The state learner

- a super learner for right-censored data

Anders Munch

Section of Biostatistics, University of Copenhagen, Copenhagen, Denmark

E-mail: a.munch@sund.ku.dk

Thomas A. Gerds

Section of Biostatistics, University of Copenhagen, Copenhagen, Denmark

Summary. In survival analysis, prediction models are needed as stand-alone tools and in applications of causal inference to estimate nuisance parameters. The super learner is a machine learning algorithm which combines a library of prediction models into a meta learner based on cross-validated loss. In right censored data, loss needs to be estimated somehow. We introduce the state learner, a new super learner for survival analysis, which simultaneously evaluates libraries of prediction models for the event of interest and the censoring distribution. The state learner can be applied to all types of survival models, works in the presence of competing risks, and does not require a single pre-specified estimator of the conditional censoring distribution. We establish an oracle inequality for the state learner and investigate its performance through numerical experiments. We illustrate how the state learner allows us to estimate causal effects in a competing risks setting without having to pre-specify models for neither the cause-specific hazards of interest nor the censoring distribution.

Keywords: Competing risks, cross-validation, loss based estimation, right-censored data, super learner

1. Introduction

Medical decision making is traditionally based on the discrepancy between intuitive judgment and the law of chance (Redelmeier et al., 1995). We are concerned with the statistical models that provide probabilistic predictions about the medical future of a patient. We have two different applications of these models in mind. One is as a standalone tool where the model provides information for a freshly diagnosed patient. The other is as input to an algorithm which estimates a causal target parameter such as an average treatment effect. The models are learned on data from previous patients. A super learner is a machine learning algorithm that combines a finite set of learners into a meta learner by estimating prediction performance in hold-out samples using a pre-specified loss function (van der Laan et al., 2007). When the aim is to make a prediction model, super learners combine strong learners, such as Cox regression models and random survival forests (Gerds and Kattan, 2021).

While the general idea of combining strong learners based on cross-validation data stems from earlier work (Wolpert, 1992; Breiman, 1996), the name super learner is

2 Munch et al.

justified by an oracle inequality (van der Laan and Dudoit, 2003; van der Vaart et al., 2006): The super learner is guaranteed to perform almost as well as the model which minimises the expected performance, i.e., the model we would select if we could evaluate the prediction performance in an infinite hold-out sample. The choice of the loss function for super learning in survival analysis deserves a closer look. Existing super learner algorithms for right-censored data use partial log-likelihood loss or inverse probability of censoring weighted loss (Polley and van der Laan, 2011; Keles et al., 2004; Golmakani and Polley, 2020; Westling et al., 2021). The use of the partial log-likelihood loss, however, restricts the class of learners and excludes for example simple Kaplan-Meier based learners and also more complex random survival forest algorithms. For this reason Golmakani and Polley (2020) restrict their learners to Cox proportional hazard models. A lesser known fact is that a super learner constructed with the negative partial loglikelihood loss implicitly depends on the censoring distribution (Hjort, 1992; Whitney et al., 2019). Inverse probability of censoring weighted loss has been studied for decades (Graf et al., 1999; Hothorn et al., 2006; Gerds and Schumacher, 2006). A disadvantage of inverse probability of censoring weighted loss estimators is that they require a prespecified model for the censoring distribution. Recently Westling et al. (2021) proposed a procedure which iterates between super learning of the censoring distribution and super learning of the event time distribution. We pursue a different route.

In this article we define the state learner, a new super learner for right-censored data, which simultaneously estimates the expected loss of learners of the event time distribution and the censoring distribution. The loss function which is used to define the state learner is based on the observed variables, that is the minimum and the order of the censoring time and the event time.

The state learner can be applied to all types of survival models, works in the presence of competing risks, and does not require a single pre-specified estimator of the conditional censoring distribution. To analyze the theoretical properties of the state learner we focus on the discrete super learner which combines the library of learners by picking the one that minimises the cross-validated loss (van der Laan et al., 2007). In the presence of competing risks, our algorithm uses separate libraries to model the cumulative hazard functions of the competing risks and the censoring distribution. We show that the oracle selector of the state learner is consistent if all libraries contain a consistent learner and prove a finite sample oracle inequality. A result of the state learner algorithm is a model which predicts the probability of an event based on covariates in the presence of competing risks (Gerds and Kattan, 2021). The other application is in targeted learning where conditional event probabilities occur as high-dimensional nuisance parameters which need to be estimated at a certain rate (van der Laan and Rose, 2011; Rytgaard et al., 2021; Rytgaard and van der Laan, 2022). We show for a targeted estimator, which uses the state learner to estimate nuisance parameters, that a second order product structure holds for the asymptotic bias term.

The article is organised as follows. We introduce our notation and framework in Section 2. In Section 3 we define general super learning for right-censored data, and in Section 4 we review existing methods. Section 5 introduces our novel proposal, the state learner, and Section 6 provides theoretical guarantees for the state learner. In Section 7 we discuss the use of the state learner in the context of targeted learning. We report

results of our numerical experiments in Section 8 and analyze a prostate cancer data set in Section 9. Section 10 contains a discussion of the merits and limitations of our proposal. Appendices A and B contain proofs.

2. Notation and framework

In a competing risk framework (Andersen et al., 2012), let T be a time to event variable, $D \in \{1,2\}$ the cause of the event, and $X \in \mathcal{X}$ a vector of baseline covariates taking values in a bounded subset $\mathcal{X} \subset \mathbb{R}^p$, $p \in \mathbb{N}$. Let $\tau < \infty$ be the maximal length of follow-up. We use \mathcal{Q} to denote the collection of all probability measures on $[0,\tau] \times \{1,2\} \times \mathcal{X}$ such that $(T,D,X) \sim Q$ for some unknown $Q \in \mathcal{Q}$. For $j \in \{1,2\}$, the cause-specific conditional cumulative hazard functions are defined by $\Lambda_j : [0,\tau] \times \mathcal{X} \to \mathbb{R}_+$ such that

$$\Lambda_j(t\mid x) = \int_0^t \frac{Q(T\in \mathrm{d} s, D=j\mid X=x)}{Q(T\geq s\mid X=x)}.$$

For ease of presentation we assume throughout that $\Lambda_j(\cdot \mid x)$ is continuous for all x and j. This is not a limitation as everything caries over directly to the general case. We denote by S the conditional event-free survival function:

$$S(t \mid x) = \exp\left\{-\Lambda_1(t \mid x) - \Lambda_2(t \mid x)\right\}. \tag{1}$$

Let \mathcal{M}_{τ} denote the space of all conditional cumulative hazard functions on $[0,\tau] \times \mathcal{X}$. Any distribution $Q \in \mathcal{Q}$ can be characterised by

$$Q(dt, j, dx) = \{S(t - | x)\Lambda_1(dt | x)H(dx)\}^{1_{\{j=1\}}}$$
$$\{S(t - | x)\Lambda_2(dt | x)H(dx)\}^{1_{\{j=2\}}},$$

where $\Lambda_j \in \mathcal{M}_{\tau}$ for j = 1, 2 and H is the marginal distribution of the covariates.

We consider the right-censored setting in which we observe data O = (T, D, X), where $\tilde{T} = \min(T, C)$ for a right-censoring time C, $\Delta = \mathbbm{1}\{T \leq C\}$, and $\tilde{D} = \Delta D$. Let \mathcal{P} denote a set of probability measures on the sample space $\mathcal{O} = [0, \tau] \times \{0, 1, 2\} \times \mathcal{X}$ such that $O \sim P$ for some unknown $P \in \mathcal{P}$. We assume that the event times and the censoring times are conditionally independent given covariates, $T \perp C \mid X$. This implies that any distribution $P \in \mathcal{P}$ is characterised by a distribution $Q \in \mathcal{Q}$ and a conditional cumulative hazard function for C given X (c.f., Begun et al., 1983; Gill et al., 1997). We use $\Gamma \in \mathcal{M}$ to denote the conditional cumulative hazard function for censoring. For ease of presentation we now also assume that $\Gamma(\cdot \mid x)$ is continuous for all x. We let $G(t \mid x) = \exp\{-\Gamma(t \mid x)\}$ denote the survival function of the conditional censoring distribution. The distribution is characterized by

$$P(dt, j, dx) = \{G(t - | x)S(t - | x)\Lambda_{1}(dt | x)H(dx)\}^{\mathbb{1}\{j=1\}}$$

$$\{G(t - | x)S(t - | x)\Lambda_{2}(dt | x)H(dx)\}^{\mathbb{1}\{j=2\}}$$

$$\{G(t - | x)S(t - | x)\Gamma(dt | x)H(dx)\}^{\mathbb{1}\{j=0\}}$$

$$= \{G(t - | x)Q(dt, j, dx)\}^{\mathbb{1}\{j\neq0\}}$$

$$\{G(t - | x)S(t - | x)\Gamma(dt | x)H(dx)\}^{\mathbb{1}\{j=0\}} .$$
(2)

Hence, we may write $\mathcal{P} = \{P_{Q,\Gamma} : Q \in \mathcal{Q}, \Gamma \in \mathcal{G}\}$ for some $\mathcal{G} \subset \mathcal{M}_{\tau}$. We also have H-almost everywhere

$$P(\tilde{T} > t \mid X = x) = S(t \mid x)G(t \mid x) = \exp\{-\Lambda_1(t \mid x) - \Lambda_2(t \mid x) - \Gamma(t \mid x)\}.$$

We further assume that there exists $\kappa < \infty$ such that $\Lambda_j(\tau - \mid x) < \kappa$, for $j \in \{1, 2\}$, and $\Gamma(\tau - \mid x) < \kappa$ for almost all $x \in \mathcal{X}$. Note that this implies that $G(\tau - \mid x)$ is bounded away from zero for almost all $x \in \mathcal{X}$. Under these assumptions, the conditional cumulative hazard functions Λ_j and Γ can be identified from P by

$$\Lambda_j(t \mid x) = \int_0^t \frac{P(\tilde{T} \in \mathrm{d}s, \tilde{D} = j \mid X = x)}{P(\tilde{T} \ge s \mid X = x)},\tag{3}$$

$$\Gamma(t \mid x) = \int_0^t \frac{P(\tilde{T} \in \mathrm{d}s, \tilde{D} = 0 \mid X = x)}{P(\tilde{T} \ge s \mid X = x)}.$$
 (4)

Thus, we can consider Λ_j and Γ as operators which map from \mathcal{P} to \mathcal{M} .

