

Estimating the optimal dynamic treatment regime from longitudinal observational data

Liliana Orellana¹, Andrea Rotnitzky^{2,3} and James Robins²

¹Universidad de Buenos Aires, ²Harvard University, ³Universidad Di Tella

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Motivation

- ▶ Highly active antiretroviral therapy (HAART) dramatically decreased morbidity and mortality due to infection with HIV.
- ▶ Eradication of HIV infection cannot be achieved with available antiretroviral regimens.
- ▶ Late initiation of HAART has both risks and benefits:
 - ▶ Risks: Irreversible damage of the immune system; AIDS.
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QUESTION \implies When to start HAART?

- ▶ Decision on "when to start" for *asymptomatic* HIV+ subjects is essentially based on CD4 cell count.

When to start HAART?

U.S. Treatment Guidelines for HIV-1 Infected Adults and Adolescents (October 2006)

Recommendations on when to start for asymptomatic HIV+ subjects:

- ▶ Definitively start if CD4 count < 200 cells/mm³.
- ▶ *Unclear* if CD4 count > 200 cells/mm³.
 - ▶ Offer trx if $200 < \text{CD4} \leq 350$ cells/mm³.
 - ▶ Preferably defer trx if CD4 > 350 cells/mm³.
- ▶ A treatment strategy based on CD4 counts is an example of a *dynamic treatment regime*.

Dynamic treatment regimes

Data

$$L_0, A_0, L_1, A_1, \dots, L_K, A_K, L_{K+1}$$

L_k = clinical and laboratory variables measured during the k^{th} clinic visit,

A_k = treatment prescription at visit k

$\bar{L}_k = (L_0, L_1, \dots, L_k)$ and $\bar{A}_k = (A_0, A_1, \dots, A_k)$

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Dynamic treatment regime

- ▶ Sequential rule for determining, at each time k , the next treatment prescription A_k .
- ▶ Rule inputs the recorded health information up to time k and returns a treatment recommendation

$$\left(\bar{L}_k, \bar{A}_{k-1}\right) \rightarrow d_k \left(\bar{L}_k, \bar{A}_{k-1}\right) \in \mathcal{A}_k, k = 0, \dots, K.$$

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Optimal dynamic regime

- ▶ Maximizes the expectation of some utility function $Y \equiv u\left(\bar{L}_{K+1}, \bar{A}_K\right)$ among the set of candidate regimes.

An ideal randomized study to compare two regimes

Suppose we want to compare two dynamic regimes:

- ▶ start HAART when CD4 falls below 500 (d^{500})
- ▶ start HAART when CD4 falls below 200 (d^{200})

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DESIGN

- ▶ Follow patients periodically, say every 6 months, from HIV diagnosis
- ▶ When CD4 first falls below 500 randomize to
 - ▶ start immediately (say, $p = 1/2$)
 - ▶ start when CD4 first seen to fall below 200 (say, $p = 1/2$)
- ▶ Let Y be the outcome, a utility function of the health and treatment history (higher values are preferable)
- ▶ Compare outcome in the two groups after a number of years of follow-up (e.g., 5 years)

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Estimation goal

We want to compare the expected utility in:

1. a hypothetical world where regime d^{500} was enforced (μ^{500}) versus
2. a hypothetical world where regime d^{200} was enforced (μ^{200}).

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- ▶ randomization generates exchangeable groups and
- ▶ each subject can be assigned to any regime.

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It is difficult to conduct such a trial to compare many regimes.
We must then rely on observational data

Observational analogue of a randomized trial

Interview HIV+ subjects periodically (say, every 6 months)

- ▶ record treatment modifications over the last time interval,
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Naive analysis

- ▶ Define baseline as time when CD4 first falls below 500.
- ▶ Regard subject is in:
 - ▶ Group I: if he initiates HAART when first seen to fall below 500.
 - ▶ Group II: if he starts HAART when first seen to fall below 200.
- ▶ Because treatment was not randomized we compare groups after adjusting for baseline potential confounding factors.

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PROBLEM \Rightarrow SELECTION BIAS

- ▶ Subjects not included in Group I or II can't be ignored.

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PROBLEM \Rightarrow SELECTION BIAS

- ▶ Subjects not included in Group I or II can't be ignored.
- ▶ Selection bias can be corrected using [Inverse Probability of Censoring Weighted](#) (IPCW) methods.

Inverse probability of censoring weighted estimation

Suppose we want to estimate

$$\mu^{200} = E(Y^{200})$$

μ^{200} = expected utility in a world where all subjects followed regime d^{200} (start HAART first time CD4 falls below 200).

