

A Statistical Framework for Understanding Causal Effects that Vary by Treatment Initiation Time in EHR-based Studies

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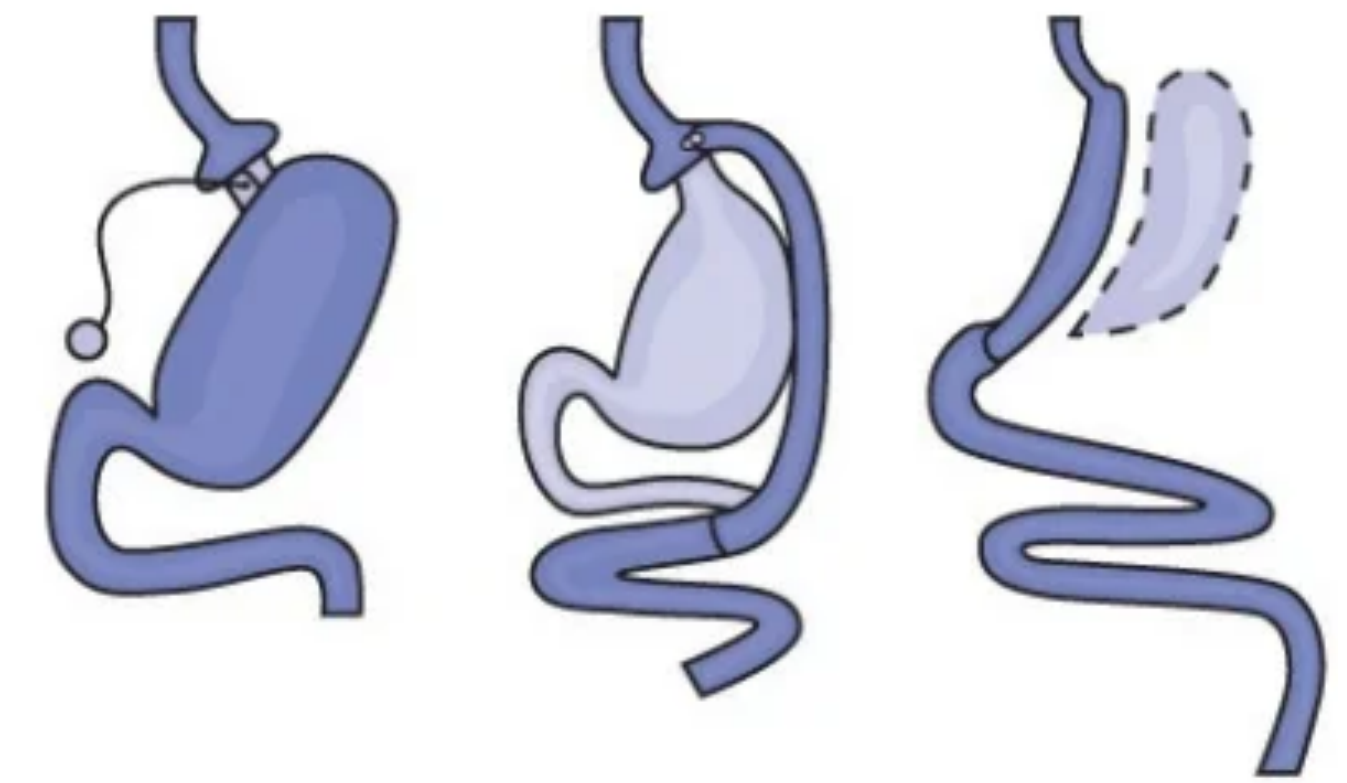
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Bariatric Surgery

- Bariatric surgery is a weight-loss surgery
 - Typical candidates have BMI $\geq 35\text{kg/m}^2$
- Sleeve Gastrectomy (SG) is a newer procedure than Roux-en-Y Gastric Bypass (RYGB)
 - SG surpassed RYGB in popularity in late 2000s/early 2010s
 - Less invasive and technically less complex
- DURABLE: NIH funded study of long-term outcomes following bariatric surgery, particularly in relation to non-surgical patients
 - Kaiser Permanente (Washington, N. California, S. California)
 - 1997-2015; ~45,000 surgical patients and 1.7 million non surgical patients