3. The concept of super learning

In survival analysis, a super learner can be used to estimate a parameter Ψ which can be identified from the observed data distribution $P \in \mathcal{P}$. In this section, to introduce the discrete super learner and the oracle learner, we consider estimation of the function-valued parameter $\Psi : \mathcal{P} \to \mathcal{M}$, given by $\Psi(P) = \Lambda_j$. This parameter is identified via equation (3) on $[0, \tau]$.

As input to the super learner we need a data set $\mathcal{D}_n = \{O_i\}_{i=1}^n$ of i.i.d. observations from $P \in \mathcal{P}$ and a finite collection of candidate learners \mathcal{A} . Each learner $a \in \mathcal{A}$ is a map $a : \mathcal{O}^n \to \mathcal{M}_{\tau}$ which takes a data set as input and returns an estimate $a(\mathcal{D}_n) \in \mathcal{M}_{\tau}$ of Λ_j . In what follows, we use the short-hand notation $P[f] = \int f(o)P(\mathrm{d}o)$. A super learner evaluates the performance of $a \in \mathcal{A}$ with a loss function $L : \mathcal{M}_{\tau} \times \mathcal{O} \to \mathbb{R}_+$ and estimates the expected loss $P[L(a(\mathcal{D}_n), \cdot)]$ using cross-validation. Specifically, the expected loss of $a \in \mathcal{A}$ is estimated by splitting the data set \mathcal{D}_n into K disjoint approximately equally sized subsets $\mathcal{D}_n^1, \mathcal{D}_n^2, \ldots, \mathcal{D}_n^K$ and then calculating the cross-validated loss

$$\hat{R}_n(a; L) = \frac{1}{K} \sum_{k=1}^K \frac{1}{|\mathcal{D}_n^k|} \sum_{O_i \in \mathcal{D}_n^k} L(a(\mathcal{D}_n^{-k}), O_i), \quad \text{with} \quad \mathcal{D}_n^{-k} = \mathcal{D}_n \setminus \mathcal{D}_n^k.$$

The subset \mathcal{D}_n^{-k} is referred to as the k'th training sample, while \mathcal{D}_n^k is referred to as the k'th test or hold-out sample. The discrete super learner is defined as

$$\hat{a}_n = \operatorname*{argmin}_{a \in \mathcal{A}} \hat{R}_n(a; L).$$

The oracle learner is defined as the learner that minimises the average loss according to the data-generating distribution P, i.e.,

$$\tilde{a}_n = \operatorname*{argmin}_{a \in \mathcal{A}} \tilde{R}_n(a; L), \quad \text{with} \quad \tilde{R}_n(a; L) = \frac{1}{K} \sum_{k=1}^K P \Big[L\Big(a(\mathcal{D}_n^{-k}), \cdot\Big) \Big].$$

Note that both the discrete super learner and the oracle learner depend on the library of learners and on the number of folds K, and that the oracle learner is a function of the data and the unknown data-generating distribution. However, these dependencies are suppressed in the notation.

4. Existing methods

Machine learning based on right-censored data commonly use the negative partial log-likelihood as loss function (e.g., Li et al., 2016; Yao et al., 2017; Lee et al., 2018; Katzman et al., 2018; Gensheimer and Narasimhan, 2019; Lee et al., 2021; Kvamme and Borgan, 2021). This loss function is however not perfectly suited for super learning. The reason is that it cannot be applied to important survival learners: The Kaplan-Meier estimator, the random survival forest, and the semi-parametric Cox regression model provide cumulative hazard functions that are piece-wise constant in the time argument. Suppose these methods are applied to a learning data set and the aim is to evaluate the partial log-likelihood loss in an independent test data set. Suppose further that the test data set contains event times that were not also seen in the learning data set, as it would naturally happen when the events are observed in continuous time. The problem is that these methods assign zero probability to the test set event times that were not also in the training data set, and hence (the logarithm of zero is infinity) the partial log-likelihood loss in the test set would be infinity.

When a proportional hazards model is assumed, the baseline hazard function can be profiled out of the likelihood (Cox, 1972). The cross-validated partial log-likelihood loss (Verweij and van Houwelingen, 1993) has therefore been suggested as a loss function for super learning which however restricts the library of learners to include only Cox proportional hazards models (Golmakani and Polley, 2020).

Another approach for super learning with right-censored data is to use an inverse probability of censoring weighted (IPCW) loss function (Graf et al., 1999; van der Laan and Dudoit, 2003; Molinaro et al., 2004; Keles et al., 2004; Hothorn et al., 2006; Gerds and Schumacher, 2006; Gonzalez Ginestet et al., 2021). An IPCW loss function is attractive because the associated risk does not depend on the censoring distribution but describes a feature of the population of interested governed by the measure $Q \in \mathcal{Q}$. Similar results can be obtained using censoring unbiased transformations (Fan and Gijbels, 1996; Steingrimsson et al., 2019) or pseudo-values (Andersen et al., 2003; Mogensen and Gerds, 2013; Sachs et al., 2019). All these methods rely on an estimator of the censoring distribution, and their drawback is that this estimator has to be pre-specified. When the data-generating mechanism is complex and not well-understood, pre-specification of the censoring distribution is a challenge.

An attempt to avoid a pre-specified censoring model was proposed independently by Han et al. (2021) and Westling et al. (2021). In both articles, the authors suggest to iterate between learning of Λ and learning of Γ using IPCW loss functions. However, no general theoretical guarantees seem to exist for this procedure, and it has not yet been extended to the situation with competing risks.

5. The state learner

The main idea of the state learner is to jointly use learners of Λ_1 , Λ_2 , and Γ , and the relations in equation (2), to learn a feature of the observed data distribution P. The discrete state learner ranks a tuple of learners of $(\Lambda_1, \Lambda_2, \Gamma)$ based on how well they jointly model the observed data. Risk predictions can then be obtained from Λ_1 and Λ_2 (Benichou and Gail, 1990; Ozenne et al., 2017). To formally introduce the state learner, we define the multi-state process

$$\eta(t) = \mathbb{1}\{\tilde{T} \le t, \tilde{D} = 1\} + 2\mathbb{1}\{\tilde{T} \le t, \tilde{D} = 2\} - \mathbb{1}\{\tilde{T} \le t, \tilde{D} = 0\}, \text{ for } t \in [0, \tau].$$

At time t, we observe that each individual is in one of four mutually exclusive states (Figure 1). The conditional distribution of $\eta(t)$ given X is determined by the function

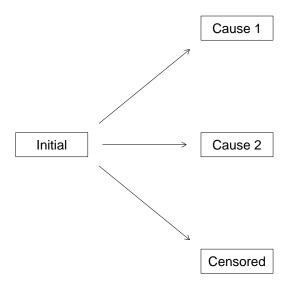


Fig. 1. Illustration of the multi-state process η used by the state learner. Note that 'censored' is a state, hence the process is always observed at any time.

$$F(t, k, x) = P(\eta(t) = k \mid X = x), \text{ for all } t \in [0, \tau], k \in \{-1, 0, 1, 2\}, x \in \mathcal{X}.$$
 (5)

The function F describes the conditional state occupation probabilities of the multistate process η . We construct a super learner for F. The target of this super learner is the function-valued parameter $\Psi(P) = F$ which is identified through equation (5). Under conditional independent censoring each quadruple $(\Lambda_1, \Lambda_2, \Gamma, H)$ characterizes a distribution $P \in \mathcal{P}$, c.f. equation (2), which in turn determines (F, H), and so a learner for F can be constructed from learners of Λ_1 , Λ_2 , and Γ as follows:

$$F(t,1,x) = P(\tilde{T} \le t, \Delta = 1 \mid X = x) = \int_0^t e^{\{-\Lambda_1(s|x) - \Lambda_2(s|x) - \Gamma(s|x)\}} \Lambda_1(\mathrm{d}s \mid x),$$

$$F(t,2,x) = P(\tilde{T} \le t, \Delta = 2 \mid X = x) = \int_0^t e^{\{-\Lambda_1(s|x) - \Lambda_2(s|x) - \Gamma(s|x)\}} \Lambda_2(\mathrm{d}s \mid x),$$

$$F(t,-1,x) = P(\tilde{T} \le t, \Delta = 0 \mid X = x) = \int_0^t e^{\{-\Lambda_1(s|x) - \Lambda_2(s|x) - \Gamma(s|x)\}} \Gamma(\mathrm{d}s \mid x),$$

$$F(t,0,x) = P(\tilde{T} > t \mid X = x) = 1 - F(t,1,x) - F(t,2,x) - F(t,-1,x).$$
(6)

The state learner requires three libraries of learners, \mathcal{A}_1 , \mathcal{A}_2 , and \mathcal{B} , where \mathcal{A}_1 and \mathcal{A}_2 contain learners of the conditional cause-specific cumulative hazard functions Λ_1 and Λ_2 , respectively, and \mathcal{B} contains learners of the conditional cumulative hazard function of the censoring distribution. Based on the Cartesian product of libraries of learners for $(\Lambda_1, \Lambda_2, \Gamma)$ we construct a library \mathcal{F} of learners for F:

$$\mathcal{F}(\mathcal{A}_1, \mathcal{A}_2, \mathcal{B}) = \{ \varphi_{a_1, a_2, b} : a_1 \in \mathcal{A}_1, a_2 \in \mathcal{A}_2, b \in \mathcal{B} \},$$

where in correspondence with the relations in equation (6),

$$\varphi_{a_{1},a_{2},b}(\mathcal{D}_{n})(t,1,x) = \int_{0}^{t} e^{\{-a_{1}(\mathcal{D}_{n})(s|x) - a_{2}(\mathcal{D}_{n})(s|x) - b(\mathcal{D}_{n})(s|x)\}} a_{1}(\mathcal{D}_{n})(ds \mid x),$$

$$\varphi_{a_{1},a_{2},b}(\mathcal{D}_{n})(t,2,x) = \int_{0}^{t} e^{\{-a_{1}(\mathcal{D}_{n})(s|x) - a_{2}(\mathcal{D}_{n})(s|x) - b(\mathcal{D}_{n})(s|x)\}} a_{2}(\mathcal{D}_{n})(ds \mid x),$$

$$\varphi_{a_{1},a_{2},b}(\mathcal{D}_{n})(t,-1,x) = \int_{0}^{t} e^{\{-a_{1}(\mathcal{D}_{n})(s|x) - a_{2}(\mathcal{D}_{n})(s|x) - b(\mathcal{D}_{n})(s|x)\}} b(\mathcal{D}_{n})(ds \mid x),$$

$$\varphi_{a_{1},a_{2},b}(\mathcal{D}_{n})(t,0,x) = 1 - \varphi_{a_{1},a_{2},b}(\mathcal{D}_{n})(t,1,x) - \varphi_{a_{1},a_{2},b}(\mathcal{D}_{n})(t,2,x)$$

