β by *least squares* solving

$$\sum_{i=1}^n R_i \{Y_i - m(X_i, \beta)\} m_{\beta}(X_i, \beta) = 0,$$

estimate β by a form of *weighted least squares* solving

$$\sum_{i=1}^n R_i \left\{ \frac{1 - \pi(X_i)}{\pi^2(X_i)} \right\} \{Y_i - m(X_i, \beta)\} m_{\beta}(X_i, \beta) = 0 \quad (6)$$

- Estimating equation (6) has *expectation* (5) when $\pi(X)$ is *correct*
- The resulting $\hat{\beta}$ *satisfies* the required conditions
- Can be *generalized* to case of $\pi(X, \gamma)$ with $\hat{\gamma}$ (ML)
- All this extends to *more general* μ (e.g., treatment effect)

New perspective

Details: My website and

Cao, W., Tsiatis, A.A. and Davidian, M. (2009). Improving efficiency and robustness of the doubly robust estimator for a population mean with incomplete data. *Biometrika* **96**, 723–734.

- The DR estimator using this $\hat{\beta}$ *greatly improved* on the “*usual*” DR estimator and exhibited *superior performance* (to $\hat{\mu}^{OR}$) in the Kang and Schafer and other scenarios

Longitudinal study

Extension: Longitudinal study with *drop-out*

- *Ideally*: Collect data L_j at time t_j , $j = 1, \dots, M + 1$
- *Full data*: $\bar{L} = \bar{L}_{M+1} = (L_1, \dots, L_{M+1})$
- *Dropout*: If subject is *last seen* at time t_j , *dropout indicator* $D = j$, observe only $\bar{L}_j = (L_1, \dots, L_j)$
- *Observed data*: iid (D_i, \bar{L}_{D_i}) , $i = 1, \dots, n$
- *Interest*: Parameter μ in a *semiparametric model* for the full data
- *Full data estimator for μ* : Solve

$$\sum_{i=1}^n \varphi(\bar{L}_i, \mu) = 0, \quad E\{\varphi(\bar{L}, \mu)\} = 0$$

- *MAR*: $\text{pr}(D = j | \bar{L})$ depends only on \bar{L}_j , $j = 1, \dots, M + 1$
- *Drop-out model*: $\text{pr}(D = j | \bar{L}) = \pi(j, \bar{L}_j)$, $\pi(M + 1, \bar{L}) = \pi(\bar{L})$

Longitudinal study

If drop-out model correct: All *consistent and asymptotically normal* estimators for μ solve

$$\sum_{i=1}^n \left\{ \frac{I(D_i = M + 1)\varphi(\bar{L}_i, \mu)}{\pi(\bar{L}_i)} + \sum_{j=1}^M \frac{dM_{ji}(\bar{L}_{ji})}{K_{ji}(\bar{L}_{ji})} \mathcal{L}_j(\bar{L}_{ji}) \right\} = 0$$

- $dM_{ji}(\bar{L}_{ji}), K_{ji}(\bar{L}_{ji})$ are functions of $\pi(j, \bar{L}_j)$
- These estimators are *DR*
- *Optimal* $\mathcal{L}_j(\bar{L}_j) = E\{\varphi(\bar{L}, \mu) | \bar{L}_j\}$; *model* by $\mathcal{L}_j(\bar{L}_j, \beta), j = 1, \dots, M$

Result: Can derive *optimal* estimator for β by analogy to the previous Tsiatis, A.A., Davidian, M. and Cao, W. (2011). Improved doubly robust estimation when the data are monotonely coarsened, with application to longitudinal studies with dropout. *Biometrics* **67**, 536–545.

Project 5

Specific aims:

1. Learning methods for optimal dynamic treatment regimes
2. *Identifying optimal dynamic treatment regimes from a restricted, feasible set*
3. Inferential methods for dynamic treatment regimes
4. Design of sequentially randomized trials for dynamic treatment regimes

Optimal treatment regimes from a feasible set

Motivation: *Individualized (personalized)* treatment

- *Premise*: Different subgroups of patients may respond *differently* to treatments
- Treatment decisions *tailored* to individual patients based on their *characteristics*, *disease status*, *medical history*, etc
- *Ideally*: Use *all* relevant information in *decision rules*
- *Realistically*: Use a key subset of information *feasibly* collected in *clinical practice*, simple-to-implement, *interpretable* decision rules

Goal: Methods for estimating such *feasible dynamic treatment regimes* from data from *clinical trials* or *observational databases*

The simplest setting

A single decision: *Two* treatment options

- *Observed data*: (Y_i, X_i, A_i) , $i = 1, \dots, n$, iid
- Y_i outcome, X_i baseline covariates, $A_i = 0, 1$

Treatment regime: A function $g : X \rightarrow \{0, 1\}$

- *Simple example*: $g(X) = I(X \leq 50)$
- $g \in \mathcal{G}$, the class of *all* such regimes
- *Optimal regime*: If followed by *all patients* in the population, would lead to *best average outcome* among all regimes in \mathcal{G}

