Table 1: Descriptions of variables.

Variable	Description
R	Time until start of registry
W	Baseline covariates
S	Susceptibility to effects of metalworking fluid exposure
$H(t)$	Adverse health status at time t
$N(t)$	Employment status at time t
$A(t)$	Metalworking fluid exposure at time t
$D(t)$	Mortality status at time t
$Y^*(t)$	Cancer status at time t
$Y(t)$	Observed Cancer status at time t
$t = \{1, 2, \ldots, 20\}$	Time, indexed in years after hire

Methods

Causal model

The UAW-GM Cohort data included person-year level exposure, outcome, and covariate data starting at hire. To emulate the shape of the data for this longitudinal cohort, we considered 20 years of data over time indexed by years since hire. Notation representing the variables of interest are presented in Table 1. The causal model represents hypothetical relationships between variables over time compatible with the theory underlying the HWSE in longitudinal occupational cohort studies. At each time point, the effect of cumulative exposure $\bar{A}(t)$ on cancer incidence $Y^*(t)$ is confounded by the path through employment status $N(t)$ and underlying health $H(t)$ as well as the path through past exposure $\bar{A}(t-1)$ and vital status $D(t)$. These paths follow straightforward logic: occupational exposure depends upon employment status and past exposure; mortality status is affected by past exposure and cancer history. Confounding by baseline covariates W is assumed throughout.

Assume we have $n = 50\,000$ iid units in X with

$$X_i(t) = \left(R_i = 0, W_i, S_i, \bar{H}_i(t), \bar{N}_i(t), \bar{A}_i(t), \bar{a}_i(t), \bar{Y}_i^*(t) = \bar{Y}_i(t)\right).$$

In general, we use bar notation to indicate variable history as follows $\bar{X}_i(t) = (X_i(k))_{k=1}^t$. Note that true cancer status $Y^*(t)$ is not observed until $t \geq R$, after the start of the registry. Call X the full data, where we have $R = 0$ for all. In the observed data X^{obs}, we cannot assume $R = 0$ for all. Additionally, susceptibility S and underlying health status H are not known:

$$X_i^{\text{obs}}(t) = \left(R_i, W_i, \bar{N}_i(t), \bar{A}_i(t), \bar{a}_i(t), \bar{Y}_i(t)\right).$$

Under the causal model, we assume the following non-parametric structural equations:

$$
\begin{aligned}
R &= f_R\left(U_R\right) \\
W &= f_W\left(U_W\right) \\
S &= f_S\left(U_S\right) \\
H(t) &= f_{H(t)}\left(H(t-1), U_{H(t)}\right) \\
N(t) &= f_{N(t)}\left(W, N(t-1), H(t), A(t-1), U_{N(t)}\right) \\
A(t) &= f_{A(t)}\left(W, \bar{A}(t-1), N(t), U_{A(t)}\right) \\
D(t) &= f_{D(t)}\left(W, \bar{A}(t-1), D(t-1), Y^*(t-1), U_{D(t)}\right) \\
Y^*(t) &= f_{Y^*(t)}\left(W, S, H(t), \bar{A}(t), D(t), Y^*(t-1), U_{Y^*(t)}\right) \\
Y(t) &= Y^*(t) \times \mathbb{1}\left[Y^*(\lfloor R \rfloor) = 0\right] \times \mathbb{1}\left[D(t) = 0\right] \ .
\end{aligned}
$$

The exogenous variables (errors) $U = \left(U_R, U_W, U_S, U_{H(t)}, U_{N(t)}, U_{A(t)}, U_{D(t)}, U_{Y^*(t)}\right)_{t=1}^{T}$ are mutually independent. Exposure status is a time-varying indicator, and exposure history is summarized as $\bar{A}(t) = \mathbb{1}\left[\sum_{k=1}^{t} \mathbb{1}\left[A(k) = 1\right] > 0\right]$. The outcome of interest is a survival outcome, so $Y^*(t-1) = 1 \Rightarrow Y^*(t) = 1$. The observed outcome $Y(t)$ at time t is a function of true outcome status, time of left censoring, and time of right censoring. An abbreviated directed acyclic graph (DAG) representing the causal relationships encoded in the equations above is presented in Figure 1.

Figure 1: Directed acyclic graph representing the causal relationships encoded in the non-parametric structural equation model at time t.