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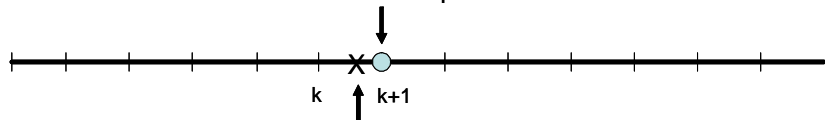
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Strategy

- ▶ Censor a subject at occasion k if he/she:
 - ▶ started HAART at occasion k prior to falling below 200 or
 - ▶ failed to start HAART at occasion k when falling below 200.
- ▶ **Redistribute** the censored subject among those still **at risk** (following regime d^{200}) who **have the same history** up to k .
- ▶ The process is repeated for $k = 0, \dots, K$.

Inverse probability of censoring weighted estimation

Subject redistributed here among all “at risk” subjects at $k+1$ with the same past from 0 to k



Subject failed to follow regime d^{200} here.
Censored at occasion k , i.e. $C^{200} = k$

Inverse probability of censoring weighted estimation

Data recorded in the cohort study

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L_k = vector of covariates measured at time k ,

A_k = HAART indicator.

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Accumulated weight through occasion k for a subject is estimated as

$$W_k^{200} = \frac{I(C^{200} > k)}{\prod_{j=1}^k \widehat{\Pr}(C^{200} > j | C^{200} > j-1, \bar{A}_{j-1}, \bar{L}_j)}$$

where C^{200} = time to censoring under regime d^{200} .

- ▶ Numerator is the indicator of following regime d^{200} through k .
- ▶ Denominator estimates the probability a subject had his observed HAART history through k .
- ▶ Usually \bar{L}_j is a high dimensional vector, so a parametrical model is assumed for the censoring probabilities.

Inverse probability of censoring weighted estimation

We estimate $\mu^{200} \equiv E(Y^{200})$ with

$$\hat{\mu}^{200} = \frac{\sum_{i=1}^n W_i^{200} Y_i}{\sum_{i=1}^n W_i^{200}}$$

where W^{200} is the accumulated weight at the end of study.

- ▶ $\hat{\mu}^{200}$ is a weighted average of the outcomes of those patients who followed regime d^{200} throughout.

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The estimator is **consistent and asymptotically normal** if:

- ▶ Model for hazard of censoring is correctly specified.
- ▶ At each time k recorded data includes all covariates used by doctors to prescribe HAART.
 - ▶ Sequential Randomization or No Unmeasured Confounders Assumption.
 - ▶ Non-testable!

Estimating the optimal regime in a candidate set

- ▶ We want to compare regimes $d^x \equiv \textit{start HAART first time CD4 falls below } x$, where $x \in \mathcal{X} = \{200, 201, \dots, 500\}$.
- ▶ In principle, we can estimate each mean $\mu^x \equiv E(Y^x)$ separately and then find \hat{x}_{opt} that maximizes $\hat{\mu}^x$.
- ▶ However, estimates $\hat{\mu}^x$ will have high variance because each regime will be followed by few subjects.
- ▶ Even in the ideal randomized trial we would also face this small cell problem.

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SOLUTION \Rightarrow parametrically model $E(Y^x)$

Estimating the optimal CD4 in a candidate set: Proposal

Assume that

$$\mu^x = E(Y^x) = h(x; \beta) \quad (1)$$

where $h(x; \beta)$ is a known smooth function of a $p \times 1$ unknown parameter β .

For example,

$$h(x; \beta) = \beta_0 + \beta_1 x + \beta_2 x^2 + \beta_3 x^3 + \beta_4 x^4 + \beta_5 x^5$$

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Given an estimate $\hat{\beta}$ of β we can find $\hat{x}_{opt} = \arg \max \left(h(x; \hat{\beta}) \right)$.

So, under model (1) the problem reduces to estimating β .

Estimating the optimal CD4 in a candidate set: Proposal

- ▶ Let $\gamma =$ number of regimes in the candidate set
 $\mathcal{X} = \{200, 201, \dots, 500\}$.
- ▶ Create an artificial data set, with each subject contributing γ observations $(W_i^{x_j}, Y_i, x_j)$, $j = 1, \dots, \gamma$.

