Limits of Estimating Heterogeneous Treatment Effects: Guidelines for Practical Algorithm Design

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Abstract
Estimating heterogeneous treatment effects from observational data is a central problem in many domains. Because counterfactual data is inaccessible, the problem differs fundamentally from supervised learning, and entails a more complex set of modeling choices. Despite a variety of recently proposed algorithmic solutions, a principled guideline for building estimators of treatment effects using machine learning algorithms is still lacking. In this paper, we provide such guidelines by characterizing the fundamental limits of estimating heterogeneous treatment effects, and establishing conditions under which these limits can be achieved. Our analysis reveals that the relative importance of the different aspects of observational data vary with the sample size. For instance, we show that selection bias matters only in small-sample regimes, whereas with a large sample size, the way an algorithm models the control and treated outcomes is what bottlenecks its performance. Guided by our analysis, we build a practical algorithm for estimating treatment effects using a non-stationary Gaussian processes with doubly-robust hyperparameters. Using a standard semi-synthetic simulation setup, we show that our algorithm outperforms the state-of-the-art, and that the behavior of existing algorithms conforms with our analysis.

1. Introduction
The problem of estimating heterogeneous (individualized) causal effects of a treatment from observational data is central in many application domains, including public health and drug development (Foster et al., 2011), computational advertising (Bottou et al., 2013), and social sciences (Xie et al., 2012). The increasing availability of observational data in all these domains has encouraged the development of various machine learning algorithms tailored for inferring treatment effects using observational data (e.g. (Li & Fu, 2017; Wager & Athey, 2017; Shalit et al., 2017; Alaa & van der Schaar, 2017)). Due to the peculiarity of the treatment effect estimation problem, these algorithms needed to address various modeling aspects that are foreign to standard supervised learning setups; such aspects include ways to handle sample selection bias (Heckman, 1977), and ways to model treated and untreated data points. Despite a variety of recent algorithmic approaches, principled guidelines for model design are lacking.

In this paper, we identify guiding principles for designing practical treatment effect estimation algorithms in the context of Bayesian nonparametric inference, and propose one an algorithm that follows these guidelines. We set these guidelines by characterizing the fundamental limits of estimating treatment effects, and studying the impact of various common modeling choices on the achievability of those limits. In what follows, we provide a brief technical background for the treatment effect estimation problem, along with a summary of our contributions.

1.1. Background and Summary of Contributions
Our analysis hinges on the Rubin-Neyman potential outcomes model (Rubin, 2005). That is, we consider an observational dataset with a population of subjects, where each subject $i$ is endowed with a $d$-dimensional feature $X_i \in \mathcal{X}$. We assume that $\mathcal{X} = [0,1]^d$, but most of our results hold for general compact metric spaces (bounded, closed sets in $\mathbb{R}^d$). A treatment assignment indicator $W_i \in \{0,1\}$ is associated with subject $i$; $W_i = 1$ if the treatment under study was applied to subject $i$, and $W_i = 0$ otherwise. Subject $i$’s responses with and without the treatment (the potential outcomes) are denoted as $Y_i^{(1)}$ and $Y_i^{(0)}$, respectively. Treatments are assigned to subjects according to an underlying policy that depends on the subjects’ features, i.e. $W_i \perp X_i$. This dependence is quantified via the conditional distribution $p(x) = \Pr(W_i = 1 | X_i = x)$, also known as the propensity score of subject $i$ (Rosenbaum & Rubin,
The response $Y_i^{(W_i)}$ is the “factual outcome” which we observe in the data, whereas $Y_i^{(1-W_i)}$ is the unrealized "counterfactual outcome" (Bottou et al., 2013). An observational dataset $D_n$ comprises $n$ samples of the form:

$$D_n = \{X_i, W_i, Y_i^{(W_i)}\}_{i=1}^n$$

The causal effect of the treatment on subject $i$ with a feature $X_i = x$ is characterized through the conditional average treatment effect (CATE) function $T(x)$, which is defined as the expected difference between the two potential outcomes (Rubin, 2005), i.e.

$$T(x) = E[Y_i^{(1)} - Y_i^{(0)} | X_i = x]$$

Our goal is to identify a set of guiding principles for building estimators of the CATE $T(x)$ using samples from $D_n$. Throughout the paper, we will assume that the joint density $dP(X_i, W_i, Y_i^{(0)}, Y_i^{(1)})$ supports the assumptions of unconfoundedness and overlap, which are necessary for causal identifiability and consistency. Unconfoundedness requires that $(Y_i^{(0)}, Y_i^{(1)}) \perp W_i | X_i$, whereas overlap requires that $0 < p(x) < 1$ (Rosenbaum & Rubin, 1984). Selection bias occurs in $D_n$ since the distribution of the treated/control subjects does not match that of the overall population.

In order to come up with principled guidelines for building estimators of $T(x)$, we characterize the fundamental (information-theoretic) limits of estimating the CATE using samples from $D_n$, and identify the modeling choices that would allow achieving those limits. To this end, in Section 3 we tackle the following question: what are the fundamental limits of CATE estimation? We answer this question by deriving the optimal minimax rate for estimating $T(x)$ using $D_n$. Interestingly, it turns out that the optimal rate does not depend on selection bias, but rather on the smoothness and sparsity of the more “complex” of the functions $E[Y_i^{(0)} | X_i = x]$ and $E[Y_i^{(1)} | X_i = x]$. We focus our analysis on Bayesian nonparametric methods, since they have the appealing properties of being robust to misspecification and are accessible for theoretical analysis.

Our analysis reveals that the relative importance of the different modeling aspects vary with the sample size. In particular, in the large-sample regime, selection bias does not pose a serious problem, and the model’s performance would be mainly determined by its structure, i.e. the way the outcomes $Y_i^{(0)}$ and $Y_i^{(1)}$ are modeled, and the impact of that on variable selection and hyperparameter tuning. On the contrary, selection bias can seriously harm a model’s generalization performance in small-sample regimes. A good model should then be carefully designed so that it operates well in both regimes by possessing the right model structure that would allow learning at a fast rate, and the right model selection (hyperparameter optimization) scheme that would account for selection bias.