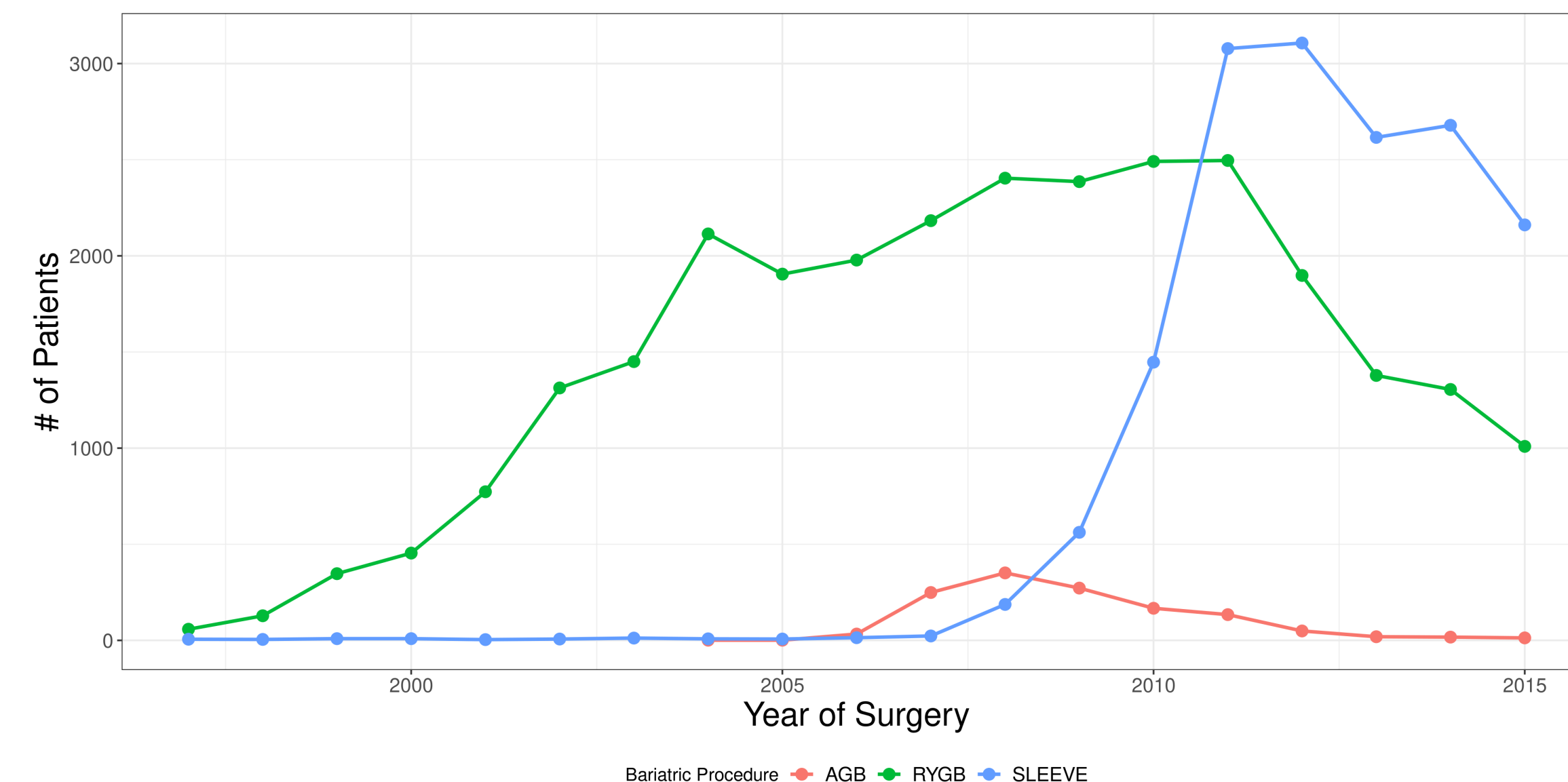


Adjustable
Gastric Band
(AGB)

Roux-en-Y
Gastric Bypass
(RYGB)

