

Day 3, Lecture 2

Longitudinal TMLE (LTMLE)

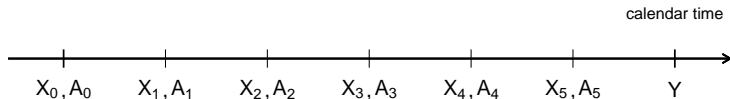
# Overview

1. Identification formulas
2. Targeting algorithm
  - ▶ Practical
3. ltmle software package
  - ▶ Practical
4. A note on right-censoring

# Longitudinal data structure

## Longitudinal data structure:

- ▶  $O = (X_0, A_0, X_1, A_1, \dots, X_K, A_K, Y = X_{K+1}) \in (\mathbb{R}^d \times \{0, 1\})^K \times \{0, 1\}$
- ▶ Covariates  $X = (X_0, X_1, \dots, X_K)$  change over time
- ▶ Treatment decisions  $A = (A_0, A_1, \dots, A_K)$  are updated over time
- ▶ Covariates and treatment decisions interact in complex ways



## Longitudinal data structure

Statistical model  $\mathcal{M}$  for  $P_0$  contains possible distributions  $P$  for the observed data  $O$ ; we assume again that  $\mathcal{M}$  is a nonparametric model.

Factorization of the density  $p$  of  $P \in \mathcal{M}$ :

$$p(o) = \mu_{X_0}(x_0) \pi_{A_0}(a \mid x_0) \prod_{k=1}^K \mu_{X_k}(x_k \mid \bar{x}_{k-1}, \bar{a}_{k-1}) \pi_{A_k}(a_k \mid \bar{x}_k, \bar{a}_{k-1}) \\ \times \mu_Y(y \mid \bar{x}_K, \bar{a}_K)$$

- ▶  $\mu_{X_0}$  is the marginal density of baseline covariates.
- ▶  $\pi_{A_0}$  is the marginal density of treatment at baseline.
- ▶  $\mu_{X_k}(x_k \mid \bar{x}_{k-1}, \bar{a}_{k-1})$  is the conditional density of  $X_k$  given the histories  $\bar{X}_{k-1} = \bar{x}_{k-1}, \bar{A}_{k-1} = \bar{a}_{k-1}$ ,  $k = 1, \dots, K$ .
- ▶  $\pi_{A_k}(a_k \mid \bar{x}_k, \bar{a}_{k-1})$  is the conditional density of  $A_k$  given the histories  $\bar{X}_k = \bar{x}_k, \bar{A}_{k-1} = \bar{a}_{k-1}$ ,  $k = 1, \dots, K$ .
- ▶  $\mu_Y(y \mid \bar{x}_K, \bar{a}_K)$  is the conditional density of  $Y$  given the histories  $\bar{X}_K = \bar{x}_K, \bar{A}_K = \bar{a}_K$ .

## Longitudinal data structure

Factorization of density allows us to write the expectation under  $P$  in terms of iterated integrals (Fubini's theorem):

$$\begin{aligned}\mathbb{E}_P[Y] &= \int_{\mathcal{O}} yp(o) d\nu(o) \\ &= \int_{\mathbb{R}^d} \sum_{a_0=0,1} \cdots \int_{\mathbb{R}^d} \sum_{a_K=0,1} \sum_{y=0,1} y \mu_Y(y \mid \bar{x}_K, \bar{a}_K) \\ &\quad \pi_K(a_K \mid \bar{x}_K, \bar{a}_K) \mu_{X_K}(x_K \mid \bar{x}_{K-1}, \bar{a}_{K-1}) d\nu_{X_K}(x_K) \\ &\quad \cdots \pi_0(a_0 \mid x_0) \mu_{X_0}(x_0) d\nu_{X_0}(x_0),\end{aligned}$$

for  $P \in \mathcal{M}$ .

# Identification

We want to identify the treatment-specific mean outcome:

$$\mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}]$$

in terms of the observed data distribution

using the assumptions:

$$Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*} = Y \quad \text{if } A_k = a_k^* \text{ for } k = 0, 1, \dots, K$$

$$Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*} \perp\!\!\!\perp A_k \mid \bar{X}_k, \bar{A}_{k-1}, \quad \text{for } k = 0, 1, \dots, K$$

$$\prod_{k=0}^K \frac{1\{A_k = a_k^*\}}{P(A_k = a_k^* \mid \bar{X}_k, \bar{A}_{k-1})} < \infty, \quad \text{for } k = 0, 1, \dots, K$$

## Identification: g-formula

The claim is that:

$$\begin{aligned} \mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}] \\ = \int_{\mathbb{R}^d} \cdots \int_{\mathbb{R}^d} \sum_{y=0,1} y \mu_Y(y \mid \bar{x}_K, \bar{a}_K^*) \\ \mu_{X_K}(x_K \mid \bar{x}_{K-1}, \bar{a}_{K-1}^*) d\nu_{X_K}(x_K) \cdots \mu_{X_0}(x_0) d\nu_{X_0}(x_0) \end{aligned}$$

## Identification: g-formula

To show the claim from the previous slide, **start from the right hand side**:

1. By consistency, replace  $Y$  by  $Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}$  in the innermost integral:

$$\begin{aligned} \sum_{y=0,1} y \mu_Y(y \mid \bar{x}_K, \bar{a}_K^*) &= \mathbb{E}_P[Y \mid \bar{X}_K = \bar{x}_K, \bar{A}_K = \bar{a}_K^*] \\ &= \mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*} \mid \bar{X}_K = \bar{x}_K, \bar{A}_K = \bar{a}_K^*] \end{aligned} \quad (1)$$

2. Drop the last conditioning variable  $A_K^* = a_K^*$  from the conditioning set by exchangeability, and then integrate out over  $L_K$ :

$$\begin{aligned} \int_{\mathbb{R}^d} \sum_{y=0,1} y \mu_Y(y \mid \bar{x}_K, \bar{a}_K^*) \mu_{X_K}(x_K \mid \bar{x}_{K-1}, \bar{a}_{K-1}^*) d\nu_{X_K}(x_K) \\ = \int_{\mathbb{R}^d} \mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*} \mid \bar{X}_K = \bar{x}_K, \bar{A}_{K-1}^* = \bar{a}_{K-1}^*] \mu_{X_K}(x_K \mid \bar{x}_{K-1}, \bar{a}_{K-1}^*) d\nu_{X_K}(x_K) \\ = \mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*} \mid \bar{X}_{K-1} = \bar{x}_{K-1}, \bar{A}_{K-1}^* = \bar{a}_{K-1}^*] \end{aligned} \quad (2)$$

3. Note that (2) is the same expression as (1), with  $K$  replaced by  $K - 1$ .
4. Repeat 2. another  $K - 1$  times which in the end **gives the left hand side** from the previous slide.