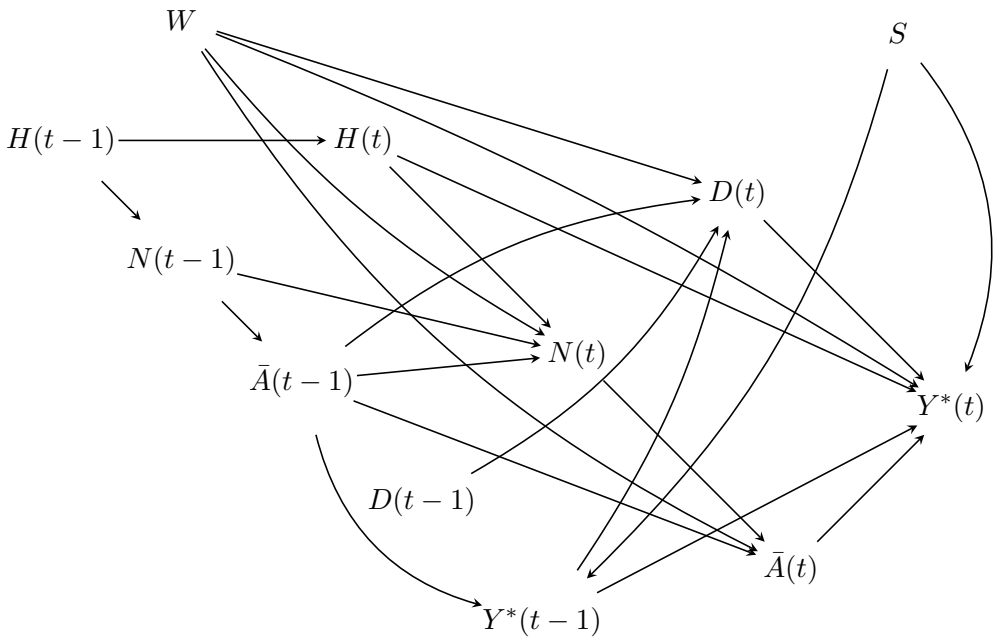


Figure 1 clarifies our conceptualization of the HWSE. At each time point t, the cumulative effect of exposure $\bar{A}(t)$ on cancer incidence $Y^*(t)$ is confounded by employment status $N(t)$ through the

backdoor path $\bar{A}(t) \leftarrow N(t) \leftarrow H(t) \rightarrow Y^*(t)$. In the absence of observed data on health status $H(t)$, an analyst may be tempted to condition on employment status $N(t)$, but doing so would introduce collider bias while blocking the causal path between past exposure $\bar{A}(t-1)$ and cancer $Y^*(t)$. Furthermore, an analysis starting at an arbitrary time point after the time origin (hire) would be tantamount to conditioning on those still alive at that time, which would also result in both collider bias and the conditioning on nodes on the causal path between the exposure and outcome of interest.

Simulation

To generate data compatible with our structural causal model, we imposed parametric relationships between the variables. For the $n = 50\,000$ units over $T = 20$ years, we have:

- $U_j \overset{\text{iid}}{\sim} \text{uniform}\,[0,1]$ for all j
- In full data $R = 0$ otherwise $R \sim \text{uniform}\,[0,30]$
- $W = \mathbb{1}\,[U_W \leq p_W] \sim \text{Bernoulli}\,(p_W)$
- $S = \mathbb{1}\,[U_S \leq p_S] \sim \text{Bernoulli}\,(p_S)$
- If $H(t-1) = 1$, then $H(t) = 1$ otherwise $H(t) = \mathbb{1}\left[U_{H(t)} \leq p_H\right] \sim \text{Bernoulli}\,(p_H)$
- if $N(t-1) = 0$ then $N(t) = 0$ otherwise

$$N(t) \sim \text{Bernoulli}\left\{\text{logit}\left(\beta_0^N + \beta_W^N W + \beta_H^N H(t) + \beta_A^N A(t-1) \times \mathbb{1}\,[t>1] + U_{N(t)}\right)\right\}$$

- If $N(t) = 0$ then $A(t) = 0$ otherwise

$$A(t) \sim \text{Bernoulli}\left\{\text{logit}\left(\left(\beta_0^A + \beta_W^A W\right) \times \mathbb{1}\,[t=1] + \beta_A^A A(t-1) \times \mathbb{1}\,[t>1] + U_{A(t)}\right)\right\}$$