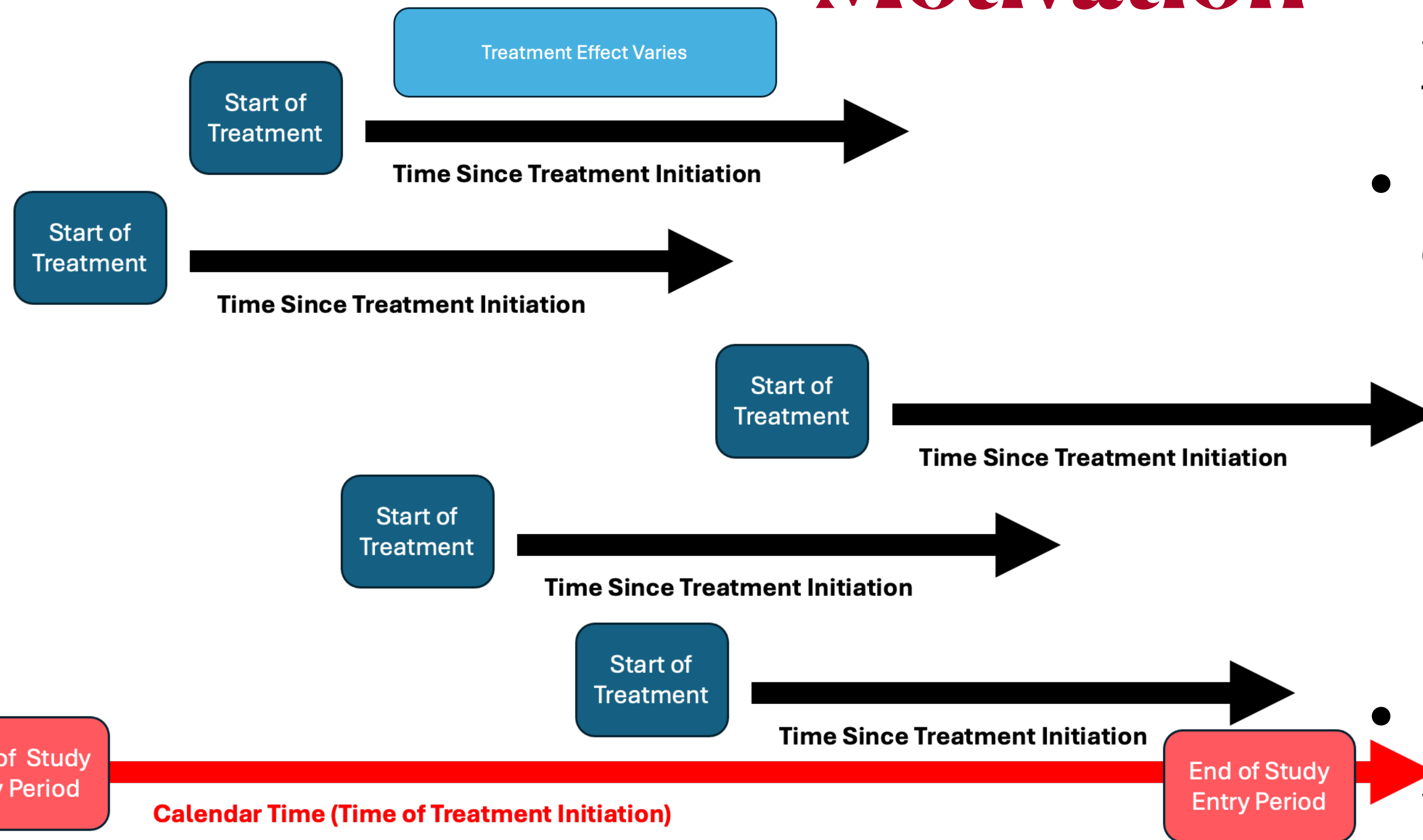
Vertical Sleeve
Gastrectomy
(VSG)

Distribution of Bariatric Surgery Procedures
DURABLE Database: 1997-2015



Motivation

- **Time-varying effects** usually speaks to time **since** starting a treatment
- Real-world treatments evolve over time
 - Standard of care (best practice)
 - Procedures where techniques can change (bariatric surgery)
- Causal effects in EHR-based studies with long study entry periods may vary over the course **of treatment initiation time (calendar time)**



Calendar Time as a Source of Bias?