## Identification: IP-weighting

We have that:

$$\begin{aligned} & \mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}] \\ &= \int_{\mathbb{R}^d} \cdots \int_{\mathbb{R}^d} \sum_{y=0,1} y \mu_Y(y \mid \bar{x}_K, \bar{a}_K^*) \\ & \quad \mu_{X_K}(x_K \mid \bar{x}_{K-1}, \bar{a}_{K-1}^*) d\nu_{X_K}(x_K) \cdots \mu_{X_0}(x_0) d\nu_{X_0}(x_0) \end{aligned}$$

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## Identification: IP-weighting

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## Identification: IP-weighting

We have that:

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## Identification: g-formula & iterated expectations

The g-formula:

$$\begin{aligned} \mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}] \\ = \int_{\mathbb{R}^d} \cdots \int_{\mathbb{R}^d} \sum_{y=0,1} y \mu_Y(y \mid \bar{x}_K, \bar{a}_K^*) \\ \mu_{X_K}(x_K \mid \bar{x}_{K-1}, \bar{a}_{K-1}^*) d\nu_{X_K}(x_K) \cdots \mu_{X_0}(x_0) d\nu_{X_0}(x_0) \end{aligned}$$

can also be written as a sequence of iterated conditional expectations.

## Identification: g-formula & iterated expectations

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$$\begin{aligned}
 & \mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}] \\
 &= \int_{\mathbb{R}^d} \cdots \int_{\mathbb{R}^d} \sum_{y=0,1} y \mu_Y(y \mid \bar{x}_K, \bar{a}_K^*) \\
 &\quad \mu_{X_K}(x_K \mid \bar{x}_{K-1}, \bar{a}_{K-1}^*) d\nu_{X_K}(x_K) \cdots \mu_{X_0}(x_0) d\nu_{X_0}(x_0) \\
 &= \int_{\mathbb{R}^d} \sum_{a_0=0,1} \cdots \int_{\mathbb{R}^d} \sum_{a_K=0,1} \mathbb{E}[Y \mid \bar{X}_K = \bar{x}_K, \bar{A}_K = \bar{a}_K] \\
 &\quad 1\{a_K = a_K^*\} \mu_{X_K}(x_K \mid \bar{x}_{K-1}, \bar{a}_{K-1}^*) d\nu_{X_K}(x_K) \\
 &\quad \cdots 1\{a_0 = a_0^*\} \mu_{X_0}(x_0) d\nu_{X_0}(x_0)
 \end{aligned}$$

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 &= \int_{\mathbb{R}^d} \sum_{a_0=0,1} \cdots \int_{\mathbb{R}^d} \sum_{a_K=0,1} \mathbb{E}[Y \mid \bar{X}_K = \bar{x}_K, \bar{A}_K = \bar{a}_K] \\
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Define:

$$\bar{Q}_{K+1}(\bar{x}_K, \bar{a}_K) = \mathbb{E}_P[Y \mid \bar{X}_K = \bar{x}_K, \bar{A}_K = \bar{a}_K]$$

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$$\begin{aligned}
 & \mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}] \\
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 &= \int_{\mathbb{R}^d} \sum_{a_0=0,1} \cdots \int_{\mathbb{R}^d} \sum_{a_K=0,1} \bar{Q}_{K+1}(\bar{x}_K, a_K^*, \bar{a}_{K-1}) \\
 &\quad 1\{a_K = a_K^*\} \mu_{X_K}(x_K \mid \bar{x}_{K-1}, \bar{a}_{K-1}) d\nu_{X_K}(x_K) \\
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 &= \int_{\mathbb{R}^d} \sum_{a_0=0,1} \cdots \int_{\mathbb{R}^d} \bar{Q}_{K+1}(\bar{x}_K, a_K^*, \bar{a}_{K-1}) \\
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 &\quad \cdots 1\{a_0 = a_0^*\} \mu_{X_0}(x_0) d\nu_{X_0}(x_0) \\
 &= \int_{\mathbb{R}^d} \sum_{a_0=0,1} \cdots \int_{\mathbb{R}^d} \sum_{a_{K-1}=0,1} \bar{Q}_K(\bar{x}_{K-1}, \bar{a}_{K-1}) \\
 &\quad 1\{a_{K-1} = a_{K-1}^*\} \mu_{X_{K-1}}(x_{K-1} \mid \bar{x}_{K-2}, \bar{a}_{K-2}) d\nu_{X_{K-1}}(x_{K-1}) \\
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## Identification: g-formula & iterated expectations

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 & \mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}] \\
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 &= \int_{\mathbb{R}^d} \sum_{a_0=0,1} \cdots \int_{\mathbb{R}^d} \bar{Q}_{K+1}(\bar{x}_K, a_K^*, \bar{a}_{K-1}) \\
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 &= \int_{\mathbb{R}^d} \sum_{a_0=0,1} \cdots \int_{\mathbb{R}^d} \bar{Q}_K(\bar{x}_{K-1}, a_{K-1}^*, \bar{a}_{K-2}) \\
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$$\begin{aligned}
 \bar{Q}_{K+1}(\bar{x}_K, \bar{a}_K) &= \mathbb{E}_P[Y \mid \bar{X}_K = \bar{x}_K, \bar{A}_K = \bar{a}_K] \\
 \bar{Q}_K(\bar{x}_{K-1}, \bar{a}_{K-1}) &= \mathbb{E}_P[\bar{Q}_{K+1}(\bar{x}_K, a_K^*, \bar{a}_{K-1}) \mid \bar{X}_{K-1} = \bar{x}_{K-1}, \bar{A}_{K-1} = \bar{a}_{K-1}]
 \end{aligned}$$

## Identification: g-formula & iterated expectations

$$\begin{aligned}
 & \mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}] \\
 &= \int_{\mathbb{R}^d} \cdots \int_{\mathbb{R}^d} \sum_{y=0,1} y \mu_Y(y \mid \bar{x}_K, \bar{a}_K^*) \\
 &\quad \mu_{X_K}(x_K \mid \bar{x}_{K-1}, \bar{a}_{K-1}^*) d\nu_{X_K}(x_K) \cdots \mu_{X_0}(x_0) d\nu_{X_0}(x_0) \\
 &= \int_{\mathbb{R}^d} \sum_{a_0=0,1} \cdots \int_{\mathbb{R}^d} \bar{Q}_{K+1}(\bar{x}_K, a_K^*, \bar{a}_{K-1}) \\
 &\quad \mu_{X_K}(x_K \mid \bar{x}_{K-1}, \bar{a}_{K-1}) d\nu_{X_K}(x_K) \\
 &\quad \cdots 1\{a_0 = a_0^*\} \mu_{X_0}(x_0) d\nu_{X_0}(x_0) \\
 &= \int_{\mathbb{R}^d} \sum_{a_0=0,1} \cdots \int_{\mathbb{R}^d} \bar{Q}_K(\bar{x}_{K-1}, a_{K-1}^*, \bar{a}_{K-2}) \\
 &\quad \mu_{X_{K-1}}(x_{K-1} \mid \bar{x}_{K-2}, \bar{a}_{K-2}) d\nu_{X_{K-1}}(x_{K-1}) \\
 &\quad \cdots 1\{a_0 = a_0^*\} \mu_{X_0}(x_0) d\nu_{X_0}(x_0)
 \end{aligned}$$

