

## **Paper review:**

# **A nonparametric framework for treatment effect modifier discovery in high dimensions**

Philippe Boileau, Ning Leng, Nima S. Hejazi, Mark van der Laan and Sandrine Dudoit (2025)

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# Table of contents

1. Introduction
2. Background on nonparametric asymptotic theory
3. TEM-VIPs for continuous outcomes
4. General framework for TEM-VIPs
5. Simulation studies
6. Data application
7. Discussion

# Table of contents

1. Introduction
2. Background on nonparametric asymptotic theory
3. TEM-VIPs for continuous outcomes
4. General framework for TEM-VIPs
5. Simulation studies
6. Data application
7. Discussion

# Introduction

In causal inference, discovering **treatment effect modifiers (TEMs)** is crucial for understanding heterogeneous treatment effects in a population.

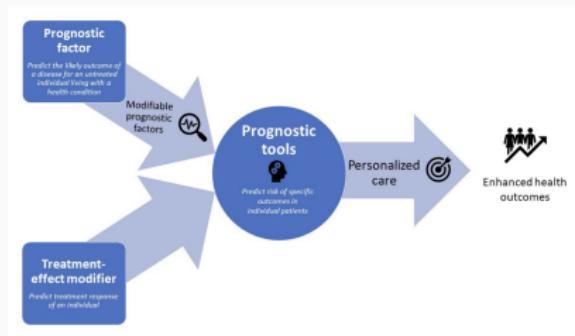


Figure reference: Tousignant-Laflamme et al. (2022)

- Precision medicine: Identifying patient subgroups exhibiting different benefits from a therapy
- Economics: Assessing government policies on diverse population strata

## Prior approaches to TEM discovery.

- **Parametric modeling:** Add treatment-covariate interaction terms in a linear outcome model, and assess the statistical significance of interaction coefficients.
- **CATE estimation:** Assess variable importance in CATE prediction.
  - Permutation-based / Dropout-based approaches
  - Penalized regression methods with feature selection properties (e.g. LASSO)

However, the above methods *rely on unverifiable assumptions* (model misspecification, sparsity, correlation structures) or produce *unreliable results in high-dimensional settings*.

# Introduction

In this paper, we explore a general framework for defining and performing inference about marginal **TEM variable importance parameters (TEM-VIPs)**.

- This work generalizes the method of Boileau et al. (2022) to continuous and binary outcomes, which directly estimates the strength of covariates' capacity to modify the treatment effect.
- The proposed framework does not rely on stringent assumptions on the DGP and accommodates high-dimensional settings.
- The framework is equipped with tools to define appropriate TEM-VIPs and corresponding EIF-based nonparametric estimators that allow valid and reliable inference.

# Table of contents

---

1. Introduction
2. Background on nonparametric asymptotic theory
3. TEM-VIPs for continuous outcomes
4. General framework for TEM-VIPs
5. Simulation studies
6. Data application
7. Discussion

# Background on nonparametric asymptotic theory

## Notation.

- **Full data:**  $\{X_i\}_{i=1}^n$ ,  $X_i = (W_i, A_i, Y_i^{(0)}, Y_i^{(1)}) \stackrel{\text{i.i.d.}}{\sim} P_{X,0} \in \mathcal{M}_X$ 
  - $W_i$ : Set of  $p$  covariates, possibly  $p \gg n$
  - $A_i$ : Binary treatment indicator
  - $Y_i^{(a)}$ : Potential outcomes; assume  $Y_i^{(a)} \in (0, 1)$
  - $\mathcal{M}_X$ : Nonparametric model of possible DGPs
- **Observed data:**  $O = (W, A, Y) \sim P_0 \in \mathcal{M}$ 
  - Consistency:  $Y = AY^{(1)} + (1 - A)Y^{(0)}$
  - Under consistency,  $\mathcal{M}$  is fully determined by  $\mathcal{M}_X$

## Background on nonparametric asymptotic theory

Recall that  $P(Y, A, W) = P(Y | A, W) P(A | W) P(W)$ . Thus, we define the following nuisance functions:

- $P_0$ : DGP of observed data
- $P_n$ : Empirical distribution
- $\hat{P}_n$ : Plug-in estimator of  $P_n$ , made up of elements of  $P_n$  and possibly nuisance parameter estimators
- Expected conditional outcome:  $\bar{Q}_0(A, W) = \mathbb{E}_{P_0}[Y | A, W]$ 
  - Full-data counterpart:  $\bar{Q}_{P_{X,0}}(a, W) = \mathbb{E}_{P_{X,0}}[Y^{(a)} | W]$
- Propensity score:  $g_0(W) = P_{P_0}[A = 1 | W]$

**Nonparametric EIF-based estimators** possess numerous attractive properties:

- Efficiency
- Root- $n$  consistency and asymptotic normality
- No assumptions on outcome/PS models (robustness, flexibility)