- If $D(t-1) = 1$ then $D(t) = 1$ otherwise

$$D(t) \sim \text{Bernoulli}\left\{\text{logit}\left(\begin{array}{r}\beta_0^D + \beta_W^D W + \beta_{\bar{A}}^D \bar{A}(t-1) \times \mathbb{1}\,[t>1] \\ +\beta_{\bar{Y}}^D \sum_{k=1}^{t-1} Y^*(k) \times \mathbb{1}\,[t>1] + U_{D(t)}\end{array}\right)\right\}$$

- If $Y^*(t-1) = 1$ then $Y^*(t) = 1$ otherwise

$$Y^*(t) \sim \text{Bernoulli}\left\{\text{logit}\left(\begin{array}{r}\beta_0^Y + \beta_W^Y W + \beta_A^Y A(t) + \beta_{\bar{A}}^Y \bar{A}(t-1) \times \mathbb{1}\,[t>1] \\ +\beta_S^Y S \times \bar{A}(t) + \beta_H^Y H(t) + U_{Y^*(t)}\end{array}\right)\right\}$$

- If $t < R$ then $Y(t) = 0$
- If $t \geq R$ then $Y(t) = Y^*(t) \times \mathbb{1}\,[Y^*(\lfloor R \rfloor) = 0] \times \mathbb{1}\,[D(t) = 0]$.

Five sets of data were generated using these equations to represent five scenarios. Scenario 1 represents the base case where 10% of workers are susceptible to exposure-related effects, the odds ratio of mortality each additional year following cancer diagnosis is about 1.6, and there is moderate time-varying confounding by health status. In scenario 2, we have greater cancer-related mortality by increasing $\beta_{\bar{Y}}^D$. In scenario 3, we increase p_S, the proportion of the study population susceptible to the carcinogenic effects of MWF exposure. In scenario 4, we consider greater time-varying confounding by health status by increasing β_H^N and β_H^Y. In the last scenario, we have greater background cancer incidence by increasing β_0^Y. The sets of parameters used in the five scenarios are presented in Table 2.

Table 2: Simulation parameters.

Parameter	Scenario 1	Scenario 2	Scenario 3	Scenario 4	Scenario 5
p_S	0.10	0.10	**0.20**	0.10	0.10
p_W	0.20	0.20	0.20	0.20	0.20
p_H	0.30	0.30	0.30	0.30	0.30
β_0^N	3.00	3.00	3.00	3.00	3.00
β_W^N	-0.10	-0.10	-0.10	-0.10	-0.10
β_H^N	-0.50	-0.50	-0.50	**-1.50**	-0.50
β_A^N	-1.50	-1.50	-1.50	-1.50	-1.50
β_0^A	-1.50	-1.50	-1.50	-1.50	-1.50
β_W^A	-0.50	-0.50	-0.50	-0.50	-0.50
β_A^A	2.50	2.50	2.50	2.50	2.50
β_0^D	-5.50	-5.50	-5.50	-5.50	-5.50
β_W^D	1.00	1.00	1.00	1.00	1.00
β_A^D	0.50	0.50	0.50	0.50	0.50
$\beta_{\bar{Y}}^D$	0.50	**2.00**	0.50	0.50	0.50
β_0^Y	-7.00	-7.00	-7.00	-7.00	**-6.00**
β_W^Y	2.00	2.00	2.00	2.00	2.00
β_A^Y	0.25	0.25	0.25	0.25	0.25
$\beta_{\bar{A}}^Y$	0.20	0.20	0.20	0.20	0.20
β_H^Y	0.70	0.70	0.70	**1.70**	0.70
β_S^Y	0.30	0.30	0.30	0.30	0.30

Interventions, potential outcomes, target parameters, and estimation

The substantive question of interest was the causal effect of occupational exposure to MWF on cancer incidence risk. Since occupational MWF exposure occurs only when individuals are at work, we defined dynamic exposure regimes that depend on employment status. Under rule a_0, set $D(t) = 0$, and set $A(t) = 0$ while $N(t) = 1$. Under rule a_1, set $D(t) = 0$, and set $A(t) = 1$ while $N(t) = 1$. Under both rules, we prevented censoring by death as if it were intervenable. The causal effect was defined by contrasting the survival function $S_{a_1}(t) = 1 - \mathbb{E}\left[Y_{a_1}(t)\right]$ under rule a_1 to $S_{a_0}(t) = 1 - \mathbb{E}\left[Y_{a_0}(t)\right]$ that under rule a_0. Note that this causal estimand was defined over *a priori counterfactuals* not observable in the real world (Frangakis and Rubin 2002). This approach is standard in epidemiologic studies, however.