In Section 4, we build a practical CATE estimation algorithm guided by the results of the analyses in Section 3. We model the outcomes $Y_i^{(0)}$ and $Y_i^{(1)}$ using a Gaussian process with a non-stationary kernel that captures the different relevant variables and different levels of smoothness of the functions $E[Y_i^{(0)} | X_i = x]$ and $E[Y_i^{(1)} | X_i = x]$. We prove that this model structure can achieve the optimal rate of CATE estimation when tuned with the right hyperparameters. We also propose a doubly-robust hyperparameter optimization scheme that accounts for selection bias in small-sample regimes, without hindering the model’s minimax-optimality in the large sample limit. We show that our algorithm outperforms state-of-the-art methods using a well-known semi-synthetic simulation setup.

1.2. Related Work

Very few works have attempted to characterize the limits of CATE estimation, or study the impact of different modeling choices on the CATE estimation performance in a principled manner. (Alaee & van der Schaaf, 2018) characterized the asymptotic “information rates” for different CATE estimators, but provided no clear guidelines on practical model design or an analysis of the impact of sample selection bias. The study in (Künzel et al., 2017) was rather empirical in nature, comparing the performance of different regression structures for the potential outcomes while ignoring selection bias. A similar study, but focusing only on random forest models, was conducted in (Lu et al., 2017).

Most of the previous works have been algorithmic in nature, focusing mainly on devising algorithms that correct for selection bias (e.g. (Johansson et al., 2016; Yoon et al., 2018; Wager & Athey, 2017; Li & Fu, 2017)). Some of these works cast the selection bias problem as a problem of covariate shift (Sugiyama et al., 2007), and use techniques from representation learning to learn feature maps that balance the biased data (e.g. (Li & Fu, 2017; Shalit et al., 2017; Johansson et al., 2016)). However, those works report much bigger improvements in CATE estimation when changing their model structure (e.g. architecture of a neural network), as compared to the gains attained by only accounting for bias (see the comparisons between the TARnet and BNN models in (Shalit et al., 2017)). Similar observations are reported in (Alaee & van der Schaaf, 2017; Atan et al., 2018), where the selection of the model structure seemed to influence the achieved CATE estimation performance even when selection bias is not accounted for. However, none of these works offer a discussion on whether selection bias is actually the main challenge in CATE estimation, or whether the outcomes’ model structure may have a bigger influence on performance.

In contrast to the works above, this paper does not attempt to develop a model by presupposing that particular model-
ing aspects are of greater importance than others, but rather provides a framework for understanding the limits on the achievable performance, and how different modeling aspects influence a model’s chance of achieving those limits. We use our analyses to both reflect on the modeling choices made in the works above, and also devise a novel, principled CATE estimation algorithms that achieves the fundamental performance limits.

2. Estimating CATE: Problem Setup

2.1. Potential Outcomes & Propensity Score

We consider the following random design regression model for the potential outcomes:

\[ Y_i^{(w)} = f_w(X_i) + \varepsilon_{i,w}, \quad w \in \{0, 1\}, \]  

(3)

where \( \varepsilon_{i,w} \sim \mathcal{N}(0, \sigma_w^2) \) is a Gaussian noise variable. It follows from (2) that the CATE is \( T(x) = f_1(x) - f_0(x) \). The response surfaces \( f_1(x) \) and \( f_0(x) \) correspond to the subjects’ responses with and without the treatment.

We assume that \( f_w(\cdot) : \mathcal{X} \to \mathbb{R}, \quad w \in \{0, 1\} \), is a totally bounded function that lives in a space of “smooth” or “regular” functions, with an unknown smoothness parameter \( \alpha_w \). We use Hölder balls for concreteness, although our results extend to other function spaces. A function \( f_w(\cdot) \) lies in the Hölder ball \( H_{\alpha_w} \), with a Hölder exponent \( \alpha_w > 0 \), if and only if it is bounded in sup-norm by a constant \( C > 0 \), all its partial derivatives up to order \( \lfloor \alpha_w \rfloor \) exist, and all its partial derivatives of order \( \lfloor \alpha_w \rfloor \) are Lipschitz with exponent \( (\alpha_w - \lfloor \alpha_w \rfloor) \) and constant \( C \). The Hölder exponents quantify the complexities of \( f_0 \) and \( f_1 \), and hence the hardness of estimating \( T(x) \) would depend on \( \alpha_0 \) and \( \alpha_1 \).

2.2. Bayesian Nonparametric Inference

Nonparametric inference is immune to misspecification of the outcomes’ and propensity models (Kennedy, 2018), and hence we focus on Bayesian nonparametric methods for inferring \( T(\cdot) \) on the basis of \( D_n \). Bayesian inference entails specifying a prior distribution \( \mathcal{P} \) over \( f_1(\cdot) \) and \( f_0(\cdot) \), i.e.

\[ f_0, f_1 \sim \mathcal{P}(\varphi_{\beta_0}, \varphi_{\beta_1}). \]  

(4)

where \( \varphi_{\beta_w} = \{\varphi_{\beta_w}^k\}_{k=1}^\infty, \quad w \in \{0, 1\} \), are complete orthonormal bases (indexed by a parameter \( \beta_w > 0 \)) with respect to the Lebesgue measure in \( \mathcal{X} \), \( f_w = \sum_k \varphi_{\beta_w}^k \), and \( \hat{f}_w = \{\hat{f}_w, \varphi_{\beta_w}^k\}_k \). Thus, for given bases \( \varphi_{\beta_0} \) and \( \varphi_{\beta_1} \), \( \mathcal{P} \) places a probability distribution on the projections \( \{\hat{f}_w\}_k \).