## Background on nonparametric asymptotic theory

**Motivation.** Let  $\Theta(P)$  be the parameter of interest and  $\Theta(\hat{P}_n)$  be its plug-in estimator.

- Let  $D(O, P)$  be the influence function  $\Theta(P)$  at  $P$ .
- The asymptotic behavior of  $\Theta(\hat{P}_n)$  is captured by the following von Mises expansion:

$$\begin{aligned}\sqrt{n}(\Theta(\hat{P}_n) - \Theta(P_0)) &= \underbrace{\sqrt{n} E_{P_n}[D(O, P_0)]}_{\text{Term 1}} - \underbrace{\sqrt{n} E_{P_n}[D(O, \hat{P}_n)]}_{\text{Term 2}} \\ &\quad + \underbrace{\sqrt{n}(E_{P_n} - E_{P_0})(D(O, \hat{P}_n) - D(O, P_0))}_{\text{Term 3}} \\ &\quad - \underbrace{\sqrt{n} R(P_0, \hat{P}_n)}_{\text{Term 4}}.\end{aligned}$$

## Background on nonparametric estimation theory

- **Term 1:**  $\sqrt{n} E_{P_n}[D(O, P_0)]$ 
  - Converges to a mean-zero Gaussian r.v. with variance  $\mathbb{E}_{P_0}[D(O, P_0)^2]$
  - **EIF-based estimators minimize this asymptotic variance**
- **Term 2:**  $\sqrt{n} E_{P_n}\left[D(O, \hat{P}_n)\right]$ 
  - Asymptotic bias
  - We construct estimators that eliminate this bias term
- **Terms 3 & 4:** Converges to 0 under standard assumptions

## Background on nonparametric estimation theory

**Estimators.** By leveraging the EIF  $D(O, P)$ , nonparametric estimators correct first-order bias and attain semiparametric efficiency.

(i) **One-step:** Adds empirical EIF correction

$$\Theta^{(OS)}(\hat{P}_n) = \Theta(\hat{P}_n) + \mathbb{E}_{P_n}[D(O, \hat{P}_n)]$$

(ii) **Estimating equation:** Solves the estimating equation

$$\mathbb{E}_{P_n}[D(O, \hat{P}_n)] = 0.$$

(iii) **Targeted maximum likelihood (TML):** Tilt  $\hat{P}_n$  to generate  $P_n^*$  such that  $\mathbb{E}_{P_n}[D(O, \hat{P}_n^*)] \approx 0$ , and define  $\Theta^{\text{TML}}(\hat{P}_n) = \Theta(P_n^*)$ .

- TML estimators constrain estimates to the parameter space.
- We will look at an example later.

## Background on nonparametric estimation theory

**Inference.** Inference about  $\Theta(P_0)$  follows naturally from *asymptotic normality*.

In particular, the  $\alpha$ -level Wald-type confidence interval for  $\Theta(P_0)$  can be constructed identically for the three estimators  $\Theta^{(*)}(\hat{P}_n)$ :

$$\Theta^{(*)}(\hat{P}_n) \pm z_{1-\alpha/2} \sqrt{\frac{\mathbb{E}_{P_0}[D(O, P_0)^2]}{n}},$$

where the term  $\mathbb{E}_{P_0}[D(O, P_0)^2]$  is substituted with  $\mathbb{E}_{P_0}[D(O, \hat{P}_n)^2]$  in practice.

## Example: EIF of ATE

Intuitively, the EIF is the *orthogonal projection* of any influence function onto the *tangent space of the observed-data model*.

- Influence function is the analogue of a *gradient vector* (in standard calculus).
- Consider, for instance, the ATE defined in terms of the full-data model

$$\Theta^F(P_{X,0}) = \mathbb{E}_{P_{X,0}}[Y^{(1)} - Y^{(0)}].$$

- If we had access to full data, the influence function would be given by  $Y^{(1)} - Y^{(0)} - \Theta(P_{X,0})$ .
- We project this IF onto the tangent space of the observed-data model to obtain

$$\begin{aligned} D(O, P_0) = & \frac{A}{g_0(W)}(Y - \bar{Q}_0(1, W)) - \frac{1-A}{1-g_0(W)}(Y - \bar{Q}_0(0, W)) \\ & + \bar{Q}_0(1, W) - \bar{Q}_0(0, W) - \Theta(P_0). \end{aligned}$$

## Example: EIF of ATE

Deriving one-step and estimating equation estimators of the ATE using the EIF, we obtain

$$\begin{aligned}\Theta_{OS} = \Theta_{EE} = \frac{1}{n} \sum_{i=1}^n & \left[ \left( \frac{A_i}{g_0(W_i)} - \frac{1 - A_i}{1 - g_0(W_i)} \right) \{ Y_i - \bar{Q}_0(A_i, W_i) \} \right. \\ & \left. + \bar{Q}_0(1, W_i) - \bar{Q}_0(0, W_i) \right].\end{aligned}$$

which coincides with the AIPW estimator for the ATE.