$$- \varphi_{a_{1},a_{2},b}(\mathcal{D}_{n})(t,-1,x)$$

To evaluate how well a function F predicts the observed multi-state process we use the integrated Brier score $\bar{B}_{\tau}(F,O) = \int_0^{\tau} B_t(F,O) dt$, where B_t is the Brier score (Brier et al., 1950) at time $t \in [0,\tau]$,

$$B_t(F, O) = \sum_{j=-1}^{2} (F(t, j, X) - \mathbb{1} \{ \eta(t) = j \})^2.$$

Based on a split of a data set \mathcal{D}_n into K disjoint approximately equally sized subsets (see Section 3), each learner $\varphi_{a_1,a_2,b}$ in the library $\mathcal{F}(\mathcal{A}_1,\mathcal{A}_2,\mathcal{B})$ is evaluated using the cross-validated loss,

$$\hat{R}_n(\varphi_{a_1,a_2,b}; \bar{B}_\tau) = \frac{1}{K} \sum_{k=1}^K \frac{1}{|\mathcal{D}_n^k|} \sum_{O_i \in \mathcal{D}_n^k} \bar{B}_\tau \Big(\varphi_{a_1,a_2,b}(\mathcal{D}_n^{-k}), O_i \Big),$$

and the discrete state learner is given by

$$\hat{\varphi}_n = \operatorname*{argmin}_{(a_1, a_2, b) \in \mathcal{A}_1 \times \mathcal{A}_2 \times \mathcal{B}} \hat{R}_n(\varphi_{a_1, a_2, b}; \bar{B}_\tau).$$

6. Theoretical results for the state learner

In this section we establish theoretical guarantees for the state learner. We show that the state learner is consistent if its library contains a consistent learner, and we establish a finite sample inequality for the excess risk of the state learner compared to the oracle. Finally we show that a certain second order structure is preserved when the state learner is used for targeted learning.

6.1. Consistency

Proposition 1 can be derived from the fact that the integrated Brier score (also called the continuous ranked probability score) is a strictly proper scoring rule (Gneiting and Raftery, 2007). This implies that if we minimise the average loss of the integrated Brier score, we recover the parameters of the data-generating distribution. Specifically, this implies that the oracle of a state learner is consistent if the library of learners contains at least one learner that is consistent for estimation of F. Recall that the function F implicitly depends on the data generating probability measure $P \in \mathcal{P}$ but that this was suppressed in the notation. We now make this dependence explicit by writing F_0 for the function which is obtained by substituting a specific $P_0 \in \mathcal{P}$ for P in equation (6). In the following we let $\mathcal{H}_{\mathcal{P}} = \{F_P : P \in \mathcal{P}\}$ where F_P is defined as in equation (5) using the measure $P \in \mathcal{P}$.

Proposition 1. If $P_0 \in \mathcal{P}$ then

$$F_0 = \operatorname*{argmin}_{F \in \mathcal{H}_{\mathcal{P}}} P_0[\bar{B}_{\tau}(F, \cdot)],$$

H-almost surely for any $j \in \{-1,0,1,2\}$ and almost any $t \in [0,\tau]$.

Proof. See Appendix A.

6.2. Oracle inequalities

We establish a finite sample oracle result for the state learner. Our Corollary 1 is in essence a special case of Theorem 2.3 in (van der Vaart et al., 2006). We assume that we split the data into equally sized folds, and for simplicity of presentation we take n to be such that $|\mathcal{D}_n^{-k}| = n/K$ with K fixed. We will allow the number of learners to grow with n and write $\mathcal{F}_n = \mathcal{F}(\mathcal{A}_{1,n}, \mathcal{A}_{2,n}, \mathcal{B}_n)$ as short-hand notation and to emphasise the dependence on n. In the following we let the space $\mathcal{H}_{\mathcal{P}}$ be equipped with the norm $\|\cdot\|_{P_0}$ defined as

$$||F||_{P_0} = \left\{ \sum_{j=-1}^2 \int_{\mathcal{X}} \int_0^{\tau} F(t,j,x)^2 \, \mathrm{d}t H_0(\mathrm{d}x) \right\}^{1/2}. \tag{7}$$

COROLLARY 1. For all $P_0 \in \mathcal{P}$, $n \in \mathbb{N}$, $k \in \{1, ..., K\}$, and $\delta > 0$,

$$\mathbb{E}_{P_0} \left[\| \hat{\varphi}_n(\mathcal{D}_n^{-k}) - F_0 \|_{P_0}^2 \right] \le (1 + 2\delta) \, \mathbb{E}_{P_0} \left[\| \tilde{\varphi}_n(\mathcal{D}_n^{-k}) - F_0 \|_{P_0}^2 \right] \\ + (1 + \delta) 16 K \tau \left(13 + \frac{12}{\delta} \right) \frac{\log(1 + |\mathcal{F}_n|)}{n}.$$

Proof. See Appendix A.

Corollary 1 has the following asymptotic consequences.

COROLLARY 2. Assume that $|\mathcal{F}_n| = O(n^q)$, for some $q \in \mathbb{N}$ and that there exists a sequence $\varphi_n \in \mathcal{F}_n$, $n \in \mathbb{N}$, such that $\mathbb{E}_{P_0} \left[\|\varphi_n(\mathcal{D}_n^{-k}) - F_0\|_{P_0}^2 \right] = O(n^{-\alpha})$, for some $\alpha \leq 1$.

(i) If $\alpha = 1$ then $\mathbb{E}_{P_0} \left[\|\hat{\varphi}_n(\mathcal{D}_n^{-k}) - F_0\|_{P_0}^2 \right] = O(\log(n)n^{-1})$.

(ii) If
$$\alpha < 1$$
 then $\mathbb{E}_{P_0} \left[\| \hat{\varphi}_n(\mathcal{D}_n^{-k}) - F_0 \|_{P_0}^2 \right] = O(n^{-\alpha})$.

Proof. See Appendix A.

6.3. Transience of the second order remainder structure

We here consider the situation where Λ_1 , Λ_2 , and Γ are nuisance parameters and an the estimator of interest is obtained from estimators of these nuisance parameters. An example is an estimator of the cumulative incidence curve, which can be obtained from estimators of Λ_1 and Λ_2 . By equations (3) and (4) and the definition of F, we have

$$\Gamma(t,x) = \int_0^t \frac{F(ds, -1, x)}{F(s, 0, x)}, \quad \text{and} \quad \Lambda_j(t, x) = \int_0^t \frac{F(ds, j, x)}{F(s, 0, x)}, \quad j \in \{1, 2\},$$
 (8)

and thus an estimator based on $\hat{\Lambda}_{1,n}$, $\hat{\Lambda}_{2,n}$, and $\hat{\Gamma}_n$ can also be obtained from an estimator of F using equation (8). A so-called targeted estimator has the key feature that it is asymptotically equivalent to a sum of i.i.d. random variables plus a second order remainder term (van der Laan and Rose, 2011; Hines et al., 2022). For the setting with competing risks, the remainder term is dominated by terms of the form

$$P\left[\int_0^{\tau} w_n(s,\cdot) \hat{M}_{1,n}(s\mid\cdot) \hat{M}_{2,n}(\mathrm{d}s\mid\cdot)\right],\tag{9}$$

where $(\hat{M}_{1,n}, \hat{M}_{2,n})$ is any of the nine combinations of $\hat{M}_{1,n} \in \{[\Gamma - \hat{\Gamma}_n], [\Lambda_1 - \hat{\Lambda}_{1,n}], [\Lambda_2 - \hat{\Lambda}_{2,n}]\}$ and $\hat{M}_{2,n} \in \{[\Gamma - \hat{\Gamma}_n], [\Lambda_1 - \hat{\Lambda}_{1,n}], [\Lambda_2 - \hat{\Lambda}_{2,n}]\}$, and w_n is some data-dependent function with domain $[0,\tau] \times \mathcal{X}$ (van der Laan and Robins, 2003). In particular, a targeted estimator will be asymptotically linear if the 'products' of the estimation errors $\hat{M}_{1,n}$ and $\hat{M}_{2,n}$ in equation (9) are $o_P(n^{-1/2})$. Proposition 2 states that if equation (9) holds for a targeted estimator based on estimators $\hat{\Lambda}_{1,n}$, $\hat{\Lambda}_{2,n}$, and $\hat{\Gamma}_n$, then a similar product structure holds for a targeted estimator based on \hat{F}_n . We state the result for the special case that $\hat{M}_{1,n} = \Gamma - \hat{\Gamma}_n$ and $\hat{M}_{2,n} = \Lambda_1 - \hat{\Lambda}_{1,n}$, but similar results hold for any combinations of $\Gamma - \hat{\Gamma}_n$, $\Lambda_1 - \hat{\Lambda}_{1,n}$, and $\Lambda_2 - \hat{\Lambda}_{2,n}$.

PROPOSITION 2. Assume that $w(s,x) \le c$, $F(s,0,x) \ge 1/c$ and $\hat{F}_n(s,0,x) \ge 1/c$ for some c > 0 for all $s \in [0,\tau]$ and $x \in \mathcal{X}$. Then there are real-valued uniformly bounded

functions w_n^a , w_n^b , w_n^c , and w_n^d with domain $[0,\tau]^2 \times \mathcal{X}$ such that

$$\begin{split} &P_0 \bigg[\int_0^\tau w(s,\cdot) \, \Big\{ \Gamma_0(s,\cdot) - \hat{\Gamma}_n(s,\cdot) \Big\} \, [\Lambda_0 - \hat{\Lambda}_n](\mathrm{d} s,\cdot) \bigg] \\ &= P_0 \bigg[\int_0^\tau \int_0^s w_n^a(s,u,\cdot) [F_0 - \hat{F}_n](u-,0,\cdot) [F_0 - \hat{F}_n](s-,0,\cdot) F_0(\mathrm{d} u,2,\cdot) F_0(\mathrm{d} s,1,\cdot) \bigg] \\ &+ P_0 \bigg[\int_0^\tau \int_0^s w_n^b(s,u,\cdot) [F_0 - \hat{F}_n](u-,0,\cdot) F_0(\mathrm{d} u,2,\cdot) [F_0 - \hat{F}_n](\mathrm{d} s,1,\cdot) \bigg] \\ &+ P_0 \bigg[\int_0^\tau \int_0^s w_n^c(s,u,\cdot) [F_0 - \hat{F}_n](\mathrm{d} u,2,\cdot) [F_0 - \hat{F}_n](s-,0,\cdot) F_0(\mathrm{d} s,1,\cdot) \bigg] \\ &+ P_0 \bigg[\int_0^\tau \int_0^s w_n^d(s,u,\cdot) [F_0 - \hat{F}_n](\mathrm{d} u,2,\cdot) [F_0 - \hat{F}_n](\mathrm{d} s,1,\cdot) \bigg]. \end{split}$$

Proof. See Appendix B.