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- ▶ Find $\hat{\beta}$ solving the weighted estimating equation

$$\mathbb{P}_n \left\{ \sum_{x \in \mathcal{X}} \frac{\partial h(x; \beta)}{\partial \beta} W^x [Y - h(x; \beta)] \right\} = 0$$

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Estimating equations can be modified to obtain estimators:

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Can allow for the possibility that optimal CD4 count depends on baseline covariates Z by considering **Parametric Marginal Structural Mean (MSM) models** of the form

$$E(Y^x | Z = z) = h_{\text{par}}(z, x; \beta)$$

For instance,

$$h_{\text{par}}(z, x; \beta) = \beta_1 + \beta_2 z + \beta_3 x + \beta_4 xz + \beta_5 x^2 + \beta_6 x^2 z$$

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$$h_{\text{par}}(z, x; \beta) = \underbrace{\beta_1 + \beta_2 z}_{q(z)} + \underbrace{\beta_3 x + \beta_4 xz + \beta_5 x^2 + \beta_6 x^2 z}_{h_{\text{sem}}(z, x; \beta)}$$

Can also consider more flexible **Semiparametric MSM models**

$$E(Y^x | Z = z) = h_{\text{sem}}(z, x; \beta) + q(z)$$

Model extensions (II)

The same approach can be used to optimize over a more complex set of candidates regimes where x is replaced by a vector (x_1, \dots, x_5) .

Example:

- ▶ Start HAART the first time that
 - ▶ CD4 falls below x_1 or
 - ▶ CD4 falls in (x_1, x_2) and current HIV RNA is greater than x_3 .
- ▶ Otherwise do not start.

The target of estimation in this approach is $(x_{1,opt}, x_{2,opt}, x_{3,opt})$.

General formulation (Summary)

- ▶ Assuming that treatment decisions are to be made at fixed times $t = 0, 1, \dots, K$.
- ▶ We considered regimes indexed by a vector $x \in \mathcal{X}$, \mathcal{X} possibly uncountable
- ▶ We developed estimators of the optimal treatment regime $x_{opt}(z)$ for subjects with baseline values $Z = z$ under:
 - ▶ Parametric Marginal Structural Mean Models for $E(Y^x|Z = z)$.
 - ▶ Semiparametric Marginal Structural Mean Models for $E(Y^x|Z = z)$.
- ▶ We established a set of assumptions for identification of $E(Y^x|Z = z)$ from the observed data distribution.
- ▶ We derived a class of consistent, doubly-robust and asymptotically normal estimators of β under each of the proposed models and the efficient estimator in the class.

Data analysis for illustrative purposes only

We applied this method to the publicly available MACS-WIHS data.

- ▶ Restricted to HIV-positive, AIDS-free participants who were antiretroviral therapy naïve by the time HAART was first available for use.
- ▶ Outcome of interest was the minimum of
 - ▶ time since baseline to death from any cause
 - ▶ time to first diagnosis of clinical AIDS
 - ▶ 7 years (five years follow-up).

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 - ▶ time since baseline to death from any cause
 - ▶ time to first diagnosis of clinical AIDS
 - ▶ 7 years (five years follow-up).
- ▶ Set of regimes $x \in [100, 400]$.
- ▶ Proportion of patients following regime d^x steadily decreased from 57% for regime d^{100} to 27% for regime d^{400} .
- ▶ We assumed a Parametric MSM polynomial model in x (5th order) with no baseline covariates.
- ▶ We obtained $\hat{x}_{opt} = 289$ cell counts/mm³ with nominal 95% CI for $x_{opt} = (266; 312)$.

Concluding remarks

- ▶ Dynamic MSM models have appealing properties
 - ▶ Easy to understand.
 - ▶ Easy to fit with standard software that allows for weighting.
 - ▶ It is possible to deal with missing outcomes (due to death for other causes or drop-out).

- ▶ We conducted simulation studies that confirmed the theoretical results.

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 - ▶ Easy to fit with standard software that allows for weighting.
 - ▶ It is possible to deal with missing outcomes (due to death for other causes or drop-out).
- ▶ We conducted simulation studies that confirmed the theoretical results.
- ▶ However... our proposal assumes that patients come to the clinic at fixed time intervals.
- ▶ This is not the realistic setting in the management of chronic diseases:
 - ▶ next visit date is decided based on patient health status and
 - ▶ patients are free to return earlier if they need to do so.

Main ideas of the talk based on:

- ▶ Orellana L.C. (2007) *Methodological challenges for the estimation of optimal dynamic treatment regimes from observational studies*. Harvard University, Dep of Biostatistics, Ph.D. Thesis.
- ▶ Robins J.M., Orellana, L., Rotnitzky A. (2008) Optimal treatment and testing strategies with possibly nonignorable observation processes. *Statistics in Medicine*, 27: 4678–4721.
- ▶ Orellana L., Rotnitzky A. and Robins J.M. Dynamic regime marginal structural mean models for estimation of optimal dynamic treatment regimes (to appear in *International Journal of Biostatistics*, 2009).