# Table of contents

---

1. Introduction
2. Background on nonparametric asymptotic theory
3. TEM-VIPs for continuous outcomes
4. General framework for TEM-VIPs
5. Simulation studies
6. Data application
7. Discussion

## TEM-VIPs for continuous outcomes

**Causal parameter.** Suppose the outcome is continuous, and the treatment effect of interest is the ATE, i.e.  $\mathbb{E}_{P_{X,0}}[Y^{(1)} - Y^{(0)}]$ .

First, we impose assumptions for computational convenience:

- (A1) Centered covariates:  $\mathbb{E}_{P_{X,0}}[W_j] = 0$ .
- (A2) Nonzero variance:  $\mathbb{E}_{P_{X,0}}[W_j^2] > 0$ .

# TEM-VIPs for continuous outcomes

## Definition (TEM-VIP, with respect to ATE)

An (*absolute*) TEM-VIP of the  $j$ -th covariate is defined as a mapping

$$\begin{aligned}\Psi_j^F(P_{X,0}) &:= \frac{\text{Cov}_{P_{X,0}}[Y^{(1)} - Y^{(0)}, W_j]}{\mathbb{V}_{P_{X,0}}[W_j]} \\ &= \frac{\mathbb{E}_{P_{X,0}}[(Y^{(1)} - Y^{(0)})W_j]}{\mathbb{E}_{P_{X,0}}[W_j^2]} \\ &= \frac{\mathbb{E}_{P_{X,0}}\left[\left(\bar{Q}_{P_{X,0}}(1, W) - \bar{Q}_{P_{X,0}}(0, W)\right)W_j\right]}{\mathbb{E}_{P_{X,0}}[W_j^2]}.\end{aligned}$$

The full estimand is then given by

$$\Psi^F : \mathcal{M}_X \rightarrow \mathbb{R}^p, \quad \Psi^F(P_{X,0}) = (\Psi_1^F(P_{X,0}), \dots, \Psi_p^F(P_{X,0})).$$

## TEM-VIPs for continuous outcomes

**Interpretation.** The TEM-VIP  $\Psi^F(P_{X,0})$  can be viewed as assessing the *correlation between the difference in potential outcomes and each potential TEM*, renormalized to be on the same scale as  $Y$ .

- Assume that the expectation of  $f(W) = \bar{Q}_{P_{X,0}}(1, W) - \bar{Q}_{P_{X,0}}(0, W)$  conditioned on  $W_j$  is linear in  $W_j$ , i.e.  $\mathbb{E}_{P_{X,0}}[f(W) | W_j] = \beta_j W_j$ . Then,  $\Psi^F(P_{X,0})$  is the vector of *simple linear regression coefficients*:

$$\Psi_j^F(P_{X,0}) = \frac{\mathbb{E}_{P_{X,0}}[f(W)W_j]}{\mathbb{E}_{P_{X,0}}[W_j^2]} = \frac{\mathbb{E}_{P_{X,0}}[\beta_j W_j^2]}{\mathbb{E}_{P_{X,0}}[W_j^2]} = \beta_j.$$

- When the relationship between  $f(W)$  and the  $W_j$ 's are nonlinear, the parameter can be interpreted as a linear model projection:

$$\Psi_j^F(P_{X,0}) = \underset{\beta_j \in \mathbb{R}}{\operatorname{argmin}} \mathbb{E}_{P_{X,0}} \left[ \left( (\bar{Q}_{P_{X,0}}(1, W) - \bar{Q}_{P_{X,0}}(0, W)) - (\alpha + \beta_j W_j) \right)^2 \right].$$

# TEM-VIPs for continuous outcomes

Under (A3) *unconfoundedness* and (A4) *positivity* assumptions, the full-data TEM-VIP  $\Psi_j^F(P_{X,0})$  is identifiable from observed data:

## Theorem 1 (Identifiability of TEM-VIP)

Assuming that (A1), (A2), (A3) and (A4) hold, we have

$$\Psi_j(P_0) := \frac{\mathbb{E}_{P_0}[(\bar{Q}_0(1, W) - \bar{Q}_0(0, W)) W_j]}{\mathbb{E}_{P_0}[W_j^2]} = \Psi_j^F(P_{X,0}).$$

Hence, the parameter  $\Psi : \mathcal{M} \rightarrow \mathbb{R}^p$ ,  $\Psi(P_0) = (\Psi_1(P_0), \dots, \Psi_p(P_0))$  is equal to the full-data estimand  $\Psi^F(P_{X,0})$ .

