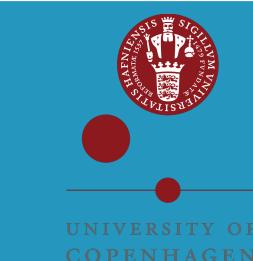
A New Iterated Conditional Expectations Estimator for Longitudinal Causal Effects in Continuous Time



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Motivation

- In medical research, the estimation of causal effects of treatments over time is often of interest.
- Continuous-time inference allows for data that is more closely aligned with the data collection process (Table 1). Moreover, discrete time approaches usually require the discretization of time, leading to a loss of information.
- There is a scarcity of (applied) literature on the estimation of longitudinal causal effects in continuous time. Rytgaard et al. (2022) considered a targeted minimum-loss based estimator based on iterated conditional expectations (Figure 1) for estimating causal effects flexibly. However, this estimator quickly becomes intractable.
- We propose a new feasible iterated conditional expectations estimator (Figure 2) for the estimation of longitudinal causal effects in continuous time.

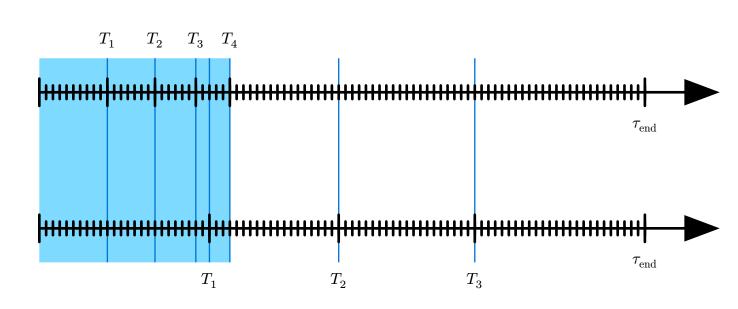


Figure 1: The figure illustrates the sequential regression approach given in Rytgaard et al. (2022) for two observations.

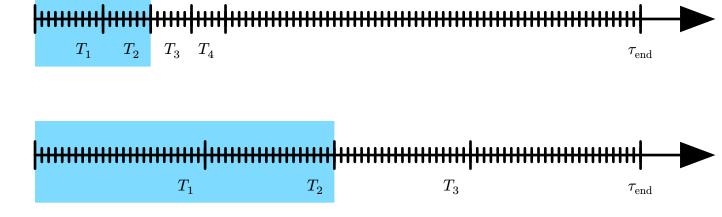


Figure 2: The figure illustrates the sequential regression approach proposed in this article.

id	time	event
1	3	side effect
1	8	primary event
2	10	primary event
3	2	side effect
3	5	treatment shift
3	7	censoring

Table 1: An example of a longitudinal dataset from electronic health records or a clinical trial. Events are registered at irregular/subject-specific time points.

Setting

Let $(N^a(t), A(t), N^\ell(t), L(t), N^y(t), N^d(t), N^c(t))^1$ be a stochastic processes observed in $[0, \tau_{\text{end}}]$, consisting of a counting process for treatment visits, treatment values, a counting process for treatment covariate measurements, covariate values, and counting processes for the primary event, competing event, and censoring, respectively. Furthermore, $A(t) \in \{0,1\}$ and $L(t) \in \mathcal{L}$, where $\mathcal{L} \subseteq$ \mathbb{R}^d is a finite set.

Assumption 1: In the time interval $[0, \tau_{\text{end}}]$ there are at most $K-1 < \infty$ many changes of treatment and covariates in total for a single individual.

Assumption 2: The counting processes N^a , N^ℓ , N^y , N^d , and N^c have with probability 1 no jump times in common.

Under these assumptions, the observed data can be written in the form

$$O=\mathcal{F}_{T_{(K)}}$$

where

$$\mathcal{F}_{T_{(k)}} = \left(T_{(k)}, \Delta_k, A\Big(T_{(k)}\Big), L\Big(T_{(k)}\Big)\right) \vee \mathcal{F}_{T_{(k-1)}} \text{ and } \mathcal{F}_0 = (L_0, A_0)$$

Assumption 3: For each $k \in \{1,...,K\}$, $T_{(k)} \mid \mathcal{F}_{T_{(k-1)}} \ll m^2$, $A(T_{(k)}) \mid T_{(k)} = t$, $\Delta_k = t$ $a, \mathcal{F}_{T_{(k-1)}} \ll
u_a, ext{ and } Lig(T_{(k)}ig) \mid T_{(k)} = t, D_{(k)} = \ell, \mathcal{F}_{T_{(k-1)}} \ll
u_\ell.$

Target parameter

Let \tilde{T}_k^1 and $\tilde{\Delta}_k^1$ be the counterfactual event time and indicator for the k'th had the patient stayed on treatment and initially received treatment (and not been censored). Our target parameter Ψ^g_{τ} : $\mathcal{M} \to \mathbb{R}$ is the mean interventional potential outcome at time τ given the intervention plan g.

$$\Psi^g_\tau(P) = \mathbb{E}_P \left[\sum_{k=1}^K \mathbb{1} \left\{ \tilde{T}^\mathbf{1}_k \leq \tau, \tilde{\Delta}^\mathbf{1}_k = y \right\} \right].$$

Identification

We consider identification conditions in Theorem 3 of Ryalen (2024). These are stated in our present uncensored setting. Let $\tilde{Y}_t = \left(\mathbb{1}\left\{\tilde{T}_1^1 \leq t, \tilde{\Delta}_1^1 = y\right\}, ..., \mathbb{1}\left\{\tilde{T}_K^1 \leq t, \tilde{\Delta}_K^1 = y\right\}\right)$ and $T^a = \inf\{t > 0: t \in \mathcal{T}_t^a = t \in \mathcal$ $A(t) \neq 1$. For each $k \in \{1, ..., K\}$, we need:

• Consistency:

$$\mathbb{1} \big\{ \tilde{T}_k^{\mathbf{1}} \leq t, \tilde{\Delta}_k^{\mathbf{1}} = y \big\} \mathbb{1} \{ T^a > t, A(0) = 1 \} = \mathbb{1} \{ T_k \leq t, \Delta_k = y \} \mathbb{1} \{ T^a > t, A(0) = 1 \}$$

for $t \in [0, \tau_{\mathrm{end}}]$.