To obtain valid effect estimates in a nonrandomized trial, all **baseline confounders** have to be appropriately measured and adjusted for in the analysis. We proceeded as if this condition was at least approximately true in the NHS non-randomized “trial” once we added the following covariates to the Cox model: parental history of myocardial infarction before age 60 (yes, no), education (graduate degree: yes, no), husband’s education (less than high school, high school graduate, college, graduate school), ethnicity (non-Hispanic white, other), age at menopause (<50, 50–53, >53), **calendar time**, high cholesterol (yes, no), high blood pressure (yes, no), diabetes (yes, no), angina (yes, no), stroke (yes, no), coronary revascularization (yes, no), osteoporosis (yes, no), body mass index (<23, 23–<25, 25–<30, ≥30), cigarette smoking (never, past, current 1–14 cigarettes per day, current 15–24 cigarettes per day, current ≥25 cigarettes per day), aspirin

studies need to address several important sources of bias, including confounding due to changes in the clinical use of treatments over time (i.e., channeling bias) [2]. Channeling occurs when users of a medication and its potential comparators are selectively prescribed a particular agent based on differences in underlying patient characteristics [3]. If these characteristics also affect the risk for the outcome of interest, channeling will lead to confounding.

The impact of **calendar time-specific channeling** is a particularly important consideration when assessing the comparative safety for drugs within a medication class targeted by a Food and Drug Administration (FDA) safety advisory. In such cases, when reported adverse effects vary among drugs within a therapeutic class, clinicians may respond by channeling patients toward or away from a particular treatment alternative based on the patient’s underlying risk for an adverse outcome.

One way to address **calendar time-specific channeling** is through the use of **calendar time-specific propensity scores** [4].

TZD use over a short period also provides an opportunity to use calendar time as an instrumental variable (IV).¹³ **Instrumental variable methods** serve as a natural experiment to compare drugs in settings where substantial unmeasured confounding is expected.^{14–16} Common in pharmacoepidemiology includes provider preference, calendar time, distance to care, and geographic variation.¹⁷ **Calendar time IV** arises from secular trends in medication use. Such trends can be a result of effectiveness or safety releases, changes in physician prefer-

Formulas for these calculations are provided in [Appendix S1](#). A further benefit of including more cases in one model is increased precision when **adjusting for calendar time** or age effects, **assuming common age/calendar-time temporal effects**.¹⁰

Common viewpoint in pharmacoepi is that this is a potential source of bias

- Towards estimating an implicitly assumed **constant** effect (across calendar time)
- Reasonable viewpoint if treatments (e.g., drugs or medications) are not changing themselves
- But what if our treatment strategies have to be more loosely defined?
 - Calendar-time varying effects = fundamental changes in treatment efficacy?
- In this project, we introduce a framework for
 1. Efficient estimation of causal effects that may vary by calendar time
 2. Determining **how** effects vary by treatment time
 3. Determining **why** effects vary by treatment time