**Efficient influence function.** The EIF provides the basis for the construction of nonparametric estimators.

## Proposition 1 (EIF of observed-data parameter)

Under (A1) and (A2), the EIF at  $P \in \mathcal{M}$  of  $\Psi_j(P)$  is given by

$$D_j(O, P) = \frac{W_j}{\mathbb{E}_P[W_j^2]} \left( \frac{2A - 1}{Ag(W) + (1 - A)(1 - g(W))} (Y - \bar{Q}(A, W)) \right. \\ \left. + \bar{Q}(1, W) - \bar{Q}(0, W) - \Psi_j(P) W_j \right).$$

**Remark.** Note that the blue term corresponds to the *uncentered EIF* of the ATE, which is the treatment effect of interest.

## Estimators.

(i) *One-step and estimating equation estimators*: The one-step and estimating equation estimators of  $\Psi_j(P_0)$  are identical. Let  $\bar{Q}_n$  and  $g_n$  be estimators of  $\bar{Q}_0$  and  $g_0$  trained of  $P_n$  and included in  $\hat{P}_n$ . Then,

$$\begin{aligned}\Psi_j^{(\text{OS})}(\hat{P}_n) &= \Psi_j^{(\text{EE})}(\hat{P}_n) = \frac{1}{\sum_{i=1}^n W_{ij}^2} \sum_{i=1}^n W_{ij} \\ &\quad \times \left( \frac{2A_i - 1}{A_i g_n(W_i) + (1 - A_i)(1 - g_n(W_i))} (Y_i - \bar{Q}_n(A_i, W_i)) \right. \\ &\quad \left. + \bar{Q}_n(1, W_i) - \bar{Q}_n(0, W_i) \right).\end{aligned}$$

## TEM-VIPs for continuous outcomes

### (ii) Targeted maximum likelihood (TML) estimator

**Step 1.** Define the negative log-likelihood loss function for  $\bar{Q}$  as

$$L(O; \bar{Q}) = -\log \left\{ \bar{Q}(A, W)^Y (1 - \bar{Q}(A, W))^{1-Y} \right\},$$

and a parametric working submodel for  $\bar{Q}$  as

$$\bar{Q}_j(\epsilon)(A, W) = \text{logit}^{-1} \left\{ \text{logit } \bar{Q}(A, W) + \epsilon H_j(A, W) \right\},$$

where

$$H_j(A, W) := \frac{W_j}{E_P[W_j^2]} \frac{2A - 1}{Ag(W) + (1 - A)(1 - g(W))}.$$

## TEM-VIPs for continuous outcomes

**Step 2.** Denoting an initial estimator of  $\bar{Q}_0$  trained on  $P_n$  by  $\bar{Q}_n^0$ , update  $\bar{Q}_n^0$  by computing  $\epsilon_{n,j}^1$  such that

$$\epsilon_{n,j}^1 = \arg \min_{\epsilon} E_{P_n} [L(O; \bar{Q}_{n,j}^0(\epsilon))].$$

Compute the *tilted conditional outcome estimator*  $\bar{Q}_{n,j}^1 = \bar{Q}_{n,j}^0(\epsilon_{n,j}^1)$  and the *tilted distribution*  $P_n^*$ , where  $\bar{Q}_n^0$  is replaced by  $\bar{Q}_{n,j}^1$  in  $\hat{P}_n$ .

**Step 3.** The TML estimator of the  $j$ -th TEM-VIP is given by

$$\Psi_j^{\text{TML}}(\hat{P}_n) = \Psi_j(P_n^*).$$

# TEM-VIPs for continuous outcomes

**Asymptotic behavior.** Note that the asymptotic distributions of  $\Phi^{\text{OS}}(\hat{P}_n)$ ,  $\Phi^{\text{EE}}(\hat{P}_n)$  and  $\Phi^{\text{TML}}(\hat{P}_n)$  are identical due to their dependence on the common EIF  $D_j(O, P_0)$ .

First, we establish *consistency* results and its double-robustness.

(A5) Outcome model estimator consistency:  $\|\bar{Q}_n(A, W) - \bar{Q}_0(A, W)\|_2^2 = o_P(1)$ .

(A6) PS estimator consistency:  $\|g_n(W) - g_0(W)\|_2^2 = o_P(1)$ .

## Proposition 2 (Consistency)

Under (A1) and (A2), and *either (A5) or (A6)*,

$$\Phi^{(*)}(\hat{P}_n) \xrightarrow{P} \Phi(P_0), \quad * = \text{OS, EE, TML}.$$

## TEM-VIPs for continuous outcomes

Moreover, the *asymptotic normality* of the estimators' sampling distributions can be specified under the following assumptions:

(A7) Donsker conditions: There exists a  $P_0$ -Donsker class  $\mathcal{G}_0$  such that

$$P_{P_0} \left[ D_j(O, \hat{P}_n) \in \mathcal{G}_0 \right] \rightarrow 1 \text{ and}$$

$$E_{P_0} \left[ (D_j(O, \hat{P}_n) - D_j(O, P_0))^2 \mid \hat{P}_n \right] = o_P(1) \text{ for each } j.$$

(A8) Shared rate convergence:

$$\|\bar{Q}_n(A, W) - \bar{Q}_0(A, W)\|_2 \|g_n(W) - g_0(W)\|_2 = o_P(n^{-1/2}).$$

(A9) Uniformly bounded covariates:  $|W_j| \leq C$  for  $j = 1, \dots, p$ .