Define:

$$\begin{aligned}
 \bar{Q}_{K+1}(\bar{x}_K, \bar{a}_K) &= \mathbb{E}_P[Y \mid \bar{X}_K = \bar{x}_K, \bar{A}_K = \bar{a}_K] \\
 \bar{Q}_K(\bar{x}_{K-1}, \bar{a}_{K-1}) &= \mathbb{E}_P[\bar{Q}_{K+1}(\bar{x}_K, a_K^*, \bar{a}_{K-1}) \mid \bar{X}_{K-1} = \bar{x}_{K-1}, \bar{A}_{K-1} = \bar{a}_{K-1}]
 \end{aligned}$$

## Identification: g-formula & iterated expectations

Full steps that represent  $\mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}]$  as a sequence of iterated conditional expectations:

$$\bar{Q}_{K+1}(\bar{x}_K, \bar{a}_K) = \mathbb{E}_P[Y \mid \bar{X}_K = \bar{x}_K, \bar{A}_K = \bar{a}_K]$$

$$\bar{Q}_K(\bar{x}_{K-1}, \bar{a}_{K-1}) = \mathbb{E}_P[\bar{Q}_{K+1}(\bar{X}_K, a_K^*, \bar{A}_{K-1}) \mid \bar{X}_{K-1} = \bar{x}_{K-1}, \bar{A}_{K-1} = \bar{a}_{K-1}]$$

$$\vdots$$

$$\bar{Q}_k(\bar{x}_{k-1}, \bar{a}_{k-1}) = \mathbb{E}_P[\bar{Q}_{k+1}(\bar{X}_k, a_k^*, \bar{A}_{k-1}) \mid \bar{X}_{k-1} = \bar{x}_{k-1}, \bar{A}_{k-1} = \bar{a}_{k-1}]$$

$$\vdots$$

$$\bar{Q}_2(\bar{x}_1, \bar{a}_1) = \mathbb{E}_P[\bar{Q}_3(\bar{X}_2, a_2^*, \bar{A}_1) \mid \bar{X}_1 = \bar{x}_1, \bar{A}_1 = \bar{a}_1]$$

$$\bar{Q}_1(x_0, a_0) = \mathbb{E}_P[\bar{Q}_2(\bar{X}_1, a_1^*, A_0) \mid X_0 = x_0, A_0 = a_0]$$

$$\mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}] = \mathbb{E}_P[\bar{Q}_1(x_0, a_0^*)].$$



# Identification (summary)

## 1. IP-weighting:

$$\mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}] = \mathbb{E}_P\left[\frac{\prod_{k=0}^K 1\{A_k = a_k^*\}}{\prod_{k=0}^K \pi_{A_k}(a_k^* | \bar{X}_k, \bar{A}_{k-1})} Y\right]$$

## 2. Sequence of iterated conditional expectations:

$$\bar{Q}_{K+1}(\bar{x}_K, \bar{a}_K) = \mathbb{E}_P[Y | \bar{X}_K = \bar{x}_K, \bar{A}_K = \bar{a}_K]$$

and iteratively for  $k = K, K-1, \dots, 1$ ,

$$\bar{Q}_k(\bar{x}_{k-1}, \bar{a}_{k-1}) = \mathbb{E}_P[\bar{Q}_{k+1}(\bar{X}_k, a_k^*, \bar{A}_{k-1}) | \bar{X}_{k-1} = \bar{x}_{k-1}, \bar{A}_{k-1} = \bar{a}_{k-1}]$$

so that

$$\mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}] = \mathbb{E}_P[\bar{Q}_1(X_0, a_0^*)].$$

# Longitudinal targeting

(For the representation in terms of iterated conditional expectations)

# Targeting algorithm

Recall —

TMLE is a two-step procedure:

Step 1 Construct initial estimator  $\hat{P}_n$  for  $P$ .

Step 2 Update the estimator  $\hat{P}_n \mapsto \hat{P}_n^*$  such that  $\hat{P}_n^*$  solves the efficient influence curve equation.

Step 1 = "initial estimation step"

Step 2 = "targeting step"

# Targeting algorithm

Recall — for the ATE:

TMLE is a two-step procedure:

Step 1 Construct initial estimators  $\hat{f}_n, \hat{\pi}_n$  for  $f, \pi$ .

Step 2 Update the estimator  $\hat{f}_n \mapsto \hat{f}_n^*$  for  $f$  such that  $\hat{f}_n^*$  for the fixed  $\hat{\pi}_n$  solves the efficient influence curve equation.

Step 1 = "initial estimation step"

Step 2 = "targeting step"

## Targeting algorithm

The relevant part of  $P$  needed to evaluate our target parameter:

$$\mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}] = \tilde{\Psi}(\bar{Q}),$$

with  $\bar{Q} = (\bar{Q}_k)_{1 \leq k \leq K+1}$ .

Starting backwards from the last time-point:

$$\bar{Q}_{K+1}(\bar{x}_K, \bar{a}_K) = \mathbb{E}_P[Y \mid \bar{X}_K = \bar{x}_K, \bar{A}_K = \bar{a}_K]$$

and iteratively for  $k = K, K-1, \dots, 1$ ,

$$\bar{Q}_k(\bar{x}_{k-1}, \bar{a}_{k-1}) = \mathbb{E}_P[\bar{Q}_{k+1}(\bar{X}_k, a_k^*, \bar{A}_{k-1}) \mid \bar{X}_{k-1} = \bar{x}_{k-1}, \bar{A}_{k-1} = \bar{a}_{k-1}]$$

so that

$$\mathbb{E}_P[Y^{A_0=a_0^*, A_1=a_1^*, \dots, A_K=a_K^*}] = \mathbb{E}_P[\bar{Q}_1(X_0, a_0^*)] = \tilde{\Psi}(\bar{Q}).$$

# Targeting algorithm

We need the **efficient influence function**:

- ▶ Tells us what we need to estimate (to construct TMLE)
- ▶ Guides the construction of the targeting step

**Construction of the targeting step** for a given target parameter  $\Psi : \mathcal{M} \rightarrow \mathbb{R}$  with efficient influence function  $\phi^*(P)$  involves:

(i) A parametric submodel  $\{\bar{Q}_\varepsilon : \varepsilon \in \mathbb{R}\} \subset \mathcal{M}$

(ii) A loss function  $(O, \bar{Q}) \mapsto \mathcal{L}(\bar{Q})(O)$

such that: (1)  $\bar{Q}_{\varepsilon=0} = \bar{Q}$ , and, (2)  $\left. \frac{d}{d\varepsilon} \right|_{\varepsilon=0} \mathcal{L}(\bar{Q}_\varepsilon)(O) = \phi^*(P)(O)$

