

Targeted Learning for Data Adaptive Causal Inference in Observational and Randomized Studies

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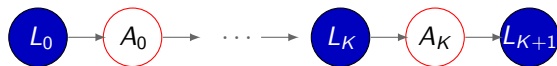
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Course Outline

- Part 1
 - Targeted Learning Overview
 - Estimation Roadmap
 - Super Learning
- Part 2
 - Targeted Minimum Loss-Based Estimation (TMLE)
- Part 3
 - TMLE for longitudinal data analysis
 - Concluding Remarks

TMLE for Longitudinal Data Analysis

- Goal: Assess Impact of Treatment at Multiple Timepoints
- Longitudinal Data (K time points)



Covariate and Outcome nodes ($L_0, \dots, L_{K+1}$)

Intervention nodes ($A_0, \dots, A_K$) indicate treatment and censoring

Challenges to Analyzing Longitudinal Data

- Common default approach
 - View as time-to-event data
 - Impose a Cox Proportional Hazards Model
- However
 - Hazard ratio may not be the most relevant target parameter
 - Cox model is misspecified
 - Cox PH model ignores informative right censoring
 - Time-dependent Cox model does not appropriately handle time-varying covariates affected by prior treatment
- Longitudinal TMLEs appropriately address these challenges

Statistical Estimation Problem

- **Data:** n i.i.d. copies
 $O = (L_0, A_0, \dots, L_K, A_K, Y = L_{K+1}) \sim P_0$
- **Statistical Model:** $\mathcal{M}$
Collection of possible probability distributions of O
- **Target Parameter:** $E(Y_{\bar{a}})$
Mean outcome under specified intervention $\bar{a} = (a_0, \dots, a_K)$
- **Mapping:** $\Psi : \mathcal{M} \rightarrow \mathbb{R}$, such that $\Psi(P_{\bar{a}}) = E(Y_{\bar{a}})$
($P_{\bar{a}}$) is post-intervention distribution identified by G-computation formula when causal assumptions are met

Note: contrasts (e.g. ATE, RR, RD) are functions of intervention-specific means

Factorization of Likelihood

- Probability distribution P_0 of O factorizes according to time-ordering as

$$\begin{aligned} P_0(O) &= \prod_{k=0}^{K+1} P_0 [L_k \mid Pa(L_k)] \prod_{k=0}^K P_0 [A_k \mid Pa(A_k)] \\ &\equiv \prod_{k=0}^{K+1} Q_{0,L_k}(O) \prod_{k=0}^K g_{0,A_k}(O) \\ &\equiv Q_0 g_0 \end{aligned}$$

where $Pa(L_k) \equiv (\bar{L}_{k-1}, \bar{Q}_{k-1})$ and $Pa(A_k) \equiv (\bar{L}_k, \bar{A}_{k-1})$ denote parents of L_k and A_k in the time-ordered sequence, respectively

- g_0 -factor represents the intervention mechanism, e.g., treatment and right-censoring mechanisms.

G-Computation Formula for Post-Intervention Distribution

- $\bar{a}_K$ is a specific treatment regime of interest
- Consider an intervention that sets $\bar{A}_K = \bar{a}_K$ in the NPSEM
- The post-intervention distribution is given by Robins' G-computation formula

$$P^a(\bar{l}) = \prod_{k=0}^{K+1} Q_{L_k}^a(\bar{l}_k),$$

where $Q_{L_k}^a(\bar{l}_k) = Q_{L_k}(l_k \mid \bar{l}_{k-1}, \bar{A}_{k-1} = \bar{a}_{k-1})$.

Statistical Target Parameter

- Let $L^a = (L_0, L_1^a, \dots, Y^a = L_{K+1}^a)$ denote the random variable with probability distribution P^a
- Our statistical target parameter is the mean of Y^a : $\Psi(P) = E_{P^a} Y^a$, where $\Psi : \mathcal{M} \rightarrow \mathbb{R}$.
 - depends on P only through $Q = Q(P)$.
 - Equivalently denoted by the mapping $\Psi : \mathcal{Q} = \{Q(P) : P \in \mathcal{M}\} \rightarrow \mathbb{R}$ so that $\psi_0 = \Psi(Q_0)$.