### Theorem 2 (Asymptotic normality)

Under (A1), (A2), (A7), (A8) and (A9),

$$\sqrt{n} (\Psi_j^{(*)}(\hat{P}_n) - \Psi_j(P_0)) \xrightarrow{D} N(0, E_{P_0}[D_j(O, P_0)^2]), \quad * = \text{OS, EE, TML}.$$

# Table of contents

---

1. Introduction
2. Background on nonparametric asymptotic theory
3. TEM-VIPs for continuous outcomes
4. General framework for TEM-VIPs
5. Simulation studies
6. Data application
7. Discussion

# General framework for TEM-VIPs

The workflow presented in the previous section for continuous outcomes can be unified into a general framework.

For instance, suppose we are interested in causal effects on *right-censored time-to-event* outcomes, which are common in clinical trials.

- Causal parameters of interest are often built upon the conditional survival function  $S_{P_{X,0}}(t | a, W) := P_{P_{X,0}}[T^{(a)} > t | W]$ , where  $T^{(a)}$  denotes the (potential) event time.
- **CATE of survival probability at time  $t$ :**

$$\mathbb{E}_{P_{X,0}}[S_{P_{X,0}}(t | 1, W) - S_{P_{X,0}}(t | 0, W) | W].$$

- **Difference in conditional restricted mean survival times (RMSTs):**

$$\begin{aligned} \mathbb{E}_{P_{X,0}} & \left[ \min\{T^{(1)}, t\} - \min\{T^{(0)}, t\} \mid W \right] \\ &= \mathbb{E}_{P_{X,0}} \left[ \int_0^t \{S_{P_{X,0}}(u | 1, W) - S_{P_{X,0}}(u | 0, W)\} \, du \mid W \right]. \end{aligned}$$

# General framework for TEM-VIPs

**Step 1.** Select a full-data, pathwise differentiable parameter  $\Phi^F(P_{X,0})$  of some treatment effect that is relevant to the problem at hand.

- ▶ ATE (for continuous/binary outcomes)

$$\Phi^F(P_{X,0}) = \mathbb{E}_{P_{X,0}}[Y^{(1)} - Y^{(0)}]$$

- ▶ Difference in RMSTs (for right-censored time-to-event outcomes)

$$\Phi^F(P_{X,0}) = \mathbb{E}_{P_{X,0}}[\min\{T^{(1)}, t\} - \min\{T^{(0)}, t\}]$$

# General framework for TEM-VIPs

**Step 2.** Define  $f(W)$  such that  $\mathbb{E}_{P_{X,0}}[f(W)] = \Phi^F(P_{X,0})$ .

- CATE (for estimation of ATE)

$$f(W) = \bar{Q}_{P_{X,0}}(1, W) - \bar{Q}_{P_{X,0}}(0, W)$$

- Conditional difference in RMSTs (for difference in RMSTs)

$$f(W) = \int_0^t \{S_{P_{X,0}}(u \mid 1, W) - S_{P_{X,0}}(u \mid 0, W)\} \, du$$

Under (A1) and (A2), the **TEM-VIP** of the  $j$ -th covariate  $\Theta_j^F$  is given by

$$\Theta_j^F = \frac{\mathbb{E}_{P_{X,0}}[f(W)W_j]}{\mathbb{E}_{P_{X,0}}[W_j^2]}.$$

### Step 3. Establish the *identifiability* of the TEM-VIP.

- Let  $\Theta_j$  and  $\Phi$  be the observed-data counterparts of  $\Theta_j^F$  and  $\Phi^F$ , respectively.
- The conditions establishing  $\Theta_j^F(P_{X,0}) = \Theta_j(P_0)$  are identical to the conditions for the identifiability of the treatment effect of interest, i.e.,  $\Phi^F(P_{X,0}) = \Phi(P_0)$ .
  - e.g., consistency, unconfoundedness, positivity.
  - The only additional assumption is that  $W_j$  has bounded variance.