## Targeting algorithm

The efficient influence function is given by:

$$\begin{aligned}\phi^*(P)(O) &= \tilde{\phi}^*(\bar{Q}, \pi)(O) \\ &= \sum_{k=1}^{K+1} \left( \prod_{l=0}^{k-1} \frac{1\{A_l = a_l^*\}}{\pi_{A_l}(a_l^* \mid \bar{x}_l, \bar{a}_{l-1}^*)} \right) \{ \bar{Q}_{k+1}(\bar{X}_k, a_k^*, \bar{A}_{k-1}) - \bar{Q}_k(\bar{X}_{k-1}, \bar{A}_{k-1}) \} \\ &\quad + \bar{Q}_1(X_0, a_0^*) - \Psi(P) \\ &\text{(with } \bar{Q}_{K+2} := Y)\end{aligned}$$

# Targeting algorithm

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- ▶ Need initial estimators for:  $\pi = (\pi_{A_k})_{0 \leq k \leq K}$ ,  $\bar{Q} = (\bar{Q}_k)_{1 \leq k \leq K+1}$



# Targeting algorithm

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(with  $\bar{Q}_{K+2} := Y$ )

- ▶ Need initial estimators for:  $\pi = (\pi_{A_k})_{0 \leq k \leq K}$ ,  $\bar{Q} = (\bar{Q}_k)_{1 \leq k \leq K+1}$
- ▶ Submodel and loss function **for each  $\bar{Q}_k$  in turn** to solve the *k-specific part of the efficient influence curve equation*,

$$\tilde{\phi}_k^*(\bar{Q}, \pi)(O) = \left( \prod_{l=0}^{k-1} \frac{1\{A_l = a_l^*\}}{\pi_{A_l}(a_l^* \mid \bar{X}_l, \bar{a}_{l-1}^*)} \right) \{ \bar{Q}_{k+1}(\bar{X}_k, a_k^*, \bar{A}_{k-1}) - \bar{Q}_k(\bar{X}_{k-1}, \bar{A}_{k-1}) \}$$

## Targeting algorithm

We construct a submodel  $\bar{Q}_{k,\varepsilon}$  through a given  $\bar{Q}_k$  and a loss function  $(O, \bar{Q}_k) \mapsto \mathcal{L}_{\bar{Q}_{k+1}}(\bar{Q}_k)(O)$  such that

$$(1) \quad \bar{Q}_{k,\varepsilon=0} = \bar{Q}_k, \quad \text{and,} \quad (2) \quad \left. \frac{d}{d\varepsilon} \right|_{\varepsilon=0} \mathcal{L}_{\bar{Q}_{k+1}}(\bar{Q}_{k,\varepsilon})(O) = \phi_k^*(\bar{Q}, \pi)(O)$$

*Note that the loss function is indexed by  $\bar{Q}_{k+1}$ .*

# Targeting algorithm

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$$(1) \quad \bar{Q}_{k,\varepsilon=0} = \bar{Q}_k, \quad \text{and}, \quad (2) \quad \left. \frac{d}{d\varepsilon} \right|_{\varepsilon=0} \mathcal{L}_{\bar{Q}_{k+1}}(\bar{Q}_{k,\varepsilon})(O) = \phi_k^*(\bar{Q}, \pi)(O)$$

*Note that the loss function is indexed by  $\bar{Q}_{k+1}$ .*

We define:

$$\mathcal{L}_{\bar{Q}_{k+1}}(\bar{Q}_k) = -(\bar{Q}_{k+1} \log(\bar{Q}_k) + (1 - \bar{Q}_{k+1}) \log(1 - \bar{Q}_k))$$

and,

$$\bar{Q}_{k,\varepsilon}(O) = \text{expit}(\text{logit}(\bar{Q}_k(\bar{X}_{k-1}, \bar{A}_{k-1})) + \varepsilon H_k(\pi)(O))$$

with the "clever covariate":  $H_k(\pi)(O) := \prod_{l=0}^{k-1} \frac{1\{A_l = a_l^*\}}{\pi_{A_l}(a_l^* \mid \bar{x}_l, \bar{a}_{l-1}^*)}$

# Targeting algorithm

Another valid choice of loss function and submodel would be:

We define:

$$\mathcal{L}_{\bar{Q}_{k+1}}(\bar{Q}_k) = -H_k(\pi)(O) (\bar{Q}_{k+1} \log(\bar{Q}_k) + (1 - \bar{Q}_{k+1}) \log(1 - \bar{Q}_k))$$

and,

$$\bar{Q}_{k,\varepsilon}(O) = \text{expit}(\text{logit}(\bar{Q}_k(\bar{X}_{k-1}, \bar{A}_{k-1})) + \varepsilon)$$

using the "clever covariate"  $H_k(\pi)(O) := \prod_{l=0}^{k-1} \frac{1\{A_l = a_l^*\}}{\pi_{A_l}(a_l^* \mid \bar{X}_l, \bar{a}_{l-1}^*)}$

as a weight.

## Targeting algorithm

The targeting step becomes a targeting algorithm that proceeds iteratively along the sequence of iterated conditional expectations, *starting from the last time-point*.

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- ▶ We start by targeting an estimator  $\hat{Q}_{K+1}$
- ▶ We set  $\hat{Q}_{K+2} := Y$
- ▶ Regress  $Y$  on  $\bar{X}_K, \bar{A}_K$  and obtain the predictions  $\hat{Q}_{K+1}(\bar{X}_k, \bar{A}_k)$



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- ▶ Regress  $\hat{Q}_{K+2}$  on the clever covariate  $H_{K+1}(\hat{\pi})(O)$  and offset  $\text{logit}\{\hat{Q}_{K+1}(\bar{X}_K, \bar{A}_K)\}$  to obtain estimate  $\hat{\varepsilon}_n$  for  $\varepsilon$

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- ▶ Update according to the submodel:  $\hat{Q}_{K+1}^*(\bar{X}_k, a_k^*, \bar{A}_{k-1}) := \text{expit}(\text{logit}\{\hat{Q}_{K+1}(\bar{X}_k, a_k^*, \bar{A}_{k-1})\} + \hat{\varepsilon}_n H_{K+1}(\hat{\pi})(O))$

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We then solve:

$$\frac{1}{n} \sum_{i=1}^n H_{K+1}(\hat{\pi})(O_i) \{ \hat{Q}_{K+2} - \hat{Q}_{K+1}^*(\bar{X}_{K,i}, \bar{A}_{K,i}) \} = 0$$

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- ▶ From the  $(k + 1)$ th step, we have targeted estimator  $\hat{Q}_{k+1}^*(\bar{X}_k, \bar{A}_k)$

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- ▶ From the  $(k + 1)$ th step, we have targeted estimator  $\hat{Q}_{k+1}^*(\bar{X}_k, \bar{A}_k)$
- ▶ Regress  $\hat{Q}_{k+1}(\bar{X}_k, a_k^*, \bar{A}_{k-1})$  on  $\bar{X}_{k-1}, \bar{A}_{k-1}$  and denote the prediction by  $\hat{Q}_k(\bar{X}_{k-1}, \bar{A}_{k-1})$

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The targeting step becomes a targeting algorithm that proceeds iteratively along the sequence of iterated conditional expectations, *starting from the last time-point*.