7. Targeted learning

We are interested in a parameter $\theta \colon \mathcal{Q} \to \Theta$ that expresses a property of the uncensored population governed by the measure $Q \in \mathcal{Q}$. The parameter space Θ can be a subset of \mathbb{R}^d or a subset of a function space, for example a subset of \mathcal{M}_{τ} . In subsection 7.1 we discuss an example from causal inference where θ is the average treatment effect and $\Theta = [-1,1]$. To discuss the role of the state learner for targeted learning we briefly review some results from semiparametric efficiency theory. Extensive reviews and introductions are available elsewhere (e.g., Pfanzagl and Wefelmeyer, 1982; Bickel et al., 1993; van der Laan and Robins, 2003; Tsiatis, 2007; Kennedy, 2016). Under the assumption of conditional independent censoring and positivity, θ is identifiable from \mathcal{P} which means that there exists an operator $\Psi \colon \mathcal{P} \to \Theta$ such that $\theta(Q) = \Psi(P_{Q,\Gamma})$ for all $\Gamma \in \mathcal{M}_{\tau}$. By equation (2) this implies that we may write

$$\theta(Q) = \Psi(P) = \tilde{\Psi}^0(\Lambda_1, \Lambda_2, H),$$

for some operator $\tilde{\Psi}^0$. The state learner provides a ranking of tuples $(a_1, a_2, b) \in \mathcal{A}_1 \times \mathcal{A}_2 \times \mathcal{B}$, and we use $\hat{a}_{1,n}$, $\hat{a}_{2,n}$, and \hat{b}_n to denote the learners corresponding to the discrete state learner $\hat{\varphi}_n$. Letting $H(\mathcal{D}_n)$ denote the empirical measure of $\{X_1, \ldots, X_n\}$, we obtain a simple plug-in estimator of θ :

$$\hat{\Psi}^0(\mathcal{D}_n) = \tilde{\Psi}^0(\hat{a}_{1,n}(\mathcal{D}_n), \hat{a}_{2,n}(\mathcal{D}_n), H(\mathcal{D}_n)). \tag{10}$$

The asymptotic distribution of $\hat{\Psi}^0_{\tau,j}$ is difficult to analyze due to the cross-validated model selection involved in the estimation of the nuisance parameters Λ_1 and Λ_2 . In addition, the estimator will typically have an asymptotic bias that vanishes at a too slow rate. Using tools from semi-parametric efficiency theory, it is possible to construct a so-called targeted or debiased estimator with smaller asymptotic bias and a asymptotic distribution which we know how to estimate (Bickel et al., 1993; van der Laan and Rose, 2011; Chernozhukov et al., 2018). A targeted estimator is based on the efficient influence function for the parameter $\tilde{\Psi}^0$ and relies on an estimator of Γ in addition to

estimators of Λ_1 and Λ_2 . The efficient influence function is a P-zero mean and square integrable function indexed by the nuisance parameters $(\Lambda_1, \Lambda_2, \Gamma)$, which we denote by $\psi(\cdot; \Lambda_1, \Lambda_2, \Gamma)$. The name is justified because any regular asymptotically linear estimator that has ψ as its influence function is asymptotically efficient, meaning that it has smallest asymptotic variance among all regular asymptotically linear estimators.

An example of a targeted estimator is the one-step estimator, defined as

$$\hat{\Psi}_{OS}(\mathcal{D}_n) = \tilde{\Psi}^0(\hat{a}_{1,n}(\mathcal{D}_n), \hat{a}_{2,n}(\mathcal{D}_n), H(\mathcal{D}_n)) + \mathbb{P}_n[\psi(\cdot; \hat{a}_{1,n}(\mathcal{D}_n), \hat{a}_{2,n}(\mathcal{D}_n), \hat{b}_n(\mathcal{D}_n))], (11)$$

where \mathbb{P}_n is the empirical measure of a data set $\{O_i\}_{i=1}^n$. Under suitable regularity conditions we have the following asymptotic expansion of the one-step estimator (Pfanzagl and Wefelmeyer, 1982; van der Laan and Robins, 2003; Fisher and Kennedy, 2021; Kennedy, 2022; Rytgaard et al., 2022),

$$\hat{\Psi}_{\mathrm{OS}}(\mathcal{D}_n) - \Psi(P) = \mathbb{P}_n[\psi(\cdot; \Lambda_1, \Lambda_2, \Gamma)] + \operatorname{Rem}(\hat{\Lambda}_{1,n}, \hat{\Lambda}_{2,n}, \hat{\Gamma}_n, P) + o_P(n^{-1/2}),$$

where the remainder term has the form

$$\operatorname{Rem}(\hat{\Lambda}_{1,n}, \hat{\Lambda}_{2,n}, \hat{\Gamma}_{n}, P) = \mathcal{O}_{P} \Big\{ \|\Lambda_{1} - \hat{\Lambda}_{1,n}\|^{2} + \|\Lambda_{2} - \hat{\Lambda}_{2,n}\|^{2} + \|\Gamma - \hat{\Gamma}_{n}\|^{2} \Big\}, \tag{12}$$

for some suitable norm $\|\cdot\|$, for instance the \mathcal{L}_P^2 -norm. When equation (12) holds and the nuisance parameters Λ_1 , Λ_2 , and Γ are consistently estimated at rate $o_P(n^{-1/4})$, then

$$\sqrt{n}(\hat{\Psi}_{OS}(\mathcal{D}_n) - \Psi(P)) \rightsquigarrow \mathcal{N}(0, P[\psi(\cdot; \Lambda_1, \Lambda_2, \Gamma)^2]), \tag{13}$$

where we use \rightsquigarrow to denote weak convergence (van der Vaart, 2000). In particular, equation (13) and Slutsky's lemma imply that we can obtain asymptotically valid $(1 - \alpha) \cdot 100\%$ confidence intervals by calculating

$$\left[\hat{\Psi}_{\mathrm{OS}}(\mathcal{D}_n) - q_{\alpha/2}\hat{\sigma}(\mathcal{D}_n), \ \hat{\Psi}_{\mathrm{OS}}(\mathcal{D}_n) + q_{\alpha/2}\hat{\sigma}(\mathcal{D}_n)\right],$$

where q_{α} is the $(1 - \alpha)$ -quantile of the standard normal distribution and $\hat{\sigma}(\mathcal{D}_n)^2 = \mathbb{P}_n \Big[\psi(\cdot; \hat{a}_{1,n}(\mathcal{D}_n), \hat{a}_{2,n}(\mathcal{D}_n), \hat{b}_n(\mathcal{D}_n))^2 \Big].$

7.1. Average treatment effect on the absolute risk of an event

In the following we demonstrate how the state learner and the general estimation strategy outlined above can be used to construct an estimator of the cause-specific average treatment effect. We assume that the covariate vector $X \in \mathbb{R}^p$ contains a binary treatment indicator A and a vector of potential confounders, $W \in \mathcal{W} \subset \mathbb{R}^{p-1}$. We use μ to denote the marginal distribution of W and π to denote the conditional probability of treatment,

$$\pi(w) = P(A = 1 \mid W = w).$$

We assume throughout that π is uniformly bounded away from 0 and 1 on \mathcal{W} and that both A and W are fully observed for all individuals. We use a super learner to estimate π (Polley et al., 2023), and we denote this estimator by $\hat{\pi}_n$. We use the empirical measure of $\{W_1, \ldots, W_n\}$ to estimate μ , and denote this estimator by $\hat{\mu}_n$. As parameter of interest

we consider the standardised difference in the absolute risk of an event with cause 1 at time τ :

$$\theta_{\tau}(Q) = \int_{\mathcal{W}} \left\{ \int_{0}^{\tau} S(s - | w, 1) \Lambda_{1}(\mathrm{d}s | w, 1) - \int_{0}^{\tau} S(s - | w, 0) \Lambda_{1}(\mathrm{d}s | w, 0) \right\} \mu(\mathrm{d}w).$$

Under the usual assumptions for causal inference (consistency, positivity, no unmeasured confounding) θ_{τ} can be given the causal interpretation

$$\theta_{\tau}(Q) = P(T^1 \le \tau, D^1 = 1) - P(T^0 \le \tau, D^0 = 1),$$

where (T^a, D^a) , $a \in \{0, 1\}$, denotes potential outcomes (Hernán and Robins, 2020). In this case, the interpretation of θ_{τ} is the difference in the average risk of cause 1 occurring before time τ in the population if everyone had been given treatment (A = 1) compared to if no one had been given treatment (A = 0).

Using equation (1) we may write $\theta_{\tau}(Q) = \tilde{\Psi}_{t}^{0}(\Lambda_{1}, \Lambda_{2}, \mu)$, where

$$\tilde{\Psi}_{t}^{0}(\Lambda_{1}, \Lambda_{2}, \mu) = \int_{\mathcal{W}} \int_{0}^{\tau} e^{-\Lambda_{1}(s-|w,1)-\Lambda_{2}(s-|w,1)} \Lambda_{1}(\mathrm{d}s \mid w, 1) \mu(\mathrm{d}w)
- \int_{\mathcal{W}} \int_{0}^{\tau} e^{-\Lambda_{1}(s-|w,0)-\Lambda_{2}(s-|w,0)} \Lambda_{1}(\mathrm{d}s \mid w, 0) \mu(\mathrm{d}w).$$
(14)

The efficient influence function for the parameter $\tilde{\Psi}_{\tau}$ depends on the set $(\Lambda_1, \Lambda_2, \Gamma, \pi)$ of nuisance parameters. We define

$$\omega_{a}(A, W; \pi) = \frac{\mathbb{1}\{A = a\}}{\pi(W)^{a}(1 - \pi(W))^{1 - a}},$$

$$g(t, A, W; \Lambda_{1}, \Lambda_{2}) = \int_{0}^{t} e^{-\Lambda_{1}(s - |W, A) - \Lambda_{2}(s - |W, A)} \Lambda_{1}(\mathrm{d}s \mid W, A),$$

$$M_{i}(\mathrm{d}t \mid A, W; \Lambda_{i}) = N_{i}(\mathrm{d}t) - \mathbb{1}\{\tilde{T} \geq t\}\Lambda_{i}(\mathrm{d}t \mid W, A), \quad j \in \{1, 2\}.$$

and

$$M(\mathrm{d}t \mid A, W; \Lambda_1, \Lambda_2) = M_1(\mathrm{d}t \mid A, W; \Lambda_1) + M_2(\mathrm{d}t \mid A, W; \Lambda_2).$$