Alternative target parameters

- Treatment-specific mean $E_{P^d} Y^d$ defined by the G-computation formula for a dynamic treatment d
- Dose-Response
 - Projection of a true dose-response curve $(E_{P^a} Y^a : a \in \mathcal{A})$ onto a working model $\{a \rightarrow m_\beta(a) : \beta\}$.
 - Projection of the true dose-response curve $(E_{P^d} Y^d : d \in \mathcal{D})$, $\mathcal{D}$ a collection of dynamic treatment rules, onto a working model $\{d \rightarrow m_\beta(d) : \beta\}$.
 - Summary measures of conditional dose-response curves $(E_{P^d}(Y^d|V) : d \in \mathcal{D})$, conditioning on baseline covariates of interest
- Related classes of target parameters defined by history adjusted marginal structural working models for history adjusted conditional treatment-specific means
- Effects of stochastic interventions, intention to treat interventions, etc.

A Sequential Regression G-Computation Formula

- By the iterative conditional expectation rule (tower rule), we have

$$E_{P^a} Y^a = E \dots E \left[E(Y^a \mid \bar{L}_K^a) \mid L_{K-1}^a \dots \mid L_0 \right].$$

- Conditional expectation given $\bar{L}_K^a$ is equivalent to conditioning on $\bar{L}_K, \bar{A}_{K-1} = \bar{a}_{K-1}$.
- This yields the sequential regression G-computation formula
 - Compute $\bar{Q}_Y^a = E_{Q_Y^a} Y \equiv E(Y \mid \bar{L}_K, \bar{A}_K = \bar{a}_K)$
 - Given $\bar{Q}_Y^a$, next compute $\bar{Q}_{L_K}^a = E_{Q_{L_K}^a} (\bar{Q}_Y^a \mid \bar{L}_{K-1}, \bar{A}_{K-1} = \bar{a}_{K-1})$
 - Iterate over all time points
 - given $\bar{Q}_{L_{k+1}}^a$, compute $\bar{Q}_{L_k}^a = E_{Q_{L_{k+1}}^a} (\bar{Q}_{L_{k+1}}^a \mid \bar{L}_{k-1}, \bar{A}_{k-1} = \bar{a}_{k-1})$
 - Until final step, $\bar{Q}_{L_0}^a = E_{Q_{L_0}} \bar{Q}_{L_1}^a$.

TMLE for an Intervention-Specific Mean

- Iterated conditional expectations approach (*Bang and Robins, 2005*)

$$\begin{aligned}
 E(Y_{\bar{a}}) &= E\left(E\left\{\dots E\left[\underbrace{E\{Y_{\bar{a}} \mid \bar{L}_K, \bar{A}_K = \bar{a}_K\}}_{\bar{Q}_{L(K)}^a} \mid \bar{L}_{K-1}, \bar{A}_{K-1} = \bar{a}_{K-1}\right] \dots \mid L_0\right\}\right) \\
 &\quad \underbrace{\hspace{10em}}_{\bar{Q}_{L(K)}^a} \\
 &\quad \vdots \\
 &\quad \underbrace{\hspace{10em}}_{\bar{Q}_{L(1)}^a} \\
 &\quad \underbrace{\hspace{10em}}_{\bar{Q}_{L(0)}^a = \Psi(Q)}
 \end{aligned}$$

- TMLE target parameter mapping: target parameter is function of iteratively defined sequence of conditional means, $\Psi(\bar{Q}^a)$

$$\bar{Q}^a = \left(\bar{Q}_Y^a, \bar{Q}_{L(K)}^a, \dots, \bar{Q}_{L(0)}^a\right)$$

Efficient Influence Curve of Target Parameter

- Efficient influence curve representation as sum of iteratively defined scores of iteratively defined conditional means

$$D^* = \sum_{k=0}^{K+1} D_k^*$$

where

$$D_{K+1}^* = \frac{I(\bar{A}_K = \bar{a}_K)}{g_{0:K}} (Y - \bar{Q}_{K+1}^a),$$

and

$$\begin{aligned} D_k^* &= \frac{I(\bar{A}_{k-1} = \bar{a}_{k-1})}{g_{0:k-1}} \left(\bar{Q}_{L_{k+1}}^a - E_{Q_{L_k}^a} \bar{Q}_{L_{k+1}}^a \right), \\ &= \frac{I(\bar{A}_{k-1} = \bar{a}_{k-1})}{g_{0:k-1}} \left(\bar{Q}_{L_{k+1}}^a - \bar{Q}_{L_k}^a \right), k = K, \dots, 1, \end{aligned}$$