- ▶ Proceed iteratively from  $k + 1$  to  $k$
- ▶ From the  $(k + 1)$ th step, we have targeted estimator  $\hat{Q}_{k+1}^*(\bar{X}_k, \bar{A}_k)$
- ▶ Regress  $\hat{Q}_{k+1}(\bar{X}_k, a_k^*, \bar{A}_{k-1})$  on  $\bar{X}_{k-1}, \bar{A}_{k-1}$  and denote the prediction by  $\hat{Q}_k(\bar{X}_{k-1}, \bar{A}_{k-1})$
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- ▶ Update according to the submodel:  
$$\hat{Q}_k^* := \text{expit}(\text{logit}(\hat{Q}_k(\bar{X}_{k-1}, \bar{A}_{k-1})) + \hat{\varepsilon}_n H_k(\hat{\pi})(O))$$

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The targeting step becomes a targeting algorithm that proceeds iteratively along the sequence of iterated conditional expectations, *starting from the last time-point*.

- ▶ Proceed iteratively from  $k + 1$  to  $k$
- ▶ From the  $(k + 1)$ th step, we have targeted estimator  $\hat{Q}_{k+1}^*(\bar{X}_k, \bar{A}_k)$
- ▶ Regress  $\hat{Q}_{k+1}(\bar{X}_k, a_k^*, \bar{A}_{k-1})$  on  $\bar{X}_{k-1}, \bar{A}_{k-1}$  and denote the prediction by  $\hat{Q}_k(\bar{X}_{k-1}, \bar{A}_{k-1})$
- ▶ Regress  $\hat{Q}_{k+1}(\bar{X}_k, a_k^*, \bar{A}_{k-1})$  on the clever covariate  $H_k(\hat{\pi})(O)$  and offset  $\text{logit}(\hat{Q}_k(\bar{X}_{k-1}, \bar{A}_{k-1}))$  to obtain estimate  $\hat{\varepsilon}_n$  for  $\varepsilon$
- ▶ Update according to the submodel:  
$$\hat{Q}_k^* := \text{expit}(\text{logit}(\hat{Q}_k(\bar{X}_{k-1}, \bar{A}_{k-1})) + \hat{\varepsilon}_n H_k(\hat{\pi})(O))$$

We then solve:

$$\frac{1}{n} \sum_{i=1}^n H_k(\hat{\pi})(O_i) \{ \hat{Q}_{k+1}^*(\bar{X}_{k,i}, a_k^*, \bar{A}_{k-1,i}) - \hat{Q}_k^*(\bar{X}_{k-1,i}, \bar{A}_{k-1,i}) \} = 0$$

## Targeting algorithm

The targeting step becomes a targeting algorithm that proceeds iteratively along the sequence of iterated conditional expectations, *starting from the last time-point*.

This procedure gives a sequence of updated estimators

$\hat{\hat{Q}}^* = (\hat{\hat{Q}}_{K+1}^*, \hat{\hat{Q}}_K^*, \dots, \hat{\hat{Q}}_1^*)$  that solves the efficient influence curve equation:

$$\frac{1}{n} \sum_{i=1}^n \sum_{k=1}^{K+1} \left( \prod_{l=0}^{k-1} \frac{1\{A_{l,i} = a_l^*\}}{\hat{\pi}_{A_l}(a_l^* | \bar{X}_{l,i}, \bar{a}_{l-1}^*)} \right) \{ \hat{\hat{Q}}_{k+1}^*(\bar{X}_{k,i}, a_k^*, \bar{A}_{k-1,i}) \\ - \hat{\hat{Q}}_k^*(\bar{X}_{k-1}, \bar{A}_{k-1}) \} + \hat{\hat{Q}}_1^*(X_0, a_0^*) - \Psi(\hat{\hat{Q}}^*)$$

where

$$\tilde{\Psi}(\hat{\hat{Q}}^*) = \frac{1}{n} \sum_{i=1}^n \hat{\hat{Q}}_1^*(X_{0,i}, a_0^*).$$

# Practical

## Implementing the targeting step

- ▶ Read the first page of **day3\_practical2.pdf** and then go through the steps of **Task 1**.

ltmle software package

# ltmle software package

See also: [https:](https://cran.r-project.org/web/packages/ltmle/ltmle.pdf)

[//cran.r-project.org/web/packages/ltmle/ltmle.pdf](https://cran.r-project.org/web/packages/ltmle/ltmle.pdf)

```
install.packages(ltmle)  
library(ltmle)
```

# ltmle software package

Some useful arguments to know:

```
ltmle(data,  
      Anodes, Lnodes, Cnodes, Ynodes,  
      abar, rule,  
      Qform, gform,  
      SL.library,  
      gbounds,  
      ...)
```

# ltmle software package

- ▶ data: data frame

NB the order of columns correspond to the order of variables.

- ▶ Anodes: names of treatment variables
- ▶ Lnodes: names of covariates
- ▶ Cnodes: names of censoring variables
- ▶ Ynodes: names of outcome variables

NB All variables except baseline covariates must be specified in Anodes, Lnodes, Cnodes or Ynodes.



## ltmle software package

For the dataset simulated in Lecture 1:

	X0.1	X0.2	X0.3	A0	X1.1	X1.2	A1	Y
1:	1.2915853	-0.29726781	0	0	1	0	0	1
2:	0.8407983	-0.48032616	0	0	0	1	0	0
3:	1.8633391	-0.50152323	0	1	1	1	0	1
4:	-1.6855569	-1.57948481	0	1	0	0	1	1
5:	-1.7854517	0.04438914	0	0	1	1	0	1
6:	0.2999129	-1.41470763	0	1	0	1	0	0

```
Anodes = paste0("A",0:1)
Lnodes = c(paste0("X0.", 1:3), paste0("X1.", 1:2))
Ynodes = "Y"
```

The ordering of columns dictates the temporal ordering of variables:

$$\left\{ \begin{matrix} X_{0,1} \\ X_{0,2} \\ X_{0,3} \end{matrix} \right\} \rightarrow A_0 \rightarrow \left\{ \begin{matrix} X_{1,1} \\ X_{1,2} \end{matrix} \right\} \rightarrow A \rightarrow Y$$

# ltmle software package

## Specifying interventions:

- ▶ `abar`: a binary vector of treatment assignments of length = `length(Anodes)` **or** a list of two elements to contrast treatment regimes
  - ▶ to specify *static treatment regimes*
- ▶ `rule`: a function that can be applied to each row of the data to return a binary vector of treatment assignments of length = `length(Anodes)`
  - ▶ to specify *dynamic treatment regimes*

# ltmle software package

For example:

```
summary(ltmle(sim.data,  
             Anodes=paste0("A",0:1),  
             Lnodes=c(paste0("X0.", 1:3), paste0("X1.", 1:2)),  
             Ynodes="Y",  
             abar=list(treatment=c(1,1), control=c(0,0))))
```

[output next slide]

# ltmle software package

## Treatment Estimate:

Parameter Estimate: 0.3029  
Estimated Std Err: 0.086738  
p-value: 0.00047919  
95% Conf Interval: (0.1329, 0.4729)

## Control Estimate:

Parameter Estimate: 0.36688  
Estimated Std Err: 0.015419  
p-value:  $<2e-16$   
95% Conf Interval: (0.33666, 0.3971)