The efficient influence function can now be written as (van der Laan and Robins, 2003; Jewell et al., 2007; Rytgaard and van der Laan, 2022),

$$\psi_{\tau}(O; \Lambda_1, \Lambda_2, \Gamma, \pi) = \psi_{\tau}^1(O; \Lambda_1, \Lambda_2, \Gamma, \pi) - \psi_{\tau}^0(O; \Lambda_1, \Lambda_2, \Gamma, \pi) - \tilde{\Psi}_t^0(\Lambda_1, \Lambda_2, \mu),$$

where

$$\psi_{\tau}^{a}(O; \Lambda_{1}, \Lambda_{2}, \Gamma, \pi)
= \omega_{a}(A, W; \pi) \int_{0}^{\tau} e^{\Gamma(t-|A,W)} M_{1}(\mathrm{d}t \mid A, W; \Lambda_{1})
- \omega_{a}(A, W; \pi) g(\tau, A, W; \Lambda_{1}, \Lambda_{2}) \int_{0}^{\tau} e^{[\Gamma+\Lambda_{1}+\Lambda_{2}](t-|A,W)} M(\mathrm{d}t \mid A, W; \Lambda_{1}, \Lambda_{2})
+ \omega_{a}(A, W; \pi) \int_{0}^{\tau} g(t, A, W; \Lambda_{1}, \Lambda_{2}) e^{[\Gamma+\Lambda_{1}+\Lambda_{2}](t-|A,W)} M(\mathrm{d}t \mid A, W; \Lambda_{1}, \Lambda_{2})
+ g(\tau, a, W; \Lambda_{1}, \Lambda_{2}).$$
(15)

Equations (14) and (15) allow us to construct a one-step estimator by using the definition given in equation (11), which gives the estimator

$$\hat{\Psi}_{t,OS}(\mathcal{D}_n) = \tilde{\Psi}_t^0(\hat{a}_1(\mathcal{D}_n), \hat{a}_2(\mathcal{D}_n), \hat{\mu}_n(\mathcal{D}_n))
+ \mathbb{P}_n[\psi_{\tau}(\cdot; \hat{a}_{1,n}(\mathcal{D}_n), \hat{a}_{2,n}(\mathcal{D}_n), \hat{b}_n(\mathcal{D}_n), \hat{\pi}_n(\mathcal{D}_n))]
= \mathbb{P}_n[\psi_{\tau}^1(\cdot; \hat{a}_{1,n}(\mathcal{D}_n), \hat{a}_{2,n}(\mathcal{D}_n), \hat{b}_n(\mathcal{D}_n), \hat{\pi}_n(\mathcal{D}_n))]
- \mathbb{P}_n[\psi_{\tau}^0(\cdot; \hat{a}_{1,n}(\mathcal{D}_n), \hat{a}_{2,n}(\mathcal{D}_n), \hat{b}_n(\mathcal{D}_n), \hat{\pi}_n(\mathcal{D}_n))].$$
(16)

8. Numerical experiments

In this section we report results from two settings of our simulation studies, both settings are without competing risks. In the next section, we consider a setting with competing risks.

We generate data according to a distribution motivated from a real dataset in which censoring depends on the baseline covariates. We simulate data based on a prostate cancer study described in (Kattan et al., 2000). The outcome of interest was the time to tumor recurrence, and five baseline covariates were used to predict outcome: prostatespecific antigen (PSA, ng/mL), Gleason score sum (GSS, values between 6 and 10), radiation dose (RD), hormone therapy (HT, yes/no) and clinical stage (CS, six values). The study was designed such that a patient's radiation dose depended on when the patient entered the study (Gerds et al., 2013). This in turn implied that the time of censoring depended on the radiation dose. The data were re-analyzed in (Gerds et al., 2013) where a sensitivity analysis was conducted based on simulated data. We use the same simulation setup, where event and censoring times are generated according to parametric Cox-Weibull models estimated from the original data, and the covariates are generated according to either marginal Gaussian normal or binomial distributions estimated from the original data (c.f., Gerds et al., 2013, Section 4.6). We refer to this simulation setting as "dependent censoring". We also considered a simulation setting where data were generated in the same way, except that censoring was generated completely independently. We refer to this simulation setting as "independent censoring".

For all super learners we use a library consisting of three learners: The Kaplan-Meier estimator (Kaplan and Meier, 1958; Gerds, 2019), a Cox model with main effects (Cox, 1972; Therneau, 2022), and a random survival forest (Ishwaran et al., 2008; Ishwaran and Kogalur, 2023). We use the same library to learn the outcome and the censoring mechanism. To be precise, note that given a learner for the cumulative hazard function of the outcome event, we can typically use the same method to construct a learner of the cumulative hazard function of the censoring distribution. This works by training the learner on the data set \mathcal{D}_n^c , where $\mathcal{D}_n^c = \{O_i^c\}_{i=1}^n$ with $O_i^c = (\tilde{T}_i, 1 - \Delta, X_i)$. When we say that we use a learner for the cumulative hazard function of the outcome to learn the cumulative hazard function of the censoring time, we mean that the learner is trained on \mathcal{D}_n^c .

We compare the state learner to two IPCW based super learners: The first super-learner, called IPCW(Cox), uses a Cox model with main effects to estimate the censoring probabilities, while the second super learner, called IPCW(KM), uses the Kaplan-Meier

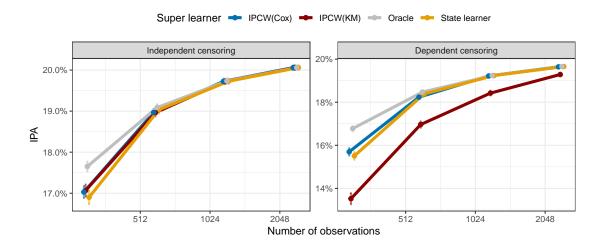


Fig. 2. For the risk prediction models provided by each of the super learners, the IPA is plotted against sample size. The results are averages across 1000 simulated data sets and the error bars are used to quantify the Monte Carlo uncertainty.

estimator to estimate the censoring probabilities. The Cox model for the censoring distribution is correctly specified in both simulation settings while the Kaplan Meier estimator only estimates the censoring model correctly in the simulation setting where censoring is independent. Both IPCW super learners are fitted using the R-package riskRegression (Gerds et al., 2023). Each super learner provides a learner for the cumulative hazard function for the outcome of interest. From the cumulative hazard function a risk prediction model can be obtained (c.f., equation (1) with $\Lambda_2 = 0$). We measure the performance of each super learner by calculating the index of prediction accuracy (IPA) (Kattan and Gerds, 2018) at a fixed time horizon (36 months) for the risk prediction model provided by the super learner. The IPA is 1 minus the ratio between the model's Brier score and the null model's Brier score, where the null model is the model that does not use any covariate information. The IPA is approximated using a large (n = 20,000) independent data set of uncensored data. As a benchmark we calculate the performance of the risk prediction model chosen by the oracle selector, which uses the large data set of uncensored event times to select the learner with the highest IPA.

The results are shown in Figure 2. We see that in the scenario where censoring depends on the covariates, using the Kaplan-Meier estimator to estimate the censoring probabilities provides a risk prediction model with an IPA that is lower than the risk prediction model provided by the state learner. The performance of the risk prediction model selected by the state learner is similar to the risk prediction model selected by the IPCW(Cox) super learner which a priori uses a correctly specified model for the censoring distribution. Both these risk prediction models are close to the performance of the oracle, except for small sample sizes.

We next compare the state learner to the super learner survSL (Westling et al., 2021).

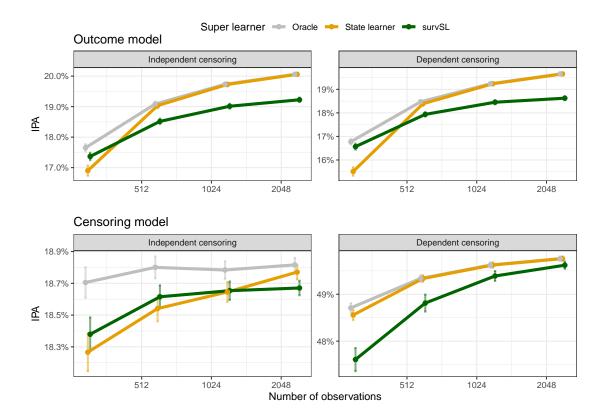


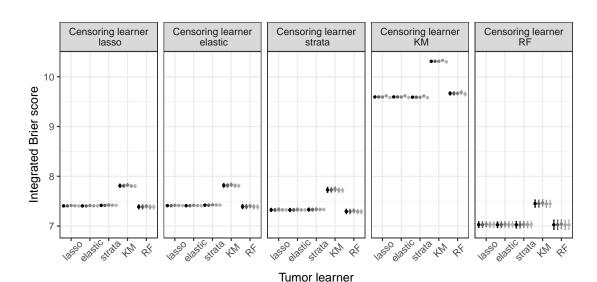
Fig. 3. For the risk prediction models of both the outcome and the censoring probability provided by each of the super learners, the IPA is plotted against sample size. The results are averages across 1000 repetitions and the error bars are used to quantify the Monte Carlo uncertainty.

This is another super learner which like the state learner works without a pre-specified censoring model. Note that both the state learner and survSL provide a prediction model for the event time outcome and also for the probability of being censored. Hence, we compare the performance of these methods with respect to both the outcome and the censoring distribution. Again we use the IPA to quantify the predictive performance.

The results are shown in Figure 3. We see that for most sample sizes, the state learner selected prediction models for both censoring and outcome which have similar or higher IPA compared to the prediction models selected by survSL.

9. Prostate cancer study

The prostate cancer data analyzed by Kattan et al. (2000), which we introduced in Section 8, include a competing event in the form of death without tumor recurrence. For illustration, we fit the state learner to the original data set consisting of 1,042 patients. We consider death without tumor recurrence and recurrence of tumor as competing



Death learner → lasso → elastic → strata → KM → RF

Fig. 4. The results of applying the 125 combinations of learners to the prostate cancer data set. The learners are KM (KM), Cox strata CS (strata), Lasso (lasso), Elastic (elastic), and RF (RF) as described Table 1. The error bars are based on five repetitions using different splits. We refer to learners of Λ_1 , Λ_2 , and Γ as 'Tumor learner', 'Death learner', and 'Censoring learner', respectively.

risks. We include the fives learner KM, Cox strata CS, Lasso, Elastic, and RF which are described in Table 1. We use the same library of learners to learn Λ_1 , Λ_2 , and Γ . In this case, Λ_1 denotes the cause-specific cumulative hazard function of tumor recurrence, and Λ_2 denotes the cause-specific cumulative hazard function of death without tumor recurrence.