and

$$D_0^* = \bar{Q}_{L_1}^a - E_{L_0} \bar{Q}_{L_1}^a = \bar{Q}_{L_1}^a - \Psi(\bar{Q}^a).$$

$$g_{0:K} = \prod_{k=1}^K g_k$$

L-TMLE Definition

- Initial estimate of $\bar{Q}_{L_k}^a$ (Assume $Y \in [0, 1]$)
super learning, parametric regression, etc.
- Submodel and Loss Function (e.g., negative log likelihood)

$$\text{logit} \bar{Q}_{L_k}^{a,*}(\epsilon_k, g) = \text{logit} \bar{Q}_{L_k}^a + \epsilon_k \frac{1}{g_{0:k-1}}, k = K + 1, \dots, 0$$

$$\begin{aligned} \mathcal{L}_{k, \bar{Q}_{L_{k+1}}^a, g}(\bar{Q}_{L_k}^a) = \\ - \frac{I(\bar{A}_{k-1} = \bar{a}_{k-1})}{g_{0:k-1}} \left\{ \bar{Q}_{L_{k+1}}^a \log \bar{Q}_{L_k}^a + (1 - \bar{Q}_{L_{k+1}}^a) \log \{1 - \bar{Q}_{L_k}^a\} \right\} \end{aligned}$$

- Mapping

$$\Psi(\bar{Q}_n^{a,*}) = \bar{Q}_{L_0, n}^{a,*} = \frac{1}{n} \sum_{i=1}^n \bar{Q}_{1, n}^{a,*}(L_{0i})$$

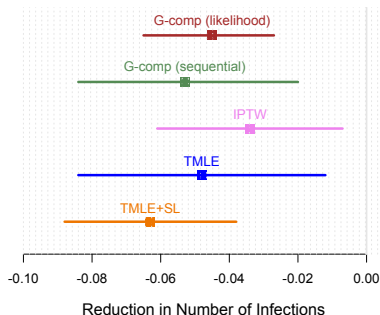
- Known bounds (e.g. rare outcome) on conditional means can be respected using logistic link.
- TMLE is double robust and asymptotically efficient if both g_0 and the conditional means $\bar{Q}_{L_k}^a$ are consistently estimated.
- Statistical inference can be based on efficient influence curve: conservative as long as g_0 is estimated well.

Example: PROBIT Study Re-Analysis

Investigate the impact of increasing duration of breastfeeding on number of GI tract infections in 1st year of life*

- Breastfeeding at time t impacts infection at time $t + 1$, which impacts decision to continue breastfeeding at $t + 2$
- TMLE + SL estimates largest effect, with variance close to that of efficient parametric G-computation estimator

**Impact of Breastfeeding for 9+ months vs. 1-2 months
on number of GI Tract Infections**



* Schnitzer, et al, 2014

Extension to Marginal Structural Models (simplified)

- Impose a MSM to smooth over areas where there is little support in the data
- Consider a working logistic MSM,
 $\text{logitm}_\beta(d, t) = \beta_1 + \beta_2 t + \beta_3 f(d, t)$
- Define target parameter as

$$\psi_0 = \underset{\beta}{\operatorname{argmin}} - E_0 \sum_{t \in \tau} \sum_{d \in D} \{Y_d(t) \log m_\beta(d, t) + (1 - Y_d(t)) \log(1 - m_\beta(d, t))\}.$$

- See Petersen, et al (2013), *ltmle* package on CRAN

Stratified TMLE for Longitudinal MSM Parameter

- Estimand $\psi_0 = \beta$ solves the equation

$$0 = E_0 \sum_{t \in \tau} \sum_{d \in D} \frac{\frac{d}{d\beta} m_\beta(d, t)}{m_\beta(1 - m_\beta)} (E_0(Y^d(t) | L_0) - m_\beta(d, t)) .$$