## General framework for TEM-VIPs

**Step 4.** Derive the *EIF* of the TEM-VIP.

If the uncentered EIF of  $\Phi(P_0)$  is given by  $d(O, P_0)$ , then the EIF of the TEM-VIP is given by

$$\frac{W_j}{\mathbb{E}_{P_0}[W_j^2]} \{d(O, P_0) - W_j \Theta_j(P_0)\}.$$

**Step 5.** Construct the EIF-based estimators for the TEM-VIP.

- These include *(i) one-step*, *(ii) estimating equation* and *(iii) TML* estimators.
- The asymptotic properties of the EIF-based estimators are identical to those of the nonparametric efficient estimators of  $\Phi$ , such as double-robustness, provided that  $W_j$  is bounded.

# Table of contents

---

1. Introduction
2. Background on nonparametric asymptotic theory
3. TEM-VIPs for continuous outcomes
4. General framework for TEM-VIPs
5. Simulation studies
6. Data application
7. Discussion

## Simulation studies

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Through simulation studies, we observe:

- (i) *Finite-sample performance* of the proposed one-step and TML estimators.
- (ii) The *estimators' capacity to recover TEMs*, compared to (augmented) modified covariates methods (Tian et al. (2014) & Chen et al. (2017)).

We consider three DGPs: (i) *continuous outcome, observational study*, (ii) *binary outcome, observational study* and (iii) *survival outcome, RCT*.

- Sample sizes:  $n = 125, 250, 500, 1,000, 2,000$
- 200 MC simulations under each setting

## Simulation setup

**DGP 1.** Continuous outcome, observational study

$$W \sim N(0, I_{500 \times 500})$$

$$A | W \sim \text{Bernoulli} \left( \text{logit}^{-1} \left( \frac{1}{4} (W_1 - W_2 + W_3) \right) \right)$$

$$Y | A, W \sim 1 + 2 \left| \sum_{j=1}^5 W_j \right| + (5A - 2) \sum_{j=1}^5 W_j + \varepsilon, \quad \varepsilon \sim N(0, 1/2)$$

**DGP 2.** Binary outcome, observational study

$$W \sim N(0, \Sigma_{100 \times 100}), \quad \Sigma_{ij} = \begin{cases} 1, & i = j, \\ 0.1|i - j|^{-1.8}, & \text{otherwise} \end{cases}$$

$$A | W \sim \text{Bernoulli} \left( \text{logit}^{-1} \left( \frac{1}{4} (W_1 + W_2 + W_3) \right) \right)$$

$$Y | A, W \sim \text{Bernoulli} \left( \text{logit}^{-1} \left( 1 - 2A + \sum_{j=1}^5 W_j + \left( A - \frac{1}{2} \right) \sum_{j=1}^5 W_j \right) \right)$$

## Simulation setup

**DGP 3.** Right-censored time-to-event outcome, randomized control trial

$$W \sim N(0, \Sigma_{300 \times 300})$$

$$A \sim \text{Bernoulli}\left(\frac{1}{2}\right)$$

$$C \mid A, W \sim \min\left\{ \text{NegBin}\left(1, \text{logit}^{-1}(5 + A + W_1)\right), 10 \right\}$$

$$T \mid A, W, C \sim \text{NegBin}\left(1, \text{logit}^{-1}\left(-2 - A + (10A - 5) \sum_{j=1}^{10} W_j\right)\right)$$

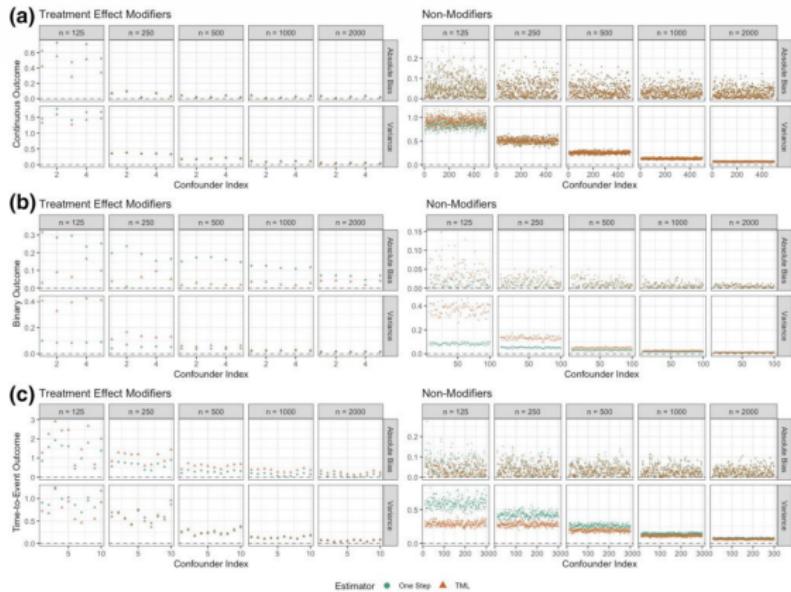
$$\tilde{T} = \min\{T, C\}$$

$$\Delta = \mathbb{I}(T > c)$$

- The covariance matrix  $\Sigma$  is block-diagonal, with each block corresponding to ten moderately correlated features.
- The target estimand is defined based on difference in RMSTs at  $t = 9$  (total duration consists of 10 time units).