[Table 1 about here.]

This gives a library consisting of $5^3 = 125$ learners for the conditional state occupation probability function F defined in equation (5). We use five folds for training and testing the models, and we repeat training and evaluation five times with different splits. The integrated Brier score (defined in Section 5) for all learners are shown in Figure 4, and the top 10 combinations of learners are displayed in Table 2. We see that the prediction performance is mostly affected by the choice of learner for the censoring distribution. Several combinations of learners give similar performance as measured by the integrated Brier score, as long as a random forest is used to model the censoring distribution.

[Table 2 about here.]

As an illustration, we use the learners of the three cumulative hazard functions selected by the state learner and the estimator defined in Section 7.1, equation (16), to

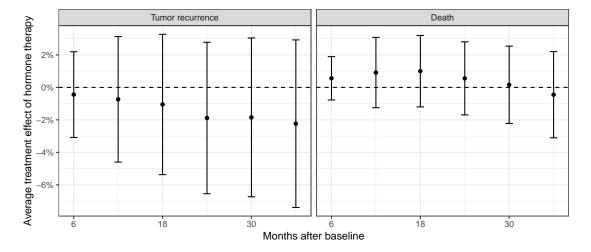


Fig. 5. Estimates of the average treatment effect of hormone therapy on the risk of tumor recurrence and death obtained from the prostate cancer study analyzed by Kattan et al. (2000). The estimates are based on the estimator defined in equation (16) with point wise 95% confidence intervals calculated as described in Section 7. The estimates of the nuisance parameters are provided by the state learner.

estimate the average treatment effect of hormone therapy on risk of tumor recurrence and death. The propensity score is estimated with a lasso model that includes all levels of interaction. The results are shown in Figure 5 for 6 month intervals after baseline with point wise 95% confidence intervals. We see that hormone therapy appears to decrease the risk of tumor recurrence and increase the risk of death without tumor recurrence, but that none of the estimated effects are statistically significant from zero.

10. Discussion

We have proposed a new super learner that can be used with right-censored data and competing events. Compared to existing IPCW-based methods, the advantage of the state learner is that it does not depend on a pre-specified estimator of the censoring distribution, but selects one automatically based on a library of learners for the censoring distribution. Furthermore, the state learner neither requires evaluation of the density of Λ_j nor assumes a (semi-)parametric form for Λ_j . In the remainder of this section we point to some limitations of our proposal and discuss avenues for further research.

A major advantage of the state learner is that the performance of each combination of learners can be estimated without (semi)parametric assumptions or additional nuisance parameters. The drawback with this approach is that we are rarely interested in features of the observed data distribution when the data are right-censored: we are only interested in the uncensored event time distribution. The finite sample oracle inequality in Corollary 1 concerns the function F, which is a feature of $P \in \mathcal{P}$, while what we are typically interested in is Λ_i or S, which are features of $Q \in \mathcal{Q}$. We emphasise that

while the state learner provides us with estimates of Λ_j and Γ based on libraries \mathcal{A}_j and \mathcal{B} , the performance of these learners is not assessed directly for their respective target parameters, but only indirectly via the performance of F. For settings without competing risks, our numerical studies suggest that measuring the performance of F also leads to good performance for estimation of S. However, further research on this issue, both numerical and theoretical, is warranted.

Our proposed super learner can be implemented with a broad library of learners and using existing software. Furthermore, while the library $\mathcal{F}(\mathcal{A}_1, \mathcal{A}_2, \mathcal{B})$ consists of $|\mathcal{A}_1||\mathcal{A}_2||\mathcal{B}|$ many learners, we only need to fit $|\mathcal{A}_1| + |\mathcal{A}_2| + |\mathcal{B}|$ many learners in each fold. To evaluate the performance of each learner we need to perform $|\mathcal{A}_1||\mathcal{A}_2||\mathcal{B}|$ many operations to calculate the integrated Brier score in each hold-out sample, one for each combination of the fitted models, but these operations are often negligible compared to fitting the models. Hence the state learner is essentially not more computationally demanding than any procedure that uses super learning to learn Λ_1 , Λ_2 , and Γ separately. While our proposal is based on constructing the library \mathcal{F} from libraries for learning Λ_1 , Λ_2 , and Γ , it would also be of interest to consider learners that estimate F directly.

In our numerical studies, we only considered learners of Λ_j and Γ that provide cumulative hazard functions which are piece-wise constant in the time argument. This simplifies the calculation of F as the integrals in equation (6) reduce to sums. When Λ_j or Γ are absolutely continuous in the time argument, calculating F is more involved, but we expect that a good approximation can be achieved by discretisation. In the future, we intend to investigate the performance of the state learner when using a broader library of learners in more comprehensive simulation studies.

A. Theoretical guarantees for the state learner

In this section we provide proofs of the results stated in Section 6.

Define
$$\bar{B}_{\tau,0}(F,o) = \bar{B}_{\tau}(F,o) - \bar{B}_{\tau}(F_0,o)$$
 and $R_0(F) = P_0[\bar{B}_{\tau,0}(F,\cdot)].$

LEMMA 1.
$$R_0(F) = ||F - F_0||_{P_0}^2$$
, where $||\cdot||_{P_0}$ is defined in equation (7).

PROOF. For any $t \in [0, \tau]$ and $j \in \{-1, 0, 1, 2\}$ we have

$$\begin{split} &\mathbb{E}_{P_0}\left[(F(t,j,X) - \mathbb{1}\{\eta(t) = j\})^2\right] \\ &= \mathbb{E}_{P_0}\left[(F(t,j,X) - F_0(t,j,X) + F_0(t,j,X) - \mathbb{1}\{\eta(t) = j\})^2\right] \\ &= \mathbb{E}_{P_0}\left[(F(t,j,X) - F_0(t,j,X))^2\right] + \mathbb{E}_{P_0}\left[(F_0(t,j,X) - \mathbb{1}\{\eta(t) = j\})^2\right] \\ &+ 2\,\mathbb{E}_{P_0}\left[(F(t,j,X) - F_0(t,j,X))(F_0(t,j,X) - \mathbb{1}\{\eta(t) = j\})\right] \\ &= \mathbb{E}_{P_0}\left[(F(t,j,X) - F_0(t,j,X))^2\right] + \mathbb{E}_{P_0}\left[(F_0(t,j,X) - \mathbb{1}\{\eta(t) = j\})^2\right], \end{split}$$

where the last equality follows from the tower property. Hence, using Fubini, we have

$$P[\bar{B}_{\tau}(F,\cdot)] = ||F - F_0||_{P_0}^2 + P_0[\bar{B}_{\tau}(F_0,\cdot)].$$

PROOF (PROOF OF PROPOSITION 1). The result follows from Lemma 1.

In the following, let Θ denote the function space consisting of all conditional state occupation probability functions for some measure P as defined in equation (5).

PROOF (PROOF OF COROLLARY 1). First note that minimising the loss \bar{B}_{τ} is equivalent to minimising the loss $\bar{B}_{\tau,0}$, so the discrete super learner and oracle according to \bar{B}_{τ} and $\bar{B}_{\tau,0}$ are identical. By Lemma 1, $R_0(F) \geq 0$ for any $F \in \Theta$, and so using Theorem 2.3 from (van der Vaart et al., 2006) with p = 1, we have that for all $\delta > 0$,

$$\mathbb{E}_{P_0} \left[R_0(\hat{\varphi}_n(\mathcal{D}_n^{-k})) \right] \leq (1 + 2\delta) \, \mathbb{E}_{P_0} \left[R_0(\tilde{\varphi}_n(\mathcal{D}_n^{-k})) \right]$$

$$+ (1 + \delta) \frac{16K}{n} \log(1 + |\mathcal{F}_n|) \sup_{F \in \Theta} \left\{ M(F) + \frac{v(F)}{R_0(F)} \left(\frac{1}{\delta} + 1 \right) \right\}$$

where for each $F \in \Theta$, (M(F), v(F)) is some Bernstein pair for the function $o \mapsto \bar{B}_{\tau,0}(F,o)$. As $\bar{B}_{\tau,0}(F,\cdot)$ is uniformly bounded by τ for any $F \in \Theta$, it follows from section 8.1 in (van der Vaart et al., 2006) that $(\tau, 1.5P_0[\bar{B}_{\tau,0}(F,\cdot)^2])$ is a Bernstein pair for $\bar{B}_{\tau,0}(F,\cdot)$. Now, for any $a,b,c \in \mathbb{R}$ we have

$$(a-c)^{2} - (b-c)^{2} = (a-b+b-c)^{2} - (b-c)^{2}$$

$$= (a-b)^{2} + (b-c)^{2} + 2(b-c)(a-b) - (b-c)^{2}$$

$$= (a-b)\{(a-b) + 2(b-c)\}$$

$$= (a-b)\{a+b-2c\},$$

so using this with a = F(t, j, x), $b = F_0(t, j, x)$, and $c = \mathbb{1}\{\eta(t) = j\}$, we have by Jensen's inequality

$$P_{0}[\bar{B}_{\tau,0}(F,\cdot)^{2}]$$

$$\leq 2\tau \mathbb{E}_{P_{0}} \left[\sum_{j=-1}^{2} \int_{0}^{\tau} \left\{ (F(t,j,X) - \mathbb{1}\{\eta(t) = j\})^{2} - (F_{0}(t,j,X) - \mathbb{1}\{\eta(t) = j\})^{2} \right\}^{2} dt \right]$$

$$= 2\tau \mathbb{E}_{P_{0}} \left[\sum_{j=-1}^{2} \int_{0}^{\tau} (F(t,j,X) - F_{0}(t,j,X))^{2} \times \left\{ F(t,j,X) + F_{0}(t,j,X) - 2\mathbb{1}\{\eta(t) = j\} \right\}^{2} dt \right]$$

$$\leq 8\tau \mathbb{E}_{P_{0}} \left[\sum_{j=-1}^{2} \int_{0}^{\tau} (F(t,j,X) - F_{0}(t,j,X))^{2} dt \right].$$

$$= 8\tau \|F - F_{0}\|_{P_{0}}^{2}.$$

Thus when $v(F) = 1.5P_0[\bar{B}_{\tau,0}(F,\cdot)^2]$ we have by Lemma 1

$$\frac{v(F)}{R_0(F)} = 1.5 \frac{P_0[\bar{B}_{\tau,0}(F,\cdot)^2]}{P_0[\bar{B}_{\tau,0}(F,\cdot)]} \le 12\tau,$$

and so using the Bernstein pairs $(\tau, 1.5P_0[\bar{B}_{\tau,0}(F,\cdot)^2])$ we have

$$\sup_{F\in\Theta}\left\{M(F)+\frac{v(F)}{R_0(F)}\left(\frac{1}{\delta}+1\right)\right\}\leq \tau\left(13+\frac{12}{\delta}\right),$$

For all $\delta > 0$ we thus have

$$\mathbb{E}_{P_0} \left[R_0(\hat{\varphi}_n(\mathcal{D}_n^{-k})) \right] \leq (1 + 2\delta) \, \mathbb{E}_{P_0} \left[R_0(\tilde{\varphi}_n(\mathcal{D}_n^{-k})) \right]$$
$$+ (1 + \delta) \log(1 + |\mathcal{F}_n|) \tau \frac{16K}{n} \left(13 + \frac{12}{\delta} \right),$$

and then the final result follows from Lemma 1.