The survival function expresses the probability that a person following rule a is cancer-free at the end of time point t. The expected time until cancer under rule a is $\mu_a = \sum_0^K S_a(t)\, dt$. The parameter used in the estimation of bias was the summary measure $\psi = \mu_{a_1} - \mu_{a_0}$, the difference in expected time until event under two different interventions over 20 years of follow-up under five different data generating scenarios. Bias was evaluated by comparing estimates of ψ to its true value in 250 simulations per scenario (the original analysis performed 500). The true value was calculated by simulating the full data for five hundred thousand individuals (the original analysis used one million) with rules a_0 and a_1 applied deterministically. Estimates of ψ were obtained by first estimating the survival curves $S_a(t)$ using two estimators: the inverse probability weighted Kaplan-Meier estimator (WKM) and the Aalen-filtered WKM (AWKM). These survival estimators are detailed in the following section.

Kaplan-Meier estimator and extensions

To estimate survival, we applied extensions of the widely-known Kaplan-Meier (KM) estimator for survival (Kaplan and Meier 1958). First, we review the estimator of Xie and Liu (2005), an extension of the KM estimator where units are weighted by the inverse probability of treatment. The standard KM estimator requires counting up the number of cases $c_a^0(t)$ that occurred in interval $(t-1, t]$ and the number of units at risk $R_a^0(t)$ in that interval at all event times t. Assuming cancer status was assessed at the end of regular intervals $t = 1, \ldots, K$, we have:

$$c_a^0(t) = \sum_i^n \mathbb{1}\left[Y_i(t) = 1\right] \times \mathbb{1}\left[Y_i(t-1) = 0\right] \times \mathbb{1}\left[\bar{A}_i(t) = \bar{a}(t)\right]$$
$$R_a^0(t) = \sum_i^n \mathbb{1}\left[Y_i(t-1) = 0\right] \times \mathbb{1}\left[\bar{A}_i(t) = \bar{a}(t)\right].$$

The standard survival estimator is

$$\hat{S}_a^0(t) = \begin{cases} 1 & \text{if } t < t_1 \\ \prod_{j \leq t}\left(1 - \frac{c_a^0(j)}{R_a^0(j)}\right) & \text{if } t \geq t_1 \end{cases}$$

where t_1 is the first event time.

In observational studies, survival contrasts estimated using the standard KM estimator are biased for the true causal survival contrast. However, if conditional ignorability and positivity are attained, the inverse probability weighted KM (WKM) estimator of Xie and Liu (2005) yields unbiased estimates of the true causal survival curve. The WKM estimator augments the standard KM estimator by weighting units at time t by $w_{i,a}(t)$ the inverse probability of treatment:

$$c_a^w(t) = \sum_i^n w_{i,a}(t) \times \mathbb{1}\left[Y_i(t) = 1\right] \times \mathbb{1}\left[Y_i(t-1) = 0\right] \times \mathbb{1}\left[\bar{A}_i(t) = \bar{a}(t)\right]$$
$$R_a^w(t) = \sum_i^n w_{i,a}(t) \times \mathbb{1}\left[Y_i(t-1) = 0\right] \times \mathbb{1}\left[\bar{A}_i(t) = \bar{a}(t)\right]$$

The WKM survival estimator for rule a is

$$\hat{S}_a^w(t) = \begin{cases} 1 & \text{if } t < t_1 \\ \prod_{j \leq t}\left(1 - \frac{c_a^w(j)}{R_a^w(j)}\right) & \text{if } t \geq t_1 \end{cases}$$

where t_1 is the first event time.

Finally, to account for (uninformative) left filtering, we applied the Aalen filter, which considers only the units at time t for which the outcome is observed:

$$c_a(t) = \sum_i^n w_{i,a}(t) \times \mathbb{1}\left[Y_i(t) = 1\right] \times \mathbb{1}\left[Y_i(t-1) = 0\right] \times \mathbb{1}\left[\bar{A}_i(t) = \bar{a}(t)\right] \times \mathbb{1}\left[t \geq R_i\right]$$
$$R_a(t) = \sum_i^n w_{i,a}(t) \times \mathbb{1}\left[Y_i(t-1) = 0\right] \times \mathbb{1}\left[\bar{A}_i(t) = \bar{a}(t)\right] \times \mathbb{1}\left[t \geq R_i\right]$$

The Aalen-filtered WKM (AWKM) estimator for rule a is

$$\hat{S}_a(t) = \begin{cases} 1 & \text{if } t < t_1 \\ \prod_{j \leq t}\left(1 - \frac{c_a(j)}{R_a(j)}\right) & \text{if } t \geq t_1 \end{cases}$$

where t_1 is the first event time.