# Simulation 1: Finite-sample performance

## Simulation 1. Finite-sample performance of estimators



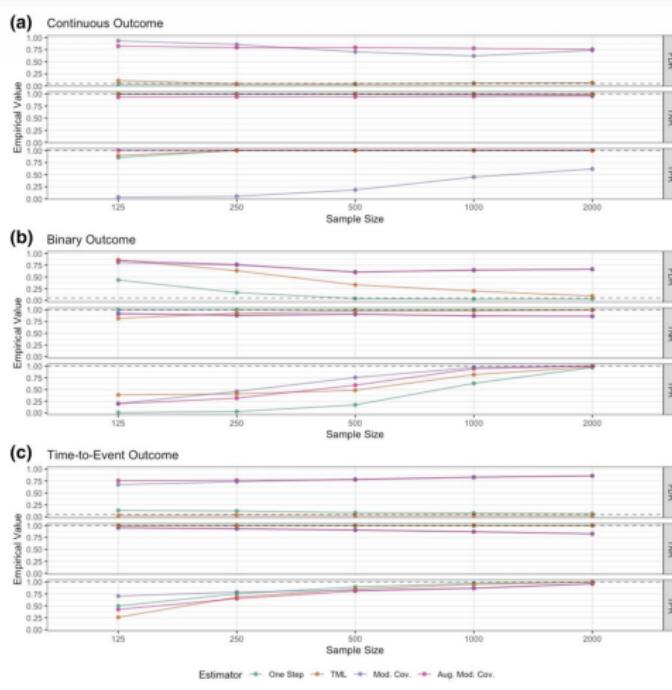
**Figure 1.** Empirical bias and variance of one-step and TML estimators in the continuous, binary, and time-to-event outcome simulation scenarios.

## Simulation 2: TEM discovery and classification

### Simulation 2. Estimators' capacity to recover TEMs

- The estimators' ability to distinguish TEMs is evaluated in terms of empirical false discovery rate (FDR), true negative rate (TNR) and true positive rate (TPR).
- Two-sided Wald-type tests at the 5% level were adjusted for multiple testing using the Benjamini–Hochberg procedure.
- The one-step and TML estimators are compared to modified covariates and augmented modified covariates methods with LASSO.
  - Modified covariates: Outcomes are transformed so that only the treatment-covariate interactions in a GLM need be modeled.
  - Augmented modified covariates: The transformed outcomes are modeled as a function of all covariates to improve efficiency.

## Simulation 2: TEM discovery and classification



**Figure 2.** TEM classification results of one-step, TML, modified covariates, augmented modified covariates estimators in the continuous, binary, and time-to-event outcome simulation scenarios.

# Table of contents

---

1. Introduction
2. Background on nonparametric asymptotic theory
3. TEM-VIPs for continuous outcomes
4. General framework for TEM-VIPs
5. Simulation studies
6. Data application
7. Discussion

## Data application

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The proposed framework is applied to a clinical trial dataset with a *right-censored time-to-event outcome*.

**Data source.** Subset of FinHER dataset provided by Loi et al. (2014)

- Trastuzumab significantly improves clinical outcomes of breast cancer patients with tumors overexpressing HER2; however, the improvement is not uniform.
- Patients with overexpressed HER2 were randomized to receive either 9 weekly trastuzumab infusions or no trastuzumab as adjuvant treatment for early-stage breast cancer.

## Variables of interest.

- **Treatment:** Trastuzumab (antibody targeting the HER2 oncogene)
- **Covariates (potential TEMs):** 500 genes selected by variability
- **Outcome:** Distant disease-free survival, i.e. time between randomization and first cancer recurrence or death

## Implementation of proposed methodology.

- **Target estimand:** RMST-based TEM-VIP
- **Estimation:** TML estimator based on conditional failure and censoring hazards estimates by a Super Learner

# Data application

## Result 1. Classification of covariates into TEMs and non-TEMs

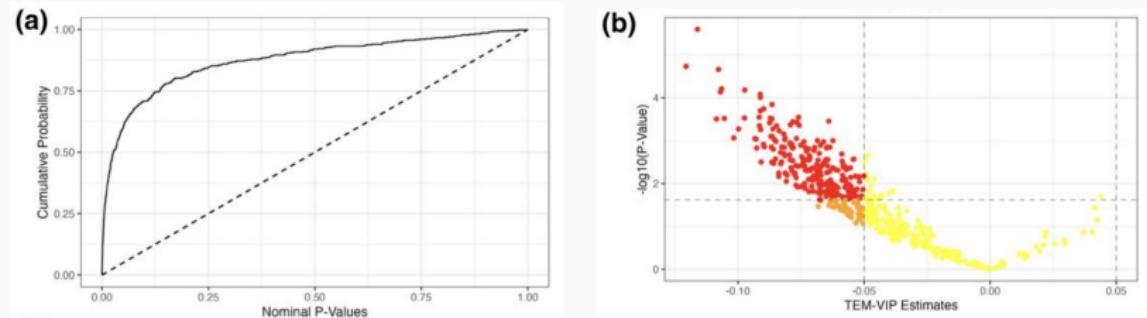


Figure 3. (a) Empirical cdf of p-values.