PROOF (PROOF OF COROLLARY 2). By definition of the oracle and Lemma 1,

$$\mathbb{E}_{P_0} \left[\| \tilde{\varphi}_n(\mathcal{D}_n^{-k}) - F_0 \|_{P_0}^2 \right] \le \mathbb{E}_{P_0} \left[\| \varphi_n(\mathcal{D}_n^{-k}) - F_0 \|_{P_0}^2 \right]$$

for all $n \in \mathbb{N}$. The results then follows from Corollary 1.

B. The state learner with targeted learning

In this section show that a product structure is preserved when an estimator $\bar{\Psi}(\hat{F}_n, \hat{H}_n)$ is used instead of $\tilde{\Psi}(\hat{\Lambda}_n, \hat{\Gamma}_n, \hat{H}_n)$.

PROOF (PROOF OF PROPOSITION 2). For notational convenience we suppress X in the following. The final result can be obtained by adding the argument X to all functions and averaging. We use the relations from equation (8) to write

$$\begin{split} &\int_{0}^{\tau} w(s) \left\{ \Gamma(s) - \hat{\Gamma}_{n}(s) \right\} [\Lambda - \hat{\Lambda}_{n}] (\mathrm{d}s) \\ &= \int_{0}^{\tau} w(s) \left\{ \int_{0}^{s} \frac{F(\mathrm{d}u, 2)}{F(u - 0)} - \int_{0}^{s} \frac{\hat{F}_{n}(\mathrm{d}u, 2)}{\hat{F}_{n}(u - 0)} - \right\} \left[\frac{F(\mathrm{d}s, 1)}{F(s - 0)} - \frac{\hat{F}_{n}(\mathrm{d}s, 1)}{\hat{F}_{n}(s - 0)} \right] \\ &= \int_{0}^{\tau} w(s) \left\{ \int_{0}^{s} \left(\frac{1}{F(u - 0)} - \frac{1}{\hat{F}_{n}(u - 0)} \right) F(\mathrm{d}u, 2) \right. \\ &\quad + \int_{0}^{s} \frac{1}{\hat{F}_{n}(u - 0)} \left[F(\mathrm{d}u, 2) - \hat{F}_{n}(\mathrm{d}u, 2) \right] \right\} \\ &\quad \times \left[\left(\frac{1}{F(s - 0)} - \frac{1}{\hat{F}_{n}(s - 0)} \right) F(\mathrm{d}s, 1) + \frac{1}{\hat{F}_{n}(s - 0)} \left(F(\mathrm{d}s, 1) - \hat{F}_{n}(\mathrm{d}s, 1) \right) \right] \\ &= \int_{0}^{\tau} \int_{0}^{s} w(s) \left(\frac{1}{F(u - 0)} - \frac{1}{\hat{F}_{n}(u - 0)} \right) \left(\frac{1}{F(s - 0)} - \frac{1}{\hat{F}_{n}(s - 0)} \right) F(\mathrm{d}u, 2) F(\mathrm{d}s, 1) \\ &\quad + \int_{0}^{\tau} \int_{0}^{s} w(s) \left(\frac{1}{F(u - 0)} - \frac{1}{\hat{F}_{n}(u - 0)} \right) \frac{F(\mathrm{d}u, 2)}{\hat{F}_{n}(u - 0)} \left(F(\mathrm{d}s, 1) - \hat{F}_{n}(\mathrm{d}s, 1) \right) \\ &\quad + \int_{0}^{\tau} \int_{0}^{s} \frac{w(s)}{\hat{F}_{n}(u - 0)} \left[F(\mathrm{d}u, 2) - \hat{F}_{n}(\mathrm{d}u, 2) \right] \left(\frac{1}{F(s - 0)} - \frac{1}{\hat{F}_{n}(s - 0)} \right) F(\mathrm{d}s, 1) \\ &\quad + \int_{0}^{\tau} \int_{0}^{s} \frac{w(s)}{\hat{F}_{n}(u - 0)} \left[F(\mathrm{d}u, 2) - \hat{F}_{n}(\mathrm{d}u, 2) \right] \frac{1}{\hat{F}_{n}(s - 0)} \left(F(\mathrm{d}s, 1) - \hat{F}_{n}(\mathrm{d}s, 1) \right). \end{split}$$

Consider the first term on the right hand side. Defining

$$w_n^*(t) = \left(F(t-,0) - \hat{F}_n(t-,0) \right) \left(\frac{1}{F(t-,0)} - \frac{1}{\hat{F}_n(t-,0)} \right),$$

we can write

$$\begin{split} & \int_0^\tau \int_0^s w(s) \left(\frac{1}{F(u-,0)} - \frac{1}{\hat{F}_n(u-,0)} \right) \left(\frac{1}{F(s-,0)} - \frac{1}{\hat{F}_n(s-,0)} \right) F(\mathrm{d}u,2) F(\mathrm{d}s,1) \\ & = \int_0^\tau \int_0^s w(s) w_n^*(u) \left(F(u-,0) - \hat{F}_n(u-,0) \right) \\ & \qquad \times w_n^*(s) \left(F(s-,0) - \hat{F}_n(s-,0) \right) F(\mathrm{d}u,2) F(\mathrm{d}s,1) \\ & = \int_0^\tau \int_0^s w_n^a(s,u) \left(F(u-,0) - \hat{F}_n(u-,0) \right) \left(F(s-,0) - \hat{F}_n(s-,0) \right) F(\mathrm{d}u,2) F(\mathrm{d}s,1), \end{split}$$

where we have defined $w_n^a(s, u) = w(s)w_n^*(s)w_n^*(u)$. By assumption, $w_n^a(s, u)$ is uniformly bounded. The same approach can be applied to the three remaining terms which gives the result.

References

- Andersen, P. K., Borgan, O., Gill, R. D. and Keiding, N. (2012) Statistical models based on counting processes. Springer Science & Business Media.
- Andersen, P. K., Klein, J. P. and Rosthøj, S. (2003) Generalised linear models for correlated pseudo-observations, with applications to multi-state models. *Biometrika*, **90**, 15–27.
- Begun, J. M., Hall, W. J., Huang, W.-M. and Wellner, J. A. (1983) Information and asymptotic efficiency in parametric-nonparametric models. *The Annals of Statistics*, **11**, 432–452.
- Benichou, J. and Gail, M. H. (1990) Estimates of absolute cause-specific risk in cohort studies. *Biometrics*, 813–826.
- Bickel, P. J., Klaassen, C. A., Ritov, Y. and Wellner, J. A. (1993) Efficient and adaptive estimation for semiparametric models, vol. 4. Johns Hopkins University Press Baltimore.
- Breiman, L. (1996) Stacked regressions. Machine learning, 24, 49–64.
- Brier, G. W. et al. (1950) Verification of forecasts expressed in terms of probability. *Monthly weather review*, **78**, 1–3.
- Chernozhukov, V., Chetverikov, D., Demirer, M., Duflo, E., Hansen, C., Newey, W. and Robins, J. (2018) Double/debiased machine learning for treatment and structural parameters.

- Cox, D. R. (1972) Regression models and life-tables. Journal of the Royal Statistical Society: Series B (Methodological), 34, 187–202.
- Fan, J. and Gijbels, I. (1996) Local polynomial modelling and its applications. Routledge.
- Fisher, A. and Kennedy, E. H. (2021) Visually communicating and teaching intuition for influence functions. *The American Statistician*, **75**, 162–172.
- Friedman, J., Hastie, T. and Tibshirani, R. (2010) Regularization paths for generalized linear models via coordinate descent. *Journal of Statistical Software*, **33**, 1–22. URL: https://www.jstatsoft.org/v33/i01/.
- Gensheimer, M. F. and Narasimhan, B. (2019) A scalable discrete-time survival model for neural networks. *PeerJ*, **7**, e6257.
- Gerds, T. A. (2019) prodlim: Product-Limit Estimation for Censored Event History Analysis. URL: https://CRAN.R-project.org/package=prodlim. R package version 2019.11.13.
- Gerds, T. A. and Kattan, M. W. (2021) Medical risk prediction models: with ties to machine learning. CRC Press.
- Gerds, T. A., Kattan, M. W., Schumacher, M. and Yu, C. (2013) Estimating a time-dependent concordance index for survival prediction models with covariate dependent censoring. *Statistics in medicine*, **32**, 2173–2184.
- Gerds, T. A., Ohlendorff, J. S. and Ozenne, B. (2023) riskRegression: Risk Regression Models and Prediction Scores for Survival Analysis with Competing Risks. URL: https://CRAN.R-project.org/package=riskRegression. R package version 2023.03.22.
- Gerds, T. A. and Schumacher, M. (2006) Consistent estimation of the expected Brier score in general survival models with right-censored event times. *Biometrical Journal*, 48, 1029–1040.
- Gill, R. D., van der Laan, M. J. and Robins, J. M. (1997) Coarsening at random: Characterizations, conjectures, counter-examples. In *Proceedings of the First Seattle Symposium in Biostatistics*, 255–294. Springer.
- Gneiting, T. and Raftery, A. E. (2007) Strictly proper scoring rules, prediction, and estimation. *Journal of the American statistical Association*, **102**, 359–378.
- Golmakani, M. K. and Polley, E. C. (2020) Super learner for survival data prediction. *The International Journal of Biostatistics*, **16**, 20190065.
- Gonzalez Ginestet, P., Kotalik, A., Vock, D. M., Wolfson, J. and Gabriel, E. E. (2021) Stacked inverse probability of censoring weighted bagging: A case study in the inf-carehiv register. *Journal of the Royal Statistical Society Series C: Applied Statistics*, 70, 51–65.