In the full data, the WKM and AWKM estimators are equivalent, and identification is achieved under overlap (positivity) and sequential ignorability (randomization) assumptions:

$$Y^*_{a,\bar{d}=0}(t') \perp\!\!\!\perp A(t) \mid W,\ \bar{A}(t-1) = \bar{a}(t-1),\ D(t-1) = 0,\ N(t) = 1$$
$$Y^*_{a,\bar{d}=0}(t') \perp\!\!\!\perp D(t) \mid W,\ D(t-1) = 0,\ Y^*(t-1) = 0,\ \bar{A}(t-1) = \bar{a}(t-1)$$

for all times $t' \geq t$, and

$$0 < \mathbb{P}\left(A(t) = 1 \mid W,\ \bar{A}(t-1) = \bar{a}(t-1),\ D(t-1) = 0,\ N(t) = 1\right) < 1$$

$$0 < \mathbb{P}\left(D(t) = 0 \mid W,\ D(t-1) = 0,\ Y^*(t-1) = 0,\ \bar{A}(t-1) = \bar{a}(t-1)\right) < 1.$$

Graphical representations of the first and second components of the ignorability assumption are presented in Figures 2 and 3 where conditioning on boxed variables are represented by the removal of edges pointing away from those variables. The resulting graphs show the fulfillment of Pearl's backdoor criterion for the estimation of the causal effects of $\bar{A}(t)$ on $Y^*(t)$ and $D(t)$ on $Y^*(t)$, respectively. Thus, the causal effect of the joint intervention on $\left(\bar{A}(t), D(t)\right)$ at each time t is identified. Causal identification is not attained when true cancer status $Y^*(t)$ is not known.

Figure 2: Directed acyclic graph representing the causal relationships encoded in the non-parametric structural equation model at time t after conditioning on $\{W,\ \bar{A}(t-1),\ D(t-1),\ N(t)\}$.