(b) Volcano plot of the 500 most variable genes' TEM-VIP estimates and associated p-values. Yellow genes are deemed unimportant due to their small estimated effect sizes and larger p-values; orange genes possess a meaningful estimated effect but fail to achieve the adjusted p-value cut-off; red genes are significant at the 5% FDR level and have large estimated TEM-VIPs.

# Data application

## Result 2. Top five selected treatment effect modifiers

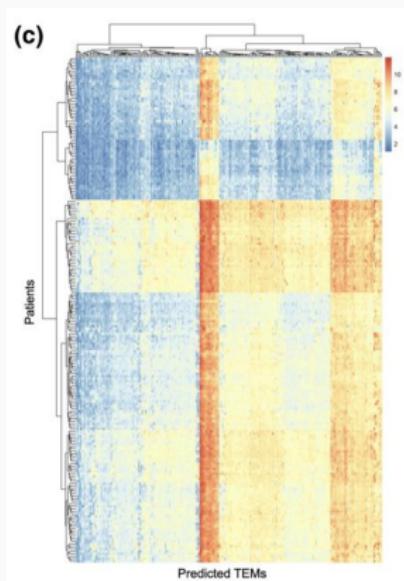
	Gene	Estimate	SE	Adj. <i>p</i> -value
1	EPPK1	-0.116	0.025	.001
2	NDUFB3	-0.121	0.028	.004
3	BNIP3L	-0.108	0.025	.004
4	PNKD	-0.106	0.027	.006
5	DUSP4	-0.097	0.024	.006

All of the genes above have previously been linked to breast cancer:

- EPPK1: Increased expression has been linked to estrogen-related receptor  $\gamma$ , which is associated with breast cancer growth suppression.
- NDUF3: A single nucleotide polymorphism in NDUFB3 promoter is reported to be significantly associated with estrogen receptor negative breast cancer.

# Data application

## Result 3. Clustering of patients with respect to predicted TEMs



**Figure 3. (c)** The log-transformed gene expression data of genes with meaningful effect estimates are used to cluster patients. Hierarchical clustering with complete linkage is used for patients and identified TEMs alike.

## Observations.

- *(Result 1)* We observe a continuum in the biomarkers' capacity to influence the treatment effect, in terms of both statistical significance and effect size.
  - Hypothesis testing alone may not be adequate.
  - One can instead deem a biomarker of clinical interest if it is significant at the 5% FDR level and its absolute estimated TEM-VIP is larger than 0.05.
- *(Result 3)* The gene expression data produce multiple distinct patient clusters.
  - However, the authors suggest that it should be considered solely as a diagnostic tool, since redefining subgroups within the same dataset based on the estimated TEM-VIPs may result in overfitting.

# Table of contents

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1. Introduction
2. Background on nonparametric asymptotic theory
3. TEM-VIPs for continuous outcomes
4. General framework for TEM-VIPs
5. Simulation studies
6. Data application
7. Discussion

# Discussion

## Contributions of the paper:

- (i) Proposed a *general workflow* for defining TEM-VIPs with respect to the treatment effect of interest and for deriving associated *nonparametric estimators based on the EIF*.
- (ii) The methodology is applicable to *various types of outcomes*, and the performance is validated under *high-dimensional settings*.

## Future directions:

- (i) Statistically rigorous subgroup discovery.
- (ii) Improving treatment effect estimation in high-dimensional settings by exploiting TEM-VIPs as variable filters.
- (iii) Deriving standardized TEM-VIPs.

## Reference

-  Boileau, P., Leng, N., Hejazi, N. S., van der Laan, M. J., & Dudoit, S. (2025). *A nonparametric framework for treatment effect modifier discovery in high dimensions*. *Journal of the Royal Statistical Society: Series B (Statistical Methodology)*, 87(1), 157–185.
-  Boileau, P., Qi, N. T., van der Laan, M. J., Dudoit, S., & Leng, N. (2022). *A flexible approach for predictive biomarker discovery*. *Biostatistics*, 24(4), 1085–1105.
-  Gruber, S. and van der Laan, M. J. (2009). *Targeted Maximum Likelihood Estimation: A Gentle Introduction*. U.C. Berkeley Division of Biostatistics Working Paper Series. Working Paper 252.