Potential choices for the basis \( \varphi_{\beta_w} \) that would give rise to implementable Bayesian inference algorithms include regular wavelet basis (Zhang, 1997), radial basis for a reproducing kernel Hilbert space (RKHS) (van der Vaart et al., 2008), etc. In general, the parameter \( \beta_w \) would determine the smoothness of the function space spanned by \( \varphi_{\beta_w} \).

2.3. Towards Principled CATE Estimation

To evaluate the predictive accuracy of the Bayesian inference procedure, we analyze the “frequentist” loss of point estimators \( \hat{T}(x) \) induced by the Bayesian posterior \( d\mu_n(T(x)|D_n) \), assuming that \( D_n \) is generated based on fixed, true response surfaces \( f_1(x) \) and \( f_0(x) \). (This type of analysis is sometimes referred to as the “Frequentist-Bayes” analysis (Sniekers et al., 2015).) In particular, we quantify the performance of a point estimator \( \hat{T}(x) = \delta(d\mu_n(T(x)|D_n)) \) by its squared-L^2(\mathcal{P}) error, which was dubbed the precision of estimating heterogeneous effects (PEHE) in (Hill, 2011), and is formally defined as:

\[ \psi(\hat{T}) \triangleq \mathbb{E} \| \hat{T} - T \|^2_{L^2(\mathcal{P})}, \]  

(5)

where \( L^2(\mathcal{P}) \) is the \( L^2 \) norm with respect to the feature distribution, i.e. \( \|f(x)\|^2_{L^2(\mathcal{P})} = \int f^2(x) \, d\mathcal{P}(X = x) \).

Not a standard supervised learning problem...

The “fundamental problem of causal inference” is that for every subject \( i \) in \( D_n \), we only observe the factual outcome \( Y_i^{(W_i)} \), whereas the counterfactual \( Y_i^{(1 - W_i)} \) remains unknown, which renders empirical evaluation of the PEHE in (5) impossible. Moreover, \( D_n \) would generally exhibit sample selection bias (Heckman, 1977), because the treatment assignment mechanism (decided by \( p(x) \)) creates a discrepancy between the feature distributions of the treated/control population and the overall population. Thus, standard supervised learning approaches based on empirical risk minimization cannot be used to learn a generalizable model for the CATE from samples in \( D_n \). This gives rise to the following fundamental modeling questions that are peculiar to the CATE estimation problem:

- [Q1]: How should the treatment assignment indicator \( W_i \) be incorporated into the learning model?
- [Q2]: How should selection bias be handled?

Adequate answers to [Q1] and [Q2] would provide guidelines for selecting the prior \( \mathcal{P}(\varphi_{\beta_0}, \varphi_{\beta_1}) \). Addressing the modeling questions above requires a profound understanding of the fundamental limits of CATE estimation, in addition to an understanding of the impact of different modeling choices on the achievability of such limits. The next Sections provide principled answers to [Q1] and [Q2] by addressing the following, more fundamental questions:

Section 3: What are the limits on the performance that can be achieved by any estimator of the CATE?

Section 4: How can we build practical algorithms that can achieve the performance limits?
3. Fundamental Limits of CATE Estimation

In this Section, we establish an information-theoretic limit on the performance of any CATE estimator. In what follows, we use the standard Bachmann-Landau order notation, and write $a \vee b = \max\{a, b\}$, $a \wedge b = \min\{a, b\}$. The notation $a \preceq b$ means that $a \leq Cb$ for a universal constant $C$, and $\asymp$ denotes asymptotic equivalence.

3.1. Optimal Minimax Rates

The “hardness” of a nonparametric estimation problem is typically characterized by its minimax risk (Stone, 1982), i.e. the minimum worst case risk achieved by any estimator when the estimand is known to live in a given function space (Yang et al., 2015). In the following Theorem, we establish the optimal minimax rate for the PEHE risk in terms of the complexity of the response surfaces $f_0$ and $f_1$.

**Theorem 1.** Suppose that $\mathcal{X} = [0, 1]^d$, and that $f_w$ depends on a subset of $d_w$ features with $d_w \leq \min\{n, d\}$ for $w \in \{0, 1\}$. If $f_0 \in H_{\alpha_0}^{\rho_0}$ and $f_1 \in H_{\alpha_1}^{\rho_1}$, then the optimal minimax rate is:

$$\inf_{\hat{T}} \sup_{f_0, f_1} \psi(\hat{T}) \asymp n^{-1} (\frac{d_{\alpha_0} \vee d_{\alpha_1}}{\rho_0 \rho_1})^{-\frac{1}{4}} \log\left(\frac{d_{\alpha_0}^2 + d_{\alpha_1}^2}{\rho_0^2 \rho_1^2}\right)^{\frac{1}{4}}.$$  

The above holds for any $p(.) \in H_{\alpha_p}^{\rho_p}, \alpha_p > 0$. □

In Theorem 1, the supremum is taken over $\alpha_w$-Hölder balls $(w \in \{0, 1\})$, whereas the infimum is taken over all possible Bayesian estimators. The minimax rate in Theorem 1 corresponds to the fast rates by which any (Bayesian) estimator $\hat{T}(.)$ can approximate the CATE function $T(.)$. The proof of Theorem 1 (provided in the supplement) uses information-theoretic techniques based on Fano’s method to derive algorithm-independent estimation rates (Yang & Barron, 1999). In the following set of remarks, we revisit [Q1] and [Q2] in the light of the results of Theorem 1.

How can Theorem 1 help us address [Q1] & [Q2]?