- Graf, E., Schmoor, C., Sauerbrei, W. and Schumacher, M. (1999) Assessment and comparison of prognostic classification schemes for survival data. *Statistics in medicine*, **18**, 2529–2545.
- Han, X., Goldstein, M., Puli, A., Wies, T., Perotte, A. and Ranganath, R. (2021) Inverse-weighted survival games. *Advances in Neural Information Processing Systems*, **34**.
- Hernán, M. and Robins, J. (2020) Causal Inference: What If. Boca Raton: Chapman & Hall/CRC.
- Hines, O., Dukes, O., Diaz-Ordaz, K. and Vansteelandt, S. (2022) Demystifying statistical learning based on efficient influence functions. *The American Statistician*, **76**, 292–304.
- Hjort, N. L. (1992) On inference in parametric survival data models. *International Statistical Review/Revue Internationale de Statistique*, 355–387.
- Hothorn, T., Bühlmann, P., Dudoit, S., Molinaro, A. and van der Laan, M. J. (2006) Survival ensembles. *Biostatistics*, **7**, 355–373.
- Η. (2023)Ishwaran, and Kogalur, U. FastUnified RandomForests(RF-SRC).forSurvival, Regression, andClassificationURL: https://cran.r-project.org/package=randomForestSRC. R package version 3.2.2.
- Ishwaran, H., Kogalur, U. B., Blackstone, E. H. and Lauer, M. S. (2008) Random survival forests. *The annals of applied statistics*, **2**, 841–860.
- Jewell, N. P., Lei, X., Ghani, A. C., Donnelly, C. A., Leung, G. M., Ho, L.-M., Cowling, B. J. and Hedley, A. J. (2007) Non-parametric estimation of the case fatality ratio with competing risks data: an application to severe acute respiratory syndrome (sars). Statistics in medicine, 26, 1982–1998.
- Kaplan, E. L. and Meier, P. (1958) Nonparametric estimation from incomplete observations. *Journal of the American statistical association*, **53**, 457–481.
- Kattan, M. W. and Gerds, T. A. (2018) The index of prediction accuracy: an intuitive measure useful for evaluating risk prediction models. *Diagnostic and prognostic research*, **2**, 1–7.
- Kattan, M. W., Zelefsky, M. J., Kupelian, P. A., Scardino, P. T., Fuks, Z. and Leibel, S. A. (2000) Pretreatment nomogram for predicting the outcome of three-dimensional conformal radiotherapy in prostate cancer. *Journal of clinical oncology*, 18, 3352–3359.
- Katzman, J. L., Shaham, U., Cloninger, A., Bates, J., Jiang, T. and Kluger, Y. (2018) Deepsurv: personalized treatment recommender system using a Cox proportional hazards deep neural network. BMC medical research methodology, 18, 1–12.
- Keles, S., van der Laan, M. and Dudoit, S. (2004) Asymptotically optimal model selection method with right censored outcomes. *Bernoulli*, **10**, 1011–1037.

- Kennedy, E. H. (2016) Semiparametric theory and empirical processes in causal inference. In Statistical causal inferences and their applications in public health research, 141–167. Springer.
- (2022) Semiparametric doubly robust targeted double machine learning: a review. arXiv preprint arXiv:2203.06469.
- Kvamme, H. and Borgan, Ø. (2021) Continuous and discrete-time survival prediction with neural networks. Lifetime Data Analysis, 27, 710–736.
- van der Laan, M. J. and Dudoit, S. (2003) Unified cross-validation methodology for selection among estimators and a general cross-validated adaptive epsilon-net estimator: Finite sample oracle inequalities and examples. Tech. rep., Division of Biostatistics, University of California.
- van der Laan, M. J., Polley, E. C. and Hubbard, A. E. (2007) Super learner. Statistical applications in genetics and molecular biology, 6.
- van der Laan, M. J. and Robins, J. M. (2003) Unified methods for censored longitudinal data and causality. Springer Science & Business Media.
- van der Laan, M. J. and Rose, S. (2011) Targeted learning: causal inference for observational and experimental data. Springer Science & Business Media.
- Lee, C., Zame, W., Yoon, J. and van der Schaar, M. (2018) Deephit: A deep learning approach to survival analysis with competing risks. In Proceedings of the AAAI conference on artificial intelligence, vol. 32.
- Lee, D. K., Chen, N. and Ishwaran, H. (2021) Boosted nonparametric hazards with time-dependent covariates. Annals of Statistics, 49, 2101.
- Li, Y., Xu, K. S. and Reddy, C. K. (2016) Regularized parametric regression for highdimensional survival analysis. In Proceedings of the 2016 SIAM International Conference on Data Mining, 765–773. SIAM.
- Mogensen, U. B. and Gerds, T. A. (2013) A random forest approach for competing risks based on pseudo-values. Statistics in medicine, 32, 3102–3114.
- Molinaro, A. M., Dudoit, S. and van der Laan, M. J. (2004) Tree-based multivariate regression and density estimation with right-censored data. Journal of Multivariate Analysis, 90, 154–177.
- Ozenne, B., Sørensen, A. L., Scheike, T., Torp-Pedersen, C. and Gerds, T. A. (2017) riskregression: Predicting the risk of an event using Cox regression models. R Journal, **9**, 440–460.
- Pfanzagl, J. and Wefelmeyer, W. (1982) Contributions to a general asymptotic statistical theory. Springer.
- Polley, E., LeDell, E., Kennedy, C. and van der Laan, M. (2023) SuperLearner: Super Learner Prediction. URL: https://CRAN.R-project.org/package=SuperLearner. R package version 2.0-28.1.

- Polley, E. C. and van der Laan, M. J. (2011) Super learning for right-censored data. In *Targeted Learning: Causal Inference for Observational and Experimental Data* (eds. M. J. van der Laan and S. Rose), 249–258. Springer.
- Redelmeier, D. A., Koehler, D. J., Liberman, V. and Tversky, A. (1995) Probability judgment in medicine: Discounting unspecified possibilities. *Medical Decision Making*, 15, 227–230.
- Rytgaard, H. C., Eriksson, F. and van der Laan, M. J. (2021) Estimation of time-specific intervention effects on continuously distributed time-to-event outcomes by targeted maximum likelihood estimation. *Biometrics*.
- Rytgaard, H. C., Gerds, T. A. and van der Laan, M. J. (2022) Continuous-time targeted minimum loss-based estimation of intervention-specific mean outcomes. *The Annals of Statistics*, **50**, 2469–2491.
- Rytgaard, H. C. and van der Laan, M. J. (2022) Targeted maximum likelihood estimation for causal inference in survival and competing risks analysis. *Lifetime Data Analysis*, 1–30.
- Sachs, M. C., Discacciati, A., Everhov, Å. H., Olén, O. and Gabriel, E. E. (2019) Ensemble prediction of time-to-event outcomes with competing risks: A case-study of surgical complications in Crohn's disease. *Journal of the Royal Statistical Society* Series C: Applied Statistics, 68, 1431–1446.
- Simon, N., Friedman, J., Hastie, T. and Tibshirani, R. (2011) Regularization paths for Cox's proportional hazards model via coordinate descent. *Journal of Statistical Software*, **39**, 1–13. URL: https://www.jstatsoft.org/v39/i05/.
- Steingrimsson, J. A., Diao, L. and Strawderman, R. L. (2019) Censoring unbiased regression trees and ensembles. *Journal of the American Statistical Association*, **114**, 370–383.
- Therneau, T. M. (2022) A Package for Survival Analysis in R. URL: https://CRAN.R-project.org/package=survival. R package version 3.4-0.
- Tsiatis, A. (2007) Semiparametric theory and missing data. Springer Science & Business Media.
- van der Vaart, A. W. (2000) Asymptotic statistics, vol. 3. Cambridge university press.
- van der Vaart, A. W., Dudoit, S. and van der Laan, M. J. (2006) Oracle inequalities for multi-fold cross validation. *Statistics & Decisions*, **24**, 351–371.
- Verweij, P. J. and van Houwelingen, H. C. (1993) Cross-validation in survival analysis. *Statistics in medicine*, **12**, 2305–2314.
- Westling, T., Luedtke, A., Gilbert, P. and Carone, M. (2021) Inference for treatment-specific survival curves using machine learning. arXiv preprint arXiv:2106.06602.

26 Munch et al.

- Whitney, D., Shojaie, A. and Carone, M. (2019) Comment: Models as (deliberate) approximations. Statistical science: a review journal of the Institute of Mathematical Statistics, 34, 591.
- Wolpert, D. H. (1992) Stacked generalization. Neural networks, 5, 241–259.
- Yao, J., Zhu, X., Zhu, F. and Huang, J. (2017) Deep correlational learning for survival prediction from multi-modality data. In *International conference on medical image computing and computer-assisted intervention*, 406–414. Springer.

Tables

Table 1. Overview of the five learners used by the state learner when applied to the prostate cancer data set. The Meier estimator was fitted using the package prodlim (Gerds, 2019). All Cox models included all five covariates in the and were fitted using the package survival (Therneau, 2022). All penalised Cox models included all five covariates as predictors and were fitted using the package glmnet (Simon et al., 2011; Friedman et al., 2010). The random forest was with the package randomForestSRC (Ishwaran and Kogalur, 2023).

Family	Model	Description	
Marginal	KM	The Kaplan-Meier estimator	
Cox	Cox	All five covariates included with additive effects	
	Cox strata CS	Cox model stratified on CS	
Penalised Cox	Lasso	Cox model with L_1 -norm penalty	
	Elastic	Cox model with L_1 - and L_2 -norm penalty	
Random forest	RF	Random forest with default settings	

Table 2. The 10 best performing models in terms of integrated Brier score. The reported standard errors a on five repetitions using different splits. The models are described in Table 1. We refer to learners of Λ_1 , Λ_2 , 'Tumor learner', 'Death learner', and 'Censoring learner', respectively.

Tumor learner	Death learner	Censoring learner	Integrated Brier score
RF	KM	RF	7.02 ± 0.05
Cox strata CT	Elastic	RF	7.03 ± 0.03
RF	Elastic	RF	7.03 ± 0.05
RF	RF	RF	7.03 ± 0.05
Cox strata CT	Lasso	RF	7.03 ± 0.03
RF	Lasso	RF	7.03 ± 0.05
Elastic	Elastic	RF	7.03 ± 0.03
Lasso	Elastic	RF	7.03 ± 0.03
Elastic	Lasso	RF	7.03 ± 0.03
Lasso	Lasso	RF	7.03 ± 0.03