## Additive Treatment Effect:

Parameter Estimate: -0.063978  
Estimated Std Err: 0.088095  
p-value: 0.46769  
95% Conf Interval: (-0.23664, 0.10868)

## ltmle software package

Here note that `Treatment Estimate` and `Control Estimate` were fitted completely separately, and that they could had been obtained with separate calls:

## ltmle software package

Here note that Treatment Estimate and Control Estimate were fitted completely separately, and that they could had been obtained with separate calls:

Treatment Estimate:

```
summary(ltmle(...,  
          abar=c(1,1)))
```

Control Estimate:

```
summary(ltmle(...,  
          abar=c(0,0)))
```

Parameter Estimate: 0.3029  
Estimated Std Err: 0.086738  
p-value: 0.00047919  
95% Conf Interval: (0.1329, 0.4729)

Parameter Estimate: 0.36688  
Estimated Std Err: 0.015419  
p-value: <2e-16  
95% Conf Interval: (0.33666, 0.3971)

- For each parameter we get the standard error,<sup>a</sup> the p-value of the null hypothesis that that quantity equals zero, and confidence intervals

---

<sup>a</sup>influence curve based if the argument `variance.method="ic"` is specified, otherwise a more robust variance estimator based on TMLE is used.

# ltmle software package

Effect of a dynamic regime:

```
summary(ltmle(sim.data,  
             Anodes=paste0("A",0:1),  
             Lnodes=c(paste0("X0.", 1:3), paste0("X1.", 1:2)),  
             Ynodes="Y",  
             rule=function(row) c(1, ifelse(row["X1.1"]==1, 0,  
1))))
```

# ltmle software package

Effect of a dynamic regime:

```
summary(ltmle(sim.data,  
  Anodes=paste0("A",0:1),  
  Lnodes=c(paste0("X0.", 1:3), paste0("X1.", 1:2)),  
  Ynodes="Y",  
  rule=function(row) c(1, ifelse(row["X1.1"]==1, 0,  
1))))
```

Static regimes can also be specified as a dynamic regime:

```
summary(ltmle(sim.data,  
  Anodes=paste0("A",0:1),  
  Lnodes=c(paste0("X0.", 1:3), paste0("X1.", 1:2)),  
  Ynodes="Y",  
  rule=list(  
    treatment=function(row) c(1,1),  
    control=function(row) c(0,0))))
```



# ltmle software package

- ▶ `Qform`: character vector of regression formulas for the outcome regressions
  - ▶ `Qform` indicates what variables are included in each outcome regression
  - ▶ default is `NULL` which means that all variables from previous time-points are included
  - ▶ (does *not* mean that GLM is used)
- ▶ `gform`: character vector of regression formulas for the propensity scores
  - ▶ `gform` indicates what variables are included in each propensity score regression
  - ▶ default is `NULL` which means that all variables from previous time-points are included
  - ▶ (does *not* mean that GLM is used)

## ltmle software package

- ▶ `SL.library`: list with names entries `Q` and `g` specifying super learner libraries to pass to `SuperLearner` for the outcome regressions and the propensities scores

You can see available models for the super learner here:

<https://cran.r-project.org/web/packages/SuperLearner/vignettes/Guide-to-SuperLearner.html> (Section 4)

**NB** Some algorithms are really very slow

**NB** Some algorithms may not converge (gives error messages)

**NB** Think about what you know about each particular algorithm and do not just blindly include a ton of heavy algorithms

## ltmle software package

We can extract the super learner weights applied to each algorithm from the ltmle object (here called `fit.ltmle.sl`):

```
fit.ltmle.sl$fit$Q[[1]]
```

\$X1.1

	Risk	Coef
SL.glm_All	0.01672344	0.96580665
SL.mean_All	0.01765778	0.03419335
SL.glm.interaction_All	0.01681181	0.00000000
SL.glmnet_All	NA	0.00000000
SL.gam_All	0.01673172	0.00000000

\$Y

	Risk	Coef
SL.glm_All	0.2092472	0.97832296
SL.mean_All	0.2346187	0.02167704
SL.glm.interaction_All	0.2116557	0.00000000
SL.glmnet_All	0.2097088	0.00000000
SL.gam_All	0.2093541	0.00000000

## ltmle software package

We can extract the super learner weights applied to each algorithm from the ltmle object (here called `fit.ltmle.sl`):

```
fit.ltmle.sl$fit$g[[1]]
```

\$A0

	Risk	Coef
SL.glm_All	0.2508915	0
SL.mean_All	0.2502013	1
SL.glmnet_All	0.2502013	0
SL.gam_All	0.2512009	0

\$A1

	Risk	Coef
SL.glm_All	0.03332285	0.000000
SL.mean_All	0.08031019	0.000000
SL.glmnet_All	0.03330272	0.822445
SL.gam_All	0.03337456	0.177555

# Practical

## Application of ltmle

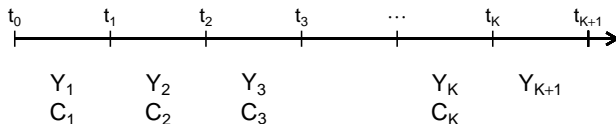
- ▶ Static and dynamic interventions
- ▶ Super learning

Proceed from **Task 2** of **day3\_practical2.pdf**.

## A note on: ltmle for right-censored data

Outcome process  $Y_k$  can jump from 0 to 1 at any time-point  $t_k$ .

Censoring process  $C_k$  can jump from 0 to 1 at any time-point  $t_k$ .



- ▶ Time to event:  $\tilde{T} = k' \wedge K + 1$  where  $k' = \min(k : Y_k = 1 \text{ or } C_k = 1)$
- ▶ Event indicator:  $\Delta = 1\{Y_{\tilde{T}} = 1\}$

## A note on: `ltmle` for right-censored data

One way that we can represent the survival data over a grid of  $K = 7$  time-points (long format), e.g., for three different individuals:

	id	k	Y	C
1:	2	1	0	0
2:	2	2	0	0
3:	2	3	0	0
4:	2	4	1	0

	id	k	Y	C
1:	5	1	0	0
2:	5	2	0	0
3:	5	3	0	0
4:	5	4	0	0
5:	5	5	0	0
6:	5	6	0	0
7:	5	7	0	0
8:	5	8	0	0

	id	k	Y	C
1:	7	1	0	0
2:	7	2	0	1

## A note on: `ltmle` for right-censored data

One way that we can represent the survival data over a grid of  $K = 7$  time-points (long format), e.g., for three different individuals:

	id	k	Y	C
1:	2	1	0	0
2:	2	2	0	0
3:	2	3	0	0
4:	2	4	1	0

	id	k	Y	C
1:	5	1	0	0
2:	5	2	0	0
3:	5	3	0	0
4:	5	4	0	0
5:	5	5	0	0
6:	5	6	0	0
7:	5	7	0	0
8:	5	8	0	0

	id	k	Y	C
1:	7	1	0	0
2:	7	2	0	1

Same data may also be presented as (wide format):

	id	Y1	C1	Y2	C2	Y3	C3	Y4	C4	Y5	C5	Y6	C6	Y7	C7	Y8
1:	2	0	0	0	0	0	0	1	0	1	0	1	0	1	0	1
2:	5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
3:	7	0	0	0	1	0	1	0	1	0	1	0	1	0	1	0