**Remark 1 (Smoothness & sparsity)**

Theorem 1 says that estimating CATE is as hard as nonparametric regression for functions with additive sparsity (Raskutti et al., 2009; Yang et al., 2015). The minimax rate in Theorem 1 decomposes into a term reflecting the complexity of CATE estimation under correct variable selection for $f_0$ and $f_1$, and a term reflecting the complexity of variable selection. Variable selection complexity remains small as long as $\log(d) = \Theta(n^\zeta)$, for some $\zeta \in (0, 1)$, and approaches the parametric rates as $\zeta \to 0$. The minimax rate will generally be dominated by the complexity of CATE estimation, and will approach the parametric rates only for very smooth response surfaces with small number of relevant dimensions, i.e. $\frac{d_{\alpha_0}}{\rho_0} \vee \frac{d_{\alpha_1}}{\rho_1} \to 0$.

The main takeaway from Theorem 1 is that the CATE learning rate is determined by the more “complex” of the surfaces $f_0$ and $f_1$, where complexity is quantified by the sparsity-to-smoothness ratio $d_w/\alpha_w$ for $w \in \{0, 1\}$. Thus, a model would achieve the optimal CATE learning rate only if it selects the correct relevant variables for $f_0$ and $f_1$, and tunes its “hyperparameters” (i.e. smoothness of the prior) to cope with a complexity of $\frac{d_{\alpha_0}}{\rho_0} \vee \frac{d_{\alpha_1}}{\rho_1}$. When $\frac{d_{\alpha_0}}{\rho_0}$ and $\frac{d_{\alpha_1}}{\rho_1}$ are very different (e.g. $f_0$ and $f_1$ have different relevant features), rate-optimal estimation is possible only if the model incorporates such differences in $I(\hat{\varphi}_{\beta_0}, \hat{\varphi}_{\beta_1})$.

The discussion above provides a concrete answer to [Q1]: the treatment assignment variable $w$ should be incorporated into the model in such a way that it encodes the different relevant dimensions and smoothness levels of $f_0$ and $f_1$ in the bases $\hat{\varphi}_{\beta_0}$ and $\hat{\varphi}_{\beta_1}$. (The simplest way to achieve this is to use two separate models for $f_0$ and $f_1$.) This is not fulfilled by many of the previous models that built a single regression function of the from $f : \mathcal{X} \times \{0, 1\} \to \mathbb{R}$, and estimated the CATE as $\hat{T}(x) = f(x, 1) - f(x, 0)$ (Hill, 2011; Johansson et al., 2016; Powers et al., 2017). This is because such models enforced the smoothness of the prior along all features to be the same for $w = 0$ and $w = 1$.

**Remark 2 (Selection bias)**

Theorem 1 gives a rather surprising answer to [Q2]: the optimal learning rate is oblivious to selection bias. Such a finding is consistent with previous results on nonparametric kernel density estimation under selection bias (Borojoo et al., 2017), and parametric Bayesian inference under covariate shift (Shimodaira, 2000; Sugiyama & Storkey, 2007). It shows that many of the recent works have missed the target; the works in (Johansson et al., 2016; Shalit et al., 2017; Alaa & van der Schaar, 2017) cast the problem of CATE estimation as one of covariate shift that results from selection bias. However, Theorem 1 says that selection bias is not a problem when we have a sufficiently large amount of data. This is because selection bias is inherently a mis-specification problem, and hence its impact on nonparametric inference is washed away in large-sample regimes.

Remarks 1 and 2 posit an explanation for various recurrent (empirical) findings reported in previous literature. For instance, (Hahn et al., 2017) found that separate modeling of $f_0$ and $f_1$ via Bayesian additive regression trees (BART) outperforms the well-known single-surface BART model developed in (Hill, 2011). Similar findings were reported for models based on Gaussian processes (Alaa & van der Schaar, 2017), and models based on deep neural networks (Shalit et al., 2017). All such findings can be explained in the light of Remark 1. On the other hand, Remark 2 may provide an explanation as to why the “TARnet” model in (Shalit et al., 2017), which models $f_0$ and $f_1$ using separate neural networks and does not account for selection
bias, outperformed the “BNN” model in (Johansson et al., 2016), which regularizes for selection bias but fits a single-output network for \( f_0 \) and \( f_1 \).

3.2. Backing off from “Asymptopia”

Theorem 1 shows that selection bias does not hinder the optimal minimax rates, and that it is only the structural properties of the prior \( \Pi(\varphi_{\beta_0}, \varphi_{\beta_1}) \) that determine a model’s rate of learning. But does the achieved learning rate suffice as a sole criterion for addressing the modeling questions [Q1] and [Q2]? The answer is “yes” only if \( D_n \) comes from a large observational dataset, in which case the learning rate suffices as a descriptor for the large-sample performance. However, if \( D_n \) is small, which is typical in post-hoc analyses of clinical trials (Foster et al., 2011), then one should make the design choices that would optimize the small-sample performance. In order to give a more complete picture of the performance in large and small-sample regimes, we derive the following bound on the PEHE:

\[
\begin{aligned}
\psi(\hat{T}) & \leq \bar{C} \cdot \exp(D_2(Q_0 || Q)) \cdot ||f_0 - \hat{f}_0||_{L^2(P_0)}^2 \\
& + \bar{C} \cdot \exp(D_2(Q_1 || Q)) \cdot ||f_1 - \hat{f}_1||_{L^2(P_1)}^2,
\end{aligned}
\]

(6)

for some \( \bar{C} > 0 \), where \( L^2(P_\omega) \), for \( \omega \in \{0, 1\} \), is the \( L^2 \) norm with respect to \( d\mathbb{P}(X = x \mid W = \omega) \), \( Q = d\mathbb{P}(X = x) \), \( Q_\omega = d\mathbb{P}(X = x \mid W = \omega) \), and \( D_m(p || q) \) is the \( m \)th order Rényi divergence. The bound in (6) holds for all \( n > 0 \), and is tight (refer to the supplement); it shows that the PEHE is a weighted linear combination of the mean squared losses for the two underlying supervised problems of learning \( f_0 \) and \( f_1 \) with no covariate shift, where the weights are determined by the extent of the mismatch between the distributions of the treated and control populations, quantified by the Rényi divergence measure. If \( D_n \) is a dataset obtained from a randomized controlled trial \((Q = Q_0 = Q_1)\), then we have \( D_2(Q_0 || Q) = D_2(Q_1 || Q) = 0 \), and the bound boils down to a sum of two supervised learning losses, i.e.,

\[
\psi(\hat{T}) \leq \bar{C} \cdot ||f_0 - \hat{f}_0||_{L^2(P_0)}^2 + \bar{C} \cdot ||f_1 - \hat{f}_1||_{L^2(P_1)}^2.
\]