An EHR-based Study of Weight Loss Following Bariatric Surgery

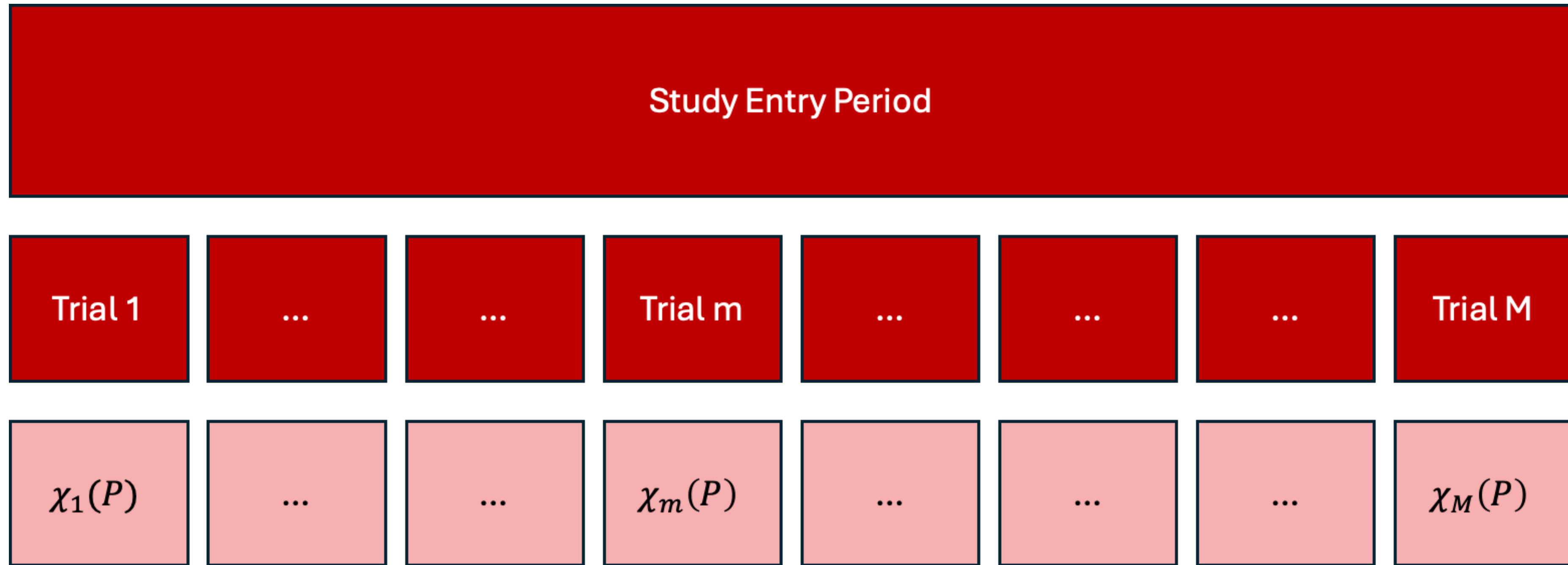
- Examine relative weight change at 4 pre-specified times post-surgery
 - {6 months, 1 year, 2 years, 3 years}
- DURABLE database between 2005-2011
 - 17,905 surgical patients
 - 933,044 non-surgical patients
- Sequence of 84 target trials (one per month between Jan. 2005 and Dec. 2011)
 - Necessary to define “time zero”
- Treatment comparisons
 - Surgery vs. No Surgery
 - {RYGB, SG} vs. No-Surgery (separately)
 - RYGB vs. SG

Notation

- Trial index $m \in \{1, \dots, M\}$ ($M = 84$ in our running example)
- $\mathbf{Z}_m = (\mathbf{L}_m, A_m, Y_m)$
 - Y_m : continuous outcome (e.g., % weight loss from baseline of trial m)
 - A_m : Binary treatment
 - $\mathbf{L}_m$: Covariates (confounders, effect modifiers)
- E_m : Eligibility indicator

$$\mathbf{O}_1, \dots, \mathbf{O}_n \stackrel{iid}{\sim} P$$
$$\mathbf{O} = (E_1, E_1 \mathbf{Z}_1, \dots, E_m, E_m \mathbf{Z}_m, \dots, E_M, E_M \mathbf{Z}_M)$$

Calendar Time-Specific Average Treatment Effect



$$\chi_m(P) = \mathbb{E}_P[Y_m(1) - Y_m(0) \mid E_m = 1]$$

- Leverage the sequential nature of the design to define initiator cohorts across calendar time
- Implicit in the definition of $\chi_m(P)$ is that comparisons are being made between whatever “versions” of treatment are in use at trial m
 - ITT analogue — don’t care what happens after trial m

Nonparametric identification + efficiency theory

1. **Consistency** $Y_m(A_m) = Y_m$ when $E_m = 1$, almost surely, for all m
2. **Positivity** $0 < \epsilon \leq P(A_m = 1 \mid \mathbf{L}_m, E_m = 1) \leq 1 - \epsilon < 1$ almost surely, for all m
3. **No Unmeasured Confounding** $