## A note on: `ltmle` for right-censored data

One way that we can represent the survival data over a grid of  $K = 7$  time-points (long format), e.g., for three different individuals:

	id	k	Y	C
1:	2	1	0	0
2:	2	2	0	0
3:	2	3	0	0
4:	2	4	1	0

	id	k	Y	C
1:	5	1	0	0
2:	5	2	0	0
3:	5	3	0	0
4:	5	4	0	0
5:	5	5	0	0
6:	5	6	0	0
7:	5	7	0	0
8:	5	8	0	0

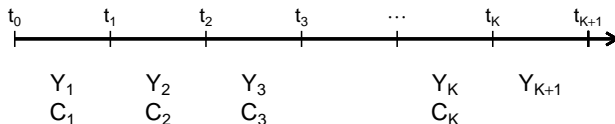
	id	k	Y	C
1:	7	1	0	0
2:	7	2	0	1

Same data may also be presented as (wide format):

	id	Y1	C1	Y2	C2	Y3	C3	Y4	C4	Y5	C5	Y6	C6	Y7	C7	Y8
1:	2	0	0	0	0	0	0	1	0	1	0	1	0	1	0	1
2:	5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
3:	7	0	0	0	1	0	1	0	1	0	1	0	1	0	1	0

(The wide format is needed for `ltmle`).

## A note on: ltmle for right-censored data



Example:

	id	X0.1	X0.2	X0.3	A0	Y1	C1	Y2	C2	Y3	C3	Y4	C4	Y5	C5	Y6	C6	Y7	C7	Y8
1:	1	0.408	-0.196	0	1	0	1	0	1	0	1	0	1	0	1	0	1	0	1	0
2:	2	-1.220	0.595	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0
3:	3	1.866	-1.609	0	1	0	0	1	0	1	0	1	0	1	0	1	0	1	0	1
4:	4	0.604	0.041	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
5:	5	-0.532	-1.251	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
6:	6	1.955	0.133	1	1	0	0	0	0	0	0	0	1	0	1	0	1	0	1	0

- ▶ Baseline covariate vector:  $X_0 = (X_{0,1}, X_{0,2}, X_{0,3})$
- ▶ Baseline treatment decision:  $A_0 \in \{0, 1\}$
- ▶ Time to event:  $\tilde{T} = k' \wedge K + 1$  where  $k' = \min(k : Y_k = 1 \text{ or } C_k = 1)$
- ▶ Event indicator:  $\Delta = 1\{Y_{\tilde{T}} = 1\}$

## A note on: ltmle for right-censored data

Generally:

$$O = (X_0, A_0, Y_1, C_1, X_1, A_1, \dots, Y_K, C_K, X_K, A_K, Y = X_{K+1}).$$

- ▶ Covariates  $X = (X_0, X_1, \dots, X_K)$  change over time.
- ▶ Treatment decisions  $A = (A_0, A_1, \dots, A_K)$  are updated over time.
- ▶ Censoring status  $C = (C_1, C_2, \dots, C_K)$  change over time.
- ▶ Outcome (death) status  $Y = (Y_1, Y_2, \dots, Y_K)$  change over time.

[After death/censoring, all variables are deterministically set to their last observed values].

## A note on: $ltmle$ for right-censored data

Inferring on the uncensored event time is like imposing a simple static intervention on all right-censoring nodes to impose 'no censoring'.

Target causal parameter:

$$\psi_{k^*}(P) = \mathbb{E}[Y_{k^*}^{A_0=a_0^*, C_1=0, A_1=a_1^*, \dots, C_K=0, A_K=a_K^*}] - \mathbb{E}[Y_{k^*}^{A_0=0, C_1=0, A_0=0, \dots, C_K=0, A_K=0}]$$

I.e., the effect of the treatment *had there been no loss to follow-up*.

= the absolute risk by time  $t_{k^*}$  if everyone had received treatment  $(a_0^*, a_1^*, \dots, a_K^*)$  contrasted to the absolute risk if everyone had not been treated.

## A note on: `ltmle` for right-censored data

The ordering of

$$O = (X_0, A_0, Y_1, C_1, X_1, A_1, \dots, Y_K, C_K, X_K, A_K, Y = X_{K+1}),$$

implies a temporal ordering:

$$X_0 \rightarrow A_0 \rightarrow \dots \rightarrow Y_k \rightarrow C_k \rightarrow X_k \rightarrow A_k \rightarrow \dots Y_{K+1}.$$

- ▶ at each time-point  $t_k$ , a patient is only at risk of dying if they did not yet die and they were not yet right-censored ( $Y_{k-1} = 0, C_{k-1} = 0$ ).
- ▶ at each time-point  $t_k$ , a patient is only at risk of being right-censoring if they did not yet die at this time and they were not yet right-censored ( $Y_k = 0, C_{k-1} = 0$ ).

## A note on: ltmle for right-censored data

Factorization of the density  $p$  of  $P \in \mathcal{M}$ :

$$\begin{aligned} p(o) &= \mu_{X_0}(x_0) \pi_{A_0}(a \mid x_0) \\ &\prod_{k=1}^K \left( \mu_{Y_k}(y_k \mid \bar{y}_{k-1}, \bar{c}_{k-1}, \bar{x}_{k-1}, \bar{a}_{k-1}) \pi_{C_k}(c_k \mid \bar{y}_k, \bar{c}_{k-1}, \bar{x}_k, \bar{a}_{k-1}) \right. \\ &\quad \times \mu_{X_k}(x_k \mid \bar{y}_k, \bar{c}_k, \bar{x}_{k-1}, \bar{a}_{k-1}) \pi_{A_k}(a_k \mid \bar{y}_k, \bar{c}_k, \bar{x}_k, \bar{a}_{k-1}) \Big) \\ &\times \mu_{Y_{K+1}}(y_{K+1} \mid \bar{y}_K, \bar{c}_K, \bar{x}_K, \bar{a}_K) \end{aligned}$$

*Without going into too many details, note for example that:*

- ▶  $\mu_{Y_k}(1 \mid \bar{0}_{k-1}, \bar{0}_{k-1}, \bar{x}_{k-1}, \bar{a}_{k-1})$  is the risk of outcome (dying) for a subject who did not yet die nor were right-censored plus had covariate and treatment history equal to  $\bar{x}_{k-1}, \bar{a}_{k-1}$ .
- ▶  $\pi_{C_k}(1 \mid \bar{0}_k, \bar{0}_{k-1}, \bar{x}_{k-1}, \bar{a}_{k-1})$  is the probability of being right-censored for a subject who did not yet die nor were right-censored plus had covariate and treatment history equal to  $\bar{x}_{k-1}, \bar{a}_{k-1}$ .