- Estimate $\bar{Q}_{L_0}^{d, t*}$, for each time point, t , and rule $d \in D$ using targeted iterated conditional expectations approach
- Finally, stack $\bar{Q}_{L_0}^{d, t*}$, and regress onto appropriate covariates in the model $(1, t, f(d, t))$

Notes on Targeting

- Multi-dimensional target parameter requires multi-dimensional fluctuation at each step, $\epsilon_k = (\epsilon_{1_k}, \epsilon_{2_k}, \epsilon_{3_k})$

$$\bar{Q}_{L_k}^{d*} = \bar{Q}_{L_k}^d + \epsilon_k \frac{h_1(d, t)}{g_{0:k-1}},$$

$$\text{with } h_1(d, t) = \frac{\frac{d}{d\beta} m_\beta(d, t)}{m_\beta(1 - m_\beta)}$$

- Fit ϵ using observations where $\bar{A}_{k-1} = \bar{a}_{k-1}$

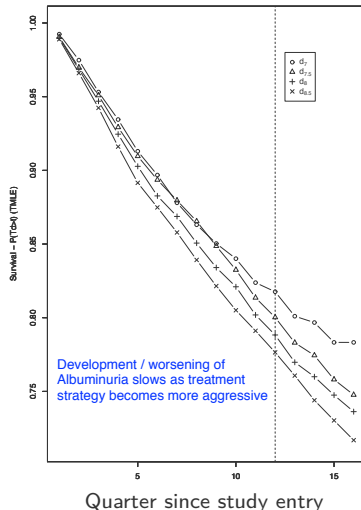
Pooled TMLE for Longitudinal MSM Parameter

- *Pooling across rules* refers to using stacking the datasets for all rules to estimate a single (multi-dimensional) ϵ_k common across all rules.
- The resulting dataset has $n \times |D|$ rows
- Pooling helps if there is sufficient support for some rules but not others
- This simultaneous targeting across all rules still solves the efficient influence curve equation $P_n D^* = 0$.
- An alternative pooled TMLE pools also over the time points t at final outcome $Y(t)$ at the targeting step. Initial estimates of $\bar{Q}_{L_k}^{t,d}$ are obtained for all k from 0 to t , across d and t , and then targeted simultaneously with a common ϵ . Updates are iterated until convergence. The dataset has $n \times |D| \times K + 1$ observations.

Example: Progression of Albuminuria in Type-2 Diabetics

- Lowering glucose levels known to prevent or slow development of Albuminuria
- Best glucose-lowering strategy is not known
- Four candidate strategies, d_θ , intensify treatment when patient's A1c level reaches $\theta = 7\%$, 7.5% , 8% , or 8.5%
- HMO Research Network EHR data
7 sites, $n = 51,179$
- Longitudinal TMLE was used to evaluate a set of increasingly aggressive dynamic strategies for lowering glucose levels

Counterfactual Survival Curves
(L-TMLE + SL)



Concluding Remarks

- TMLE provides a template for construction of efficient substitution estimators
- Three basic requirements
 - Loss function
 - Submodel for fluctuation so that its loss-based score spans the efficient influence curve
 - Procedure for iteratively minimizing the empirical risk along the fluctuation model through a current estimator

Concluding Remarks

- All TMLEs are double robust and efficient, but may have different finite sample performance
- Sequential regression
 - particularly effective representation of post-intervention distribution, and thereby causal effects
 - Estimate only smallest portion of Q needed for evaluating the parameter

Core Concepts in Targeted Learning

- Translate a scientific question and background knowledge into a formal causal model, target causal quantity, statistical model and statistical target parameter
- Target statistical parameter has a causal interpretation when assumptions are met, variable importance otherwise
- SL + TMLE for estimation
 - Optimal bias/variance trade-off for target parameter
 - Loss-based estimation using cross-validation
 - Flexible fitting of relevant components of the likelihood
 - Double robust to mitigate misspecification bias
 - Substitution estimator that respects domain knowledge

Selected Resources

- <http://www.targetedlearningbook.com>
- M.J. van der Laan and S. Rose. *Targeted Learning: Prediction and Causal Inference for Observational and Experimental Data*. Springer, New York, 2011.
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Software

- E.C. Polley, SuperLearner: Super Learner in Prediction, v2.0-19, <http://cran.r-project.org/web/packages/SuperLearner>, 2016.
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