Since the minimax rate for standard nonparametric regression is \( \|f - \hat{f}\|_2^2 \asymp C_w \cdot n^{-\frac{2}{d_0 + d_1}} \) (Stone, 1982), when \( d_0/\alpha_0 >> d_1/\alpha_1 \), the first-order Taylor approximation for the logarithm of the PEHE in (6) is given by:

\[
\begin{aligned}
\log(\psi(\hat{T})) & \approx D_2(Q_0 || Q) \cdot \log(C_0) - \frac{2\alpha_0}{2\alpha_0 + d_0} \log(n) \\
& + O\left(n^{-\frac{2\alpha_0}{2\alpha_0 + d_0} + \frac{2\alpha_0}{2\alpha_0 + d_0}}\right).
\end{aligned}
\]

(7)

That is, when viewed on a log-log scale, the behavior of the PEHE versus the number of samples can be described as follows. \( \log(\text{PEHE}) \) is a linear function of \( \log(n) \). Selection bias adds a constant offset to \( \log(\text{PEHE}) \), but does not affect its slope, which harms the performance only in the small-sample regime. In the large-sample regime, the slope of \( \log(\text{PEHE}) \), which depends solely on the smoothness and sparsity of the response surfaces, dominates the performance, and selection bias becomes less of a problem. Figure 1 depicts the PEHE in (7) on a log-log scale.

4. CATE Estimation using Non-Stationary Gaussian Process Regression

In this Section, we build on the analyses conducted in Section 3 to design a practical algorithm for CATE estimation.

4.1. Non-Stationary Gaussian Process Priors

We specify the prior \( \Pi(\varphi_{\beta_0}, \varphi_{\beta_1}) \) as a Gaussian process (GP) over functions of the form \( g : \mathcal{X} \times \{0, 1\} \rightarrow \mathbb{R} \), with a kernel \( K_\beta \), and a hyperparameter set \( \beta \) as follows:

\[
g \sim \mathcal{G}P(0, K_\beta(z, z')).
\]

(8)

where \( z = (x, w) \in \mathcal{X} \times \{0, 1\} \), and \( f_\omega(x) = g(x, w) \). The kernel \( K_\beta \) specifies the bases \( \varphi_{\beta_0} \) and \( \varphi_{\beta_1} \) through its induced canonical feature map \( K_\beta(., z) \) (Rasmussen & Williams, 2006; Alvarez et al., 2012). As pointed out in remark 1, the treatment assignment variable \( w \) should encode the different relevant dimensions and smoothness levels of \( f_0 \) and \( f_1 \). Thus, we model \( K_\beta \) as a non-stationary kernel that depends explicitly on \( w \) as follows:

\[
K_\beta(z, z') = \Gamma(w, w') \cdot k_0^w(x, x'),
\]

\[
k_\beta(x, x') = [k_{\beta_0}(x, x'), k_{\beta_1}(x, x'), k_{\beta_0}(x, x') + k_{\beta_1}(x, x')],
\]

\[
\Gamma(w, w') = [\Gamma_0(w, w'), \Gamma_1(w, w'), 1 - \Gamma_0(w, w') - \Gamma_1(w, w')],
\]

where \( \Gamma_0(w, w') = (1 - w)(1 - w') \), \( \Gamma_1(w, w') = w \cdot w' \), and \( k_{\beta_0}(x, x') \) is a Matérn kernel with a length-scale parameter...
\( \beta_w \), for \( w \in \{0, 1\} \). The kernel defined above ensures that any covariance matrix induced by points in \( \mathcal{X} \times \{0, 1\} \) is positive definite. Variable selection is implemented by using the automatic relevance determination version of the Matérn kernel (Rasmussen & Williams, 2006). The non-stationarity of \( K_\delta \) allows setting different length-scales and relevant variables for the marginal priors on \( f_0 \) and \( f_1 \) while sharing data between the two spaces, i.e.

\[
K_\delta((x, w), (x', w')) = k_{\beta_w}(x, x'), \quad w \in \{0, 1\},
K_\delta((x, w), (x', w')) = k_{\beta_0}(x, x') + k_{\beta_1}(x, x'), \quad w \neq w'.
\] (9)

That is, all draws from the prior give Matérn sample paths with different smoothness levels (\( \beta_0 \) and \( \beta_1 \)) for \( f_0 \) and \( f_1 \), respectively, and the correlations between the paths are captured via the kernel mixture \( k_{\beta_0}(x, x') + k_{\beta_1}(x, x') \).

Note that draws from a Matérn prior with length-scale \( \beta \) are almost surely \( \beta \)-Hölder for all \( \beta \leq \beta \) (Vaart & Zanten, 2011). Thus, \( G\mathcal{P}(0, K_\delta) \) specifies a \( \beta \)-Hölder ball as an a priori regularity class for response surface \( f_w \), \( w \in \{0, 1\} \).

In the following Theorem, we show that point estimators induced by the prior \( G\mathcal{P}(0, K_\delta) \) can achieve the optimal minimax rate in Theorem 1.