## A note on: ltmle for right-censored data

Identification of the target parameter as without censoring:

$$\bar{Q}_{K+1}(\bar{x}_K, \bar{a}_K, \bar{y}_K) = \mathbb{E}_P[Y_{K+1} \mid \bar{Y}_K = \bar{y}_K, \bar{C}_K = 0, \bar{X}_K = \bar{x}_K, \bar{A}_K = \bar{a}_K]$$

$$\bar{Q}_K(\bar{x}_{K-1}, \bar{a}_{K-1}, \bar{y}_{K-1}) = \mathbb{E}_P[Q_{K+1}(\bar{X}_K, a_K^*, \bar{A}_{K-1}, \bar{Y}_{K-1}) \mid \bar{Y}_{K-1} = \bar{y}_{K-1}, \bar{C}_{K-1} = 0, \bar{X}_{K-1} = \bar{x}_{K-1}, \bar{A}_{K-1} = \bar{a}_{K-1}]$$

$\vdots$

$$\bar{Q}_k(\bar{x}_{k-1}, \bar{a}_{k-1}, \bar{y}_{k-1}) = \mathbb{E}_P[Q_{k+1}(\bar{X}_k, a_k^*, \bar{A}_{k-1}, \bar{Y}_{k-1}) \mid \bar{Y}_{k-1} = \bar{y}_{k-1}, \bar{C}_{k-1} = 0, \bar{X}_{k-1} = \bar{x}_{k-1}, \bar{A}_{k-1} = \bar{a}_{k-1}]$$

$\vdots$

$$\bar{Q}_2(\bar{x}_1, \bar{a}_1, \bar{y}_1) = \mathbb{E}_P[Q_3(\bar{X}_2, a_2^*, A_1, Y_1) \mid \bar{Y}_1 = y_1, \bar{C}_1 = 0, \bar{X}_1 = \bar{x}_1, A_1 = a_1]$$

$$\bar{Q}_1(x_0, a_0) = \mathbb{E}_P[Q_2(\bar{X}_1, a_1^*) \mid X_0 = x_0, A_0 = a_0]$$

+ note that  $\bar{Q}_k(\bar{x}_{k-1}, \bar{a}_{k-1}, \bar{y}_{k-1}) = 1$  if  $y_{k'-1} = 1$  for  $k' \leq k$ .

## A note on: ltmle for right-censored data

The efficient influence function is given by:

$$\begin{aligned} & \tilde{\phi}^*(\bar{Q}, \pi)(O) \\ &= \sum_{k=1}^{K+1} 1\{Y_k = 0\} \left( \prod_{l=0}^{k-1} \frac{1\{A_l = a_l^*\} 1\{C_l = 0\}}{\pi_{A_l}(0, 0, a_l^* \mid \bar{x}_l, \bar{a}_{l-1}^*) \pi_{C_l}(0 \mid 0, 0, \bar{x}_l, \bar{a}_{l-1}^*)} \right) \\ & \quad \times \{ \bar{Q}_{k+1}(\bar{X}_k, a_k^*, \bar{A}_{k-1}, 0) - \bar{Q}_k(\bar{X}_{k-1}, \bar{A}_{k-1}, 0) \} + \bar{Q}_0(X_0) - \psi(P). \end{aligned}$$



## A note on: ltmle for right-censored data

```
ltmle(data,  
      Anodes, Lnodes, Cnodes, Ynodes,  
      abar, rule,  
      Qform, gform,  
      SL.library,  
      gbounds,  
      survivalOutcome=TRUE,  
      ...)
```

- ▶ Right-censoring nodes are specified in the Cnodes argument.
- ▶ The formatting of Cnodes is a bit peculiar — it should be a factor variable with the values 0 and 1 and the labels "uncensored" and "censored".
- ▶ Note that we further specify survivalOutcome=TRUE, so that Ynodes are treated as indicators of a terminating event.

## A note on: ltmle for right-censored data

- ▶ **Qform**: character vector of regression formulas for the outcome regressions
  - ▶ Qform indicates what variables are included in each outcome regression
  - ▶ default is NULL which means that all variables from previous time-points are included
  - ▶ (does *not* mean that GLM is used)
- ▶ **gform**: character vector of regression formulas for the propensity scores **and the hazards of censoring**
  - ▶ gform indicates what variables are included in each propensity score regression **and the hazards of censoring**
  - ▶ default is NULL which means that all variables from previous time-points are included
  - ▶ (does *not* mean that GLM is used)

## A note on: ltmle for right-censored data

For the example:

	id	X0.1	X0.2	X0.3	A0	Y1	C1	Y2	C2	Y3	C3	Y4	C4	Y5	C5	Y6	C6	Y7	C7	Y8
1:	1	0.408	-0.196	0	1	0	1	0	1	0	1	0	1	0	1	0	1	0	1	0
2:	2	-1.220	0.595	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0
3:	3	1.866	-1.609	0	1	0	0	1	0	1	0	1	0	1	0	1	0	1	0	1
4:	4	0.604	0.041	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
5:	5	-0.532	-1.251	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
6:	6	1.955	0.133	1	1	0	0	0	0	0	0	0	1	0	1	0	1	0	1	0

```
for (k in 1:7)
  sim.data[, (paste0("C", k)):=BinaryToCensoring(is.censored=get(
    paste0("C", k)))]
```

```
ltmle.fit <- ltmle(sim.data[, -"id"],
  Anodes="A0", Lnodes=paste0("X0.", 1:3),
  Cnodes=paste0("C", 1:7), Ynodes=paste0("Y", 1:8),
  abar=list(treatment=1, control=0),
  SL.library=list(Q=c("SL.glm", "SL.mean"),
    g=c("SL.glm", "SL.mean")),
  survivalOutcome=TRUE)
```

## A note on: ltmle for right-censored data

### Treatment Estimate:

Parameter Estimate: 0.40391  
Estimated Std Err: 0.031247  
p-value:  $<2e-16$   
95% Conf Interval: (0.34267, 0.46515)

### Control Estimate:

Parameter Estimate: 0.49545  
Estimated Std Err: 0.029052  
p-value:  $<2e-16$   
95% Conf Interval: (0.43851, 0.55239)

### Additive Treatment Effect:

Parameter Estimate: -0.091539  
Estimated Std Err: 0.041653  
p-value: 0.027973  
95% Conf Interval: (-0.17318, -0.0099008)

### Relative Risk:

Parameter Estimate: 0.81524  
Est Std Err log(RR): 0.095263  
p-value: 0.032009  
95% Conf Interval: (0.63620, 0.98250)

## A note on: ltmle for right-censored data

```
ltmle.fit$fit$g[[1]]
```

\$A0

	Risk	Coef
SL.glm_All	0.2502361	0.4568067
SL.mean_All	0.2501919	0.5431933

\$C1

	Risk	Coef
SL.glm_All	0.1533290	0.98494501
SL.mean_All	0.1667804	0.01505499

\$C2

	Risk	Coef
SL.glm_All	0.1365783	0.956276
SL.mean_All	0.1428211	0.043724

\$C3

	Risk	Coef
SL.glm_All	0.1373765	0.92097765
SL.mean_All	0.1440681	0.07902235