Potential outcomes

Formalize: $Y^*(1)$ = outcome if patient were to receive 1; similarly $Y^*(0)$

- *Thus*, $E\{Y^*(1)\}$ is the *average outcome* if *all patients* in the population received 1; similarly $E\{Y^*(0)\}$
- *Assume* we observe $Y = Y^*(1)A + Y^*(0)(1 - A)$
- *Assume* $Y^*(0), Y^*(1) \perp\!\!\!\perp A|X$ (*no unmeasured confounders*); automatic in a *randomized trial*
- $\implies E\{Y^*(1)\} = E\{E(Y|A = 1, X)\}$; similarly $E\{Y^*(0)\}$
- For any $g \in \mathcal{G}$, define

$$Y^*(g) = Y^*(1)g(X) + Y^*(0)\{1 - g(X)\} \quad (1)$$

- *Optimal regime*: Leads to *largest* $E\{Y^*(g)\}$ among all $g \in \mathcal{G}$; i.e.,

$$g^{opt}(X) = \arg \max_{g \in \mathcal{G}} E\{Y^*(g)\}$$

Optimal regime

$$(1): E\{Y^*(g)\} = E\left[E(Y|A=1, X)g(X) + E(Y|A=0, X)\{1-g(X)\} \right]$$

$$g^{opt}(X) = I\{E(Y|A=1, X) - E(Y|A=0, X) \geq 0\}$$

- *Thus*: If $E(Y|A, X)$ is *known* can find g^{opt}
- *Posit a model* $\mu(A, X, \beta)$ for $E(Y|A, X)$ and *estimate* β based on observed data $\implies \hat{\beta}$
- *Estimate* g^{opt} by $\hat{g}^{opt}(X) = I\{\mu(1, X, \hat{\beta}) - \mu(0, X, \hat{\beta}) \geq 0\}$
- “*Regression estimator*”
- *But*: $\mu(A, X, \beta)$ may be *misspecified*, so \hat{g}^{opt} could be far from g^{opt}

Alternative perspective: $\mu(A, X, \beta)$ defines a *class* of regimes, *indexed* by β , that *may or may not* contain g^{opt}

- But may be *feasible* and *interpretable*

Optimal restricted regime

For example: Suppose *in truth*

$$E(Y|A, X) = \exp\{1 + X_1 + 2X_2 + 3X_1X_2 + A(1 - 2X_1 + X_2)\}$$

$$\implies g^{opt}(X) = I(X_2 \geq 2X_1 - 1)$$

- *Posit* $\mu(A, X, \beta) = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + A(\beta_3 + \beta_4 X_1 + \beta_5 X_2)$
- Defines *class* \mathcal{G}_η with elements

$$I(X_2 \geq \eta_1 X_1 + \eta_0) \cup I(X_2 \leq \eta_1 X_1 + \eta_0), \quad \eta_0 = -\beta_3/\beta_5, \quad \eta_1 = -\beta_4/\beta_5$$

Thus, in general: Consider *class* $\mathcal{G}_\eta = \{g(X, \eta)\}$ *indexed* by η

- Write $g_\eta(X) = g(X, \eta)$
- *Optimal restricted regime* $g_\eta^{opt}(X) = g(X, \eta^{opt})$,

$$\eta^{opt} = \arg \max_\eta E\{Y^*(g_\eta)\}$$

Optimal restricted regime

Approach: Estimate η^{opt} by maximizing a “good” (*DR*) estimator for $E\{Y^*(g_\eta)\}$

- *Missing data* analogy: “*Full data*” are $\{Y^*(g_\eta), X\}$; “*observed data*” are $(C_\eta, C_\eta Y, X)$, where

$$C_\eta = Ag(X, \eta) + (1 - A)\{1 - g(X, \eta)\}$$

- $\pi(X) = \text{pr}(A = 1|X)$; *known* in a randomized trial; otherwise *model* and *estimate* $\pi(X, \hat{\gamma})$
- $\pi_c(X) = \text{pr}(C_\eta = 1|X) = \pi(X)g(X, \eta) + \{1 - \pi(X)\}\{1 - g(X, \eta)\}$

Optimal restricted regime

Estimator for $E\{Y^*(g_\eta)\}$:

$$n^{-1} \sum_{i=1}^n \left\{ \frac{C_{\eta,i} Y_i}{\pi_c(X_i, \hat{\gamma})} - \frac{C_{\eta,i} - \pi_c(X_i, \hat{\gamma})}{\pi_c(X_i, \hat{\gamma})} m(X_i, \hat{\beta}, \eta) \right\} \quad (2)$$

$$m(X, \beta, \eta) = \mu(1, X, \beta)g(X, \eta) + \mu(0, X, \beta)\{1 - g(X, \eta)\}$$

- *Consistent* if *either* $\pi(X, \gamma)$ or $\mu(A, X, \beta)$ is *correct*
- *Maximize* (2) in η to obtain $\hat{\eta}^{opt}$

Current work:

- Approaches to *maximizing* (2)
- *Simulations*: Almost equals performance of *correct* regression estimator and is *superior* with *misspecified* $\mu(A, X, \beta)$
- *Extension* to *multiple decision points*
- Zhang, Tsiatis, Davidian (2011), *in preparation*

Computational Resource and Dissemination Core

Goals:

- Efficient, robust, reliable code implementing project methodology
- *Creation and dissemination of public-use software* (to be made available on the IMPACT website)
- E.g., R packages, SAS macros, specialized implementations (e.g., FORTRAN, c)

Programmers: Scientific programmers at UNC-CH and NCSU dedicated to these activities