**Theorem 2.** Suppose that the \( d_w \) relevant features for \( f_w \) are known a priori for \( w \in \{0, 1\} \). If \( f_0 \in H^{\alpha_0}, f_1 \in H^{\alpha_1}, \Pi = G\mathcal{P}(0, K_\delta) \), and \( T = \mathbb{E}[T | \mathcal{D}_n] \), then we have that

\[
\psi(\hat{T}) \leq \alpha_0(\psi(\Pi))^{\alpha_0(\Pi)} + \frac{1}{n} \frac{1}{\alpha_1(\Pi)^{\alpha_1(\Pi)}},
\]

whenever \( \min\{\alpha_0, \alpha_1, \beta_0, \beta_1\} \geq d/2 \). \( \square \)

Note that posterior consistency holds for all combinations of \( (\alpha_0, \alpha_1, \beta_0, \beta_1) \) since the support of the Matérn prior is the space of bounded continuous functions\(^1\). The bound in Theorem 2 can be shown to be tight using the results in (Castillo, 2008). Theorem 2 says that the posterior induced by the prior \( G\mathcal{P}(0, K_\delta) \) contracts around the true CATE function at the optimal rate given in Theorem 1 provided that the following matching condition is met:

\[
\beta_v = \alpha_v, \\
\alpha_v \frac{d_v}{\alpha_v} \leq \beta_{1-v} \leq \alpha_{1-v} + \alpha_v \frac{d_v}{\alpha_v} = \frac{d_{1-v}}{2},
\] (10)

where \( v = 1 \) if \( d_1/\alpha_1 > d_0/\alpha_0 \), and \( v = 0 \) otherwise. The condition in (10) implies that achieving the optimal rate (steepest slope in Figure 1) via the non-stationary GP prior in Section 4.1 is only a matter of hyperparameter tuning: the smoothness of the prior needs to match the smoothness of the “more complex” of the two response surfaces. Note that Theorem 2 implies that we do not need to handle selection bias in order to achieve the optimal rate, which is consistent with the earlier discussion in remark 2.

### 4.2. Doubly-Robust Hyperparameters

Theorem 2 says that the optimal minimax rate for CATE estimation can be achieved by satisfying the smoothness matching condition in (10). However, in practice, the smoothness levels of the true response functions are unknown and need to be learned from the data. Moreover, since selection bias is impactful in small-sample regimes, ignoring it may lead to a poor generalization performance when the size of \( \mathcal{D}_n \) is small. In this Section, we propose a hyperparameter optimization algorithm that accounts for selection bias while ensuring minimax-optimality in the large-sample limit.

Previous works tend to adjust for selection bias “mechanically” using variants of importance sampling approaches based on inverse-propensity-weighting (IPW) (Sugiyama et al., 2007; Shimodaira, 2000), and kernel mean matching (Huang et al., 2007), or by learning a “balanced representation” of treated and control populations (Li & Fu, 2017). We do not attempt to explicitly adjust for selection bias using ad-hoc approaches, and rather seek the “informationally optimal” estimator of the PEHE. That is, we seek the most efficient (unbiased) estimator \( \hat{\psi}^*(\hat{T}) \) of \( \psi(T) \), which satisfies an analog of the Cramér-Rao bound (information-inequality) in parametric estimation, i.e. \( \text{Var}[\hat{\psi}^*(\hat{T})] \leq \text{Var}[\hat{\psi}(T)] \), for any estimator \( \hat{\psi}(T) \).

Classical Cramér-Rao bounds do not apply to estimators of the form \( \hat{\psi}^*(\hat{T}) \), since such estimators are functionals of nonparametric objects. There are, however, analogous information inequalities for nonparametric estimation, including Bhattacharya’s variance bound (Bhattacharya, 1946), and its generalization due to Bickel (Bickel et al., 1998). We proceed by realizing that the PEHE \( \psi(T) \) is simply a functional that belongs to the doubly-robust class of functionals analyzed by Robins in (Robins et al., 2008).

Thus, one can construct the “most” efficient estimator of \( \psi(T) \) using the most efficient influence function of \( \psi(T) \) as follows (Robins et al., 2008; Robins, 2004):

\[
\hat{\psi}^*(\hat{T}) = \sum_{i=1}^{n} \left( \frac{W_i(1-p(X_i)) - p(X_i)}{p(X_i)q(X_i)(1-p(X_i))} \right)^2.
\]

The derivation of the estimator above can be found in Theorem 9 in (Robins, 2004) and Section 5 in (Robins et al., 2008). When the propensity function \( p(.) \) is known, this estimator approximate the PEHE at its optimal minimax rate. We estimate \( p(.) \) via standard kernel density estimation methods. It can be easily shown using the results in (Dudoit & van der Laan, 2005) that when using the estimator above to tune the GP hyperparameters via cross-validation, then the learned length-scale parameters will satisfy the matching condition for minimax optimality.
5. Experiments

In this Section, we check the validity of our analyses using a synthetic simulation setup (Subsection 5.1), and then evaluate the performance of our proposed model using data from a real-world clinical trial with simulated potential outcomes (Subsection 5.2). We will use the acronym NSGP to refer to the non-stationary GP model proposed in Section 4.

5.1. Learning Brownian Response Surfaces

5.1.1. Synthetic Model

Let $\mathcal{X} = [0, 1]$, and define a $\kappa$-fold integrated Brownian motion $B_{\kappa}, \kappa \in \mathbb{N}_{+}$, on $\mathcal{X}$ as follows:

$$ B_{\kappa}(x) = \int_0^x \int_0^{x_1} \cdots \int_0^{x_{\kappa-1}} B_0(x_1) \, dx_1 \, dx_2 \cdots \, dx_{\kappa}, $$

where $B_0(.)$ is a standard Brownian motion (Wiener process). Sample paths of $B_0$ are almost surely Hölder regular with exponent $\frac{1}{2}$ (Karatzas & Shreve, 2012). Since $B_{\kappa}(x)$ is almost surely non-differentiable everywhere in $\mathcal{X}$, then sample paths of $B_{\kappa}(x)$ are Hölder with exponent $\kappa + \frac{1}{2}$, i.e., $B_{\kappa} \in H^{\kappa + \frac{1}{2}}$ with probability 1. Therefore, when the true response surfaces are $\kappa$-fold integrated Brownian paths, the optimality and achievability results in Theorems 1 and 2 should hold. To this end, we simulate the true response surfaces $f_0 \in H^{\alpha_0}$ and $f_1 \in H^{\alpha_1}$ as $f_0 \sim B_{\alpha_0 - \frac{1}{2}}$ and $f_1 \sim B_{\alpha_1 - \frac{1}{2}}$, where we set $\alpha_0 = 2.5$ and $\alpha_1 = 5.5$.

The propensity score is modeled as a parametrized logistic function $p(x \mid \eta) = \frac{1 + e^{-\eta(x - \frac{1}{2})}}{1 + e^{-\eta(x - \frac{1}{2})}}^{-1}$, where $\eta \in \mathbb{R}$ is a parameter that determines the severity of selection bias. For a pair of fixed Brownian paths $f_0$ and $f_1$, synthetic observational samples $(X_i, W_i, Y_i^{(W_i)})$ are generated as follows: $X_i \sim \text{Uniform}[0, 1], W_i \sim \text{Bernoulli}(p(x \mid \eta))$, and $Y_i^{(W_i)} \sim f_{W_i} + \mathcal{N}(0, \sigma^2)$, where $\sigma^2 = 0.1$.

5.1.2. Experiments and Results

Using the setup in Section 5.1.1, we conducted the following Monte Carlo simulations to verify our theoretical findings and highlight the merits of our NSGP model.

- **Verifying Theorems 1 and 2:** In order to check the validity of the results of Theorems 1 and 2, we use a NSGP Matérn prior $GP(0, K_\beta)$, with length-scale parameters $\beta_0$ and $\beta_1$ that are matched exactly with the regularities of the Brownian paths $f_0$ and $f_1$ (i.e. $\beta_0 = 2.5$ and $\beta_1 = 5.5$). According to Theorem 1, the optimal rate for estimating the CATE $T = f_1 - f_0$ is $n^{-\frac{1}{2}}$, and from Theorem 2, the NSGP with $\beta_0 = 2.5$ and $\beta_1 = 5.5$ should achieve that rate.

Figure 2a provides a scatter-plot for the PEHE achieved by the NSGP with respect to the number of samples on a log-log scale for different settings of $\eta$. We fit a linear regression model that describes the PEHE behavior in the log-log scale. We found the slope of the linear fit to be 0.8437, which is very close$^2$ to the slope of $\frac{1}{2}$ = 0.833 predicted by Theorem 1. Moreover, by changing the magnitude of $\eta$ from 0 to $\frac{1}{2}$, the PEHE curve did not exhibit any significant change in its slope, and was only moved upwards by a constant offset. On the contrary, Figure 2b shows the PEHE behavior when the NSGP prior is over-smoothed ($\beta_0 > \alpha_0$) for $\eta = 0$: as predicted by Theorem 2, learning becomes sluggish (slopes become less steep) as $\beta_0$ increase since the matching condition in (10) does not hold any more.

- **NSGPs do not leave any money on the table:** In this experiment, we show that the different components of the NSGP model allow it to perform well in small and large sample regimes. We set a strong selection bias of $\eta = \frac{1}{2}$ and compare the log(PEHE) characteristic of NSGP with a model that uses the same non-stationary kernel as NSGP, and another model that uses a standard stationary kernel, but both models are tuned using marginal likelihood maximization. As we can see in Figure 2c, the model with the non-stationary kernel achieves the same learning rate as NSGP, but exhibits a large offset as it does not account for selection bias, whereas the stationary model fails to learn the smoothness of the rougher Brownian motion since it assigns the same length-scale to both surfaces, and hence it over-smooths the prior, achieving a suboptimal rate.

$^2$The minor discrepancy is a result of the residual error in the linear regression fit.
5.2. The Infant Health and Development Program

We evaluated the performance of the NSGP model presented in Section 4.1 using the standard semi-synthetic experimental setup designed by Hill in (Hill, 2011). We report a state-of-the-art result in this setup, and draw connections between our experimental results and our analyses.

5.2.1. Data and Benchmarks

The Infant Health and Development Program (IHDP) is an interventional program intended to enhance the health of premature infants (Hill, 2011). (Hill, 2011) extracted features and treatment assignments from a real-world clinical trial, and introduced selection bias to the data artificially by removing a subset of the patients. The potential outcomes are simulated according to the standard non-linear "Response Surface B" setting in (Hill, 2011). The dataset comprised 747 subjects, with 25 features for each subject. Our experimental setup is identical to (Hill, 2011; Johanson et al., 2016; Shalit et al., 2017; Alaa & van der Schaar, 2017): we run 1000 experiments in which we compute the in-sample and out-of-sample √PEHE (with 80/20 training/testing splits), and report average results in Table 1.

We compared the performance of NSGP with a total of 23 CATE estimation benchmarks. We considered: tree-based algorithms (BART (Hill, 2011), Causal forests (Wager & Athey, 2017)), Bayesian causal forests (Hahn et al., 2017), methods based on deep learning (CFR Wass., CFR MMD, BNN, TARnet (Shalit et al., 2017)), multivariate additive regression splines (MARS (Powers et al., 2017)), Gaussian processes (CMGP) (Alaa & van der Schaar, 2017), nearest neighbor matching (k-NN), propensity score matching (PSM), and targeted maximum likelihood (TMLE) (Porter et al., 2011). We also compared a number of T-learners and S-learners as in (Künzel et al., 2017), using a variety of baseline machine learning algorithms (DNN stands for deep networks and OLS stands for linear regression).

### Table 1. Simulation results for the IHDP dataset. The values reported correspond to the average PEHE (±95% confidence intervals).

<table>
<thead>
<tr>
<th>[Q1]</th>
<th>[Q2]</th>
<th>Model</th>
<th>In-sample √PEHE</th>
<th>Out-of-sample √PEHE</th>
<th>[Q1]</th>
<th>[Q2]</th>
<th>Model</th>
<th>In-sample √PEHE</th>
<th>Out-of-sample √PEHE</th>
</tr>
</thead>
<tbody>
<tr>
<td>✓</td>
<td>✓</td>
<td>NSGP</td>
<td>0.51 ± 0.013</td>
<td>0.64 ± 0.030</td>
<td>✓</td>
<td></td>
<td>T-XGBoost</td>
<td>1.46 ± 0.081</td>
<td>1.98 ± 0.152</td>
</tr>
<tr>
<td>✓</td>
<td>✓</td>
<td>SGP</td>
<td>0.95 ± 0.021</td>
<td>1.21 ± 0.052</td>
<td>✓</td>
<td></td>
<td>S-XGBoost</td>
<td>2.97 ± 0.211</td>
<td>3.04 ± 0.216</td>
</tr>
<tr>
<td>✓</td>
<td>✓</td>
<td>CMGP</td>
<td>0.61 ± 0.011</td>
<td>0.76 ± 0.012</td>
<td>✓</td>
<td></td>
<td>T-AdaBoost</td>
<td>2.40 ± 0.177</td>
<td>2.79 ± 0.212</td>
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<tr>
<td>✓</td>
<td></td>
<td>TARNet</td>
<td>0.88 ± 0.021</td>
<td>0.95 ± 0.025</td>
<td>✓</td>
<td></td>
<td>S-AdaBoost</td>
<td>4.53 ± 0.317</td>
<td>4.56 ± 0.312</td>
</tr>
<tr>
<td>✓</td>
<td></td>
<td>BNN</td>
<td>2.21 ± 0.115</td>
<td>2.15 ± 0.125</td>
<td>✓</td>
<td></td>
<td>T-OLS</td>
<td>1.85 ± 0.107</td>
<td>1.94 ± 0.122</td>
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<tr>
<td>✓</td>
<td></td>
<td>CFR Wass.</td>
<td>0.71 ± 0.018</td>
<td>0.76 ± 0.032</td>
<td>✓</td>
<td></td>
<td>S-OLS</td>
<td>5.06 ± 0.357</td>
<td>5.05 ± 0.352</td>
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<tr>
<td>✓</td>
<td></td>
<td>CFR MMD</td>
<td>0.73 ± 0.021</td>
<td>0.78 ± 0.022</td>
<td>✓</td>
<td></td>
<td>T-DNN</td>
<td>3.36 ± 0.137</td>
<td>3.46 ± 0.142</td>
</tr>
<tr>
<td>✓</td>
<td></td>
<td>T-Random Forest</td>
<td>1.41 ± 0.071</td>
<td>2.21 ± 0.162</td>
<td>✓</td>
<td></td>
<td>S-DNN</td>
<td>3.56 ± 0.217</td>
<td>3.64 ± 0.212</td>
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<tr>
<td>✓</td>
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<td>S-Random Forest</td>
<td>2.72 ± 0.241</td>
<td>2.91 ± 0.252</td>
<td>✓</td>
<td></td>
<td>MARS</td>
<td>1.66 ± 0.106</td>
<td>1.74 ± 0.112</td>
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<tr>
<td>✓</td>
<td></td>
<td>Causal Forest</td>
<td>2.41 ± 0.141</td>
<td>2.82 ± 0.181</td>
<td>✓</td>
<td></td>
<td>k-NN</td>
<td>2.69 ± 0.177</td>
<td>4.06 ± 0.212</td>
</tr>
<tr>
<td>✓</td>
<td></td>
<td>BART</td>
<td>2.00 ± 0.141</td>
<td>2.22 ± 0.151</td>
<td>✓</td>
<td></td>
<td>PSM</td>
<td>4.92 ± 0.312</td>
<td>4.92 ± 0.312</td>
</tr>
<tr>
<td>✓</td>
<td></td>
<td>BCF</td>
<td>1.31 ± 0.061</td>
<td>1.71 ± 0.102</td>
<td>✓</td>
<td></td>
<td>TMLE</td>
<td>5.27 ± 0.357</td>
<td>5.27 ± 0.352</td>
</tr>
</tbody>
</table>

5.2.2. Results and Conclusions

As we can see in Table 1, the proposed NSGP model significantly outperforms all competing benchmarks. The combined benefit of the two components of an NSGP (non-stationary kernel and doubly-robust hyperparameters) is highlighted by comparing its performance to a vanilla SGP (stationary GP) with marginal likelihood maximization. The gain with respect to such a model is a 2-fold improvement in the PEHE.

Because the IHDP dataset has a “moderate” sample size, both selection bias and learning rate seem to impact the performance. Thus, our method took advantage of having addressed modeling questions [Q1] and [Q2] appropriately by being both “rate-optimal” and “bias-aware”.

The check marks in columns [Q1] and [Q2] designate methods that address modeling questions [Q1] and [Q2] “appropriately” in the light of the analysis presented in Section 3. Methods with [Q1] checked use a regression structure with “outcome-specific” hyperparameters, and methods with [Q2] checked adjust for selection bias. A general observation is that the structure of the regression model seem to matter much more than the strategy for handling selection bias. This is evident from the fact that the TAR-net model (does not handle bias but models outcomes separately) significantly outperforms BNN (handles bias but uses a single-surface model (Shalit et al., 2017)), and that all T-learners (models 2 separate response surfaces) outperformed their S-shaped counterparts (models a single surface). For parametric models, such as OLS, the issue of selecting the right regression structure is even more crucial.

To sum up, the results in Table 1 imply that selecting the right regression structure is crucial for rate-optimality in sufficiently large dataset, whereas handling selection bias provides an extra bonus. In Table 1, methods that address both [Q1] and [Q2] (NSGP, CMGP, and CFR. Wass and MMD) displayed a superior performance.
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Limits of Estimating Heterogeneous Treatment Effects