• Exchangeability:

$$A \Big(T_{(k)} \Big) \perp \Big(\mathbb{1} \Big\{ \tilde{T}_{k+1}^{\mathbf{1}} \leq t, \tilde{D}_{k+1}^{\mathbf{1}} = y \Big\}, ..., \mathbb{1} \Big\{ \tilde{T}_{K}^{\mathbf{1}} \leq t, \tilde{D}_{K}^{\mathbf{1}} = y \Big\} \Big) \mid D_{(k)} = a, \mathcal{F}_{T_{(k-1)}}$$

(and

$$\begin{split} \lambda^a \bigg(t \mid \mathcal{F}_{T_{(k-1)}} \vee \left(\tilde{Y}_t \right)_{t \in [0,\tau_{\text{end}}]} \bigg) \\ &= \lim_{h \to 0} \frac{P \bigg(t \leq T_{(k)} < t + h, \Delta_k = a \mid T_{(k)} \geq t, \mathcal{F}_{T_{(k-1)}}, \left(\tilde{Y}_t \right)_{t \in [0,\tau_{\text{end}}]} \bigg)}{h} \end{split}$$

does not depend on $\left(ilde{Y}_t
ight)_{t \in [0, au_{\mathrm{end}}]}$).

Positivity: The weights

$$w_k(f_{k-1},t_k) = \frac{\mathbbm{1}\{a_0=1\}}{\pi_0(l_0)} \prod_{j=1}^{k-1} \left(\frac{\mathbbm{1}\{a_j=1\}}{\pi_j(f_{j-1})}\right)^{\mathbbm{1}\{\delta_j=a\}} \mathbbm{1}\{t_1 < \ldots < t_k\}$$

$$\operatorname{fulfill} \mathbb{E}_P \Big[w_k \Big(\mathcal{F}_{T_{(k-1)}}, T_{(k)} \Big) \Big] = 1.$$

Identification formula

Under the assumptions of consistency, exchangeability, and positivity, the target parameter is identified via

$$\Psi^g_\tau(P) = \mathbb{E}_P\left[\sum_{k=1}^K w_k\Big(\mathcal{F}_{T_{(k-1)}}, T_{(k)}\Big)\mathbb{1}\{T_k \leq \tau, \Delta_k = y\}\right].$$

Iterated Conditional Expectation Estimator

The form of the efficient influence function (Bickel et al. (1993)) in this setting suggests the use of a iterated conditional expectations estimator. Let $S^c\Big(t\mid \mathcal{F}_{T_{(k)}}\Big)$ be the conditional survival function of the censoring time given the k previous events and $\mathcal{F}_{T_{(k)}}^{-A}$ denote the history without the treatment process.

Proposed continuous-time ICE estimator

- For each event point k = K, K 1, ..., 1 (starting with k = K):
- 1. Obtain $\hat{S}^c(t \mid \mathcal{F}_{T_{(k)}})$ by fitting a cause-specific hazard model for the censoring via the interevent time $S_{(k)} = T_{(k)} - T_{(k-1)}$, regressing on $\mathcal{F}_{T_{(k-1)}}$ (among the people who are still at risk after k events).
- 2. Define the subject-specific weight:

$$\hat{\eta}_k = \frac{\mathbb{1} \Big\{ T_{(k)} \leq \tau, \Delta_k \in \{a,\ell\}, k < K \Big\} \hat{\nu}_k \Big(\mathcal{F}_{T_{(k)}}^{-A}, \mathbf{1} \Big)}{\hat{S}^c \Big(T_{(k)} \mid \mathcal{F}_{T_{(k-1)}} \Big)}$$

Then calculate the subject-specific pseudo-outcome

$$\hat{R}_k = \frac{\mathbb{1}\big\{T_{(k)} \leq \tau, \Delta_k = y\big\}}{\hat{S}^c\big(T_{(k)} \mid \mathcal{F}_{T_{(k-1)}}\big)} + \hat{\eta}_k$$

Regress \hat{R}_k on $\mathcal{F}_{T_{(k-1)}}$ on the data with $T_{(k-1)} < \tau$ and $\Delta_k \in \{a,\ell\}$ to obtain a prediction function $\hat{\nu}_{k-1}:\mathcal{H}_{k-1}\to\mathbb{R}_+$.

• At baseline, we obtain the estimate $\hat{\Psi}_n = \frac{1}{n} \sum_{i=1}^n \hat{\nu}_0(L_i(0), 1)$.

Existing discrete-time ICE estimator

- 1. For each time point k=K,K-1,...,1 (starting with $\hat{R}_K=Y$): Regress \hat{R}_k on $\left(\bar{L}_{k-1}, \bar{A}_{k-1}\right)$ to obtain $\hat{\nu}_{k-1}$; obtain predictions $\hat{R}_{k-1} = \hat{\nu}_{k-1}(\bar{L}_{k-1}, \mathbf{1})$.

 2. At baseline, we obtain the estimate $\hat{\Psi}_n = \frac{1}{n} \sum_{i=1}^n \hat{\nu}_0(L_i(0), 1)$.
- This approach can be extended to survival outcomes.

References

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