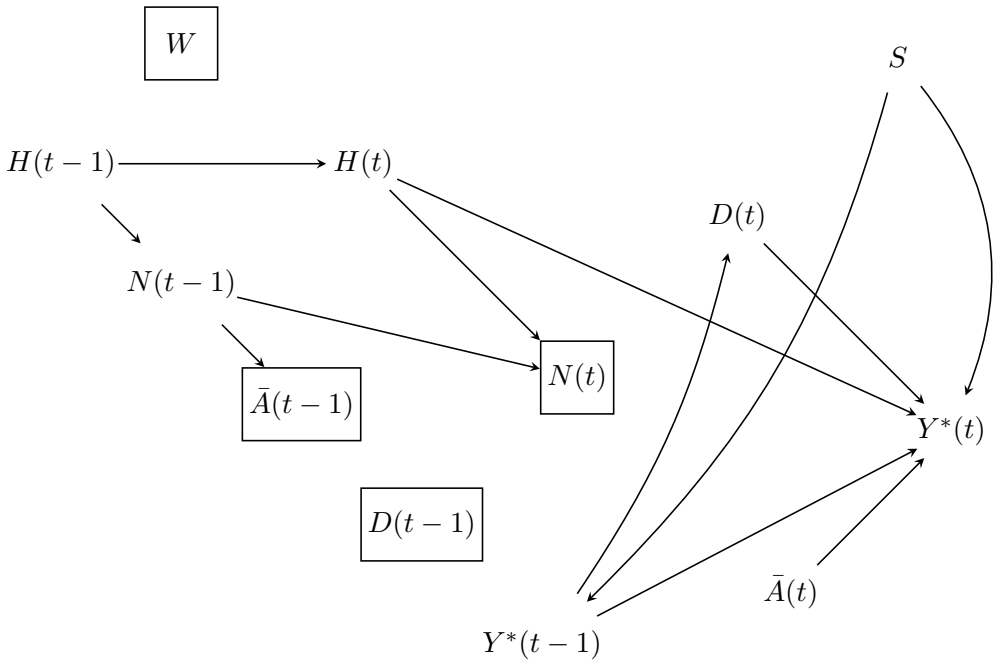
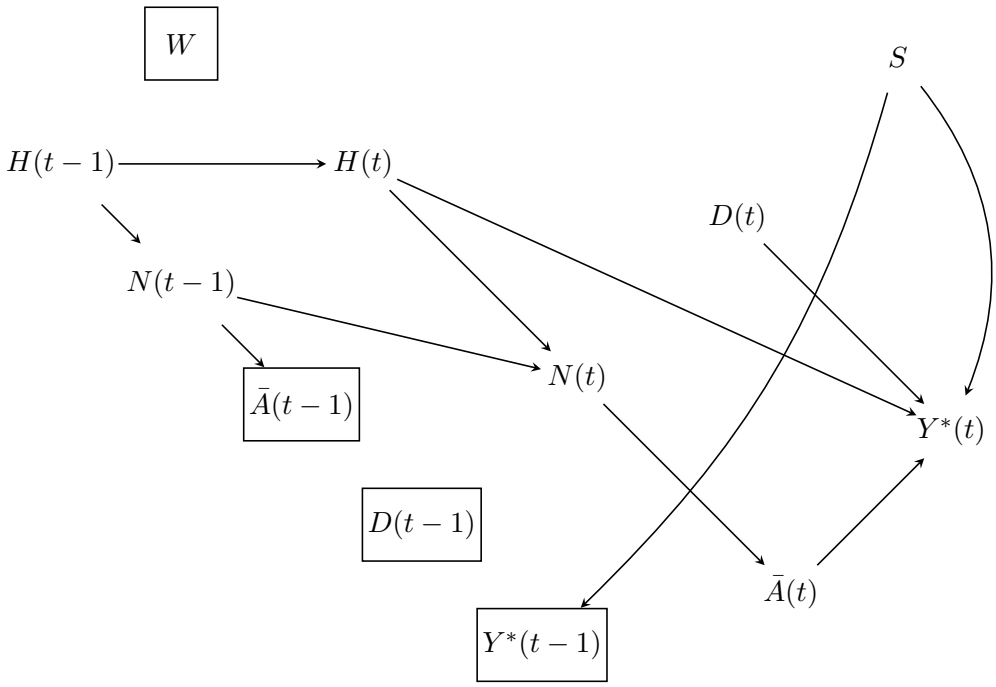


Figure 3: Directed acyclic graph representing the causal relationships encoded in the non-parametric structural equation model at time t after conditioning on $\{W,\ \bar{A}(t-1),\ D(t-1),\ Y^*(t-1)\}$.



Estimation of weights

To estimate the weights for the WKM and AWKM estimators, we fit two logistic regressions at each time point $t = 1, \ldots 20$:

$$\text{logit}\left(\mathbb{P}\left(A(t) = 1 \mid W,\ \bar{A}(t-1),\ D(t-1) = 0,\ N(t) = 1\right)\right) = \alpha_0 + W\alpha_1 + A(t-1)\alpha_2$$
$$\text{logit}\left(\mathbb{P}\left(D(t) = 1 \mid W,\ D(t-1) = 0,\ Y(t-1) = 0,\ \bar{A}(t-1)\right)\right) = \beta_0 + W\beta_1 + \bar{A}(t-1)\beta_2$$

The first was fit on data for those alive and at work at time t. The second was among those alive and (observed to be) cancer-free. For each unit at time t, the weight was calculated by taking the inverse of the cumulative probability of following the exposure rule and remaining uncensored:

$$\hat{w}_a(t) = \left[\prod_{j=1}^{t} \begin{array}{r} \widehat{\mathbb{P}}\left\{A(j) = a(j) \mid W,\ \bar{A}(j-1) = \bar{a}(j-1),\ D(j-1) = 0,\ N(j) = 1\right\} \times \\ \widehat{\mathbb{P}}\left\{D(j) = 0 \mid W,\ D(j-1) = 0,\ Y(j-1) = 0,\ \bar{A}(j) = \bar{a}(j)\right\} \end{array}\right]^{-1}.$$

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Kaplan, Edward L, and Paul Meier. 1958. "Nonparametric Estimation from Incomplete Observations." *Journal of the American Statistical Association* 53 (282): 457–81.

Xie, Jun, and Chaofeng Liu. 2005. "Adjusted Kaplan–Meier Estimator and Log-Rank Test with Inverse Probability of Treatment Weighting for Survival Data." *Statistics in Medicine* 24 (20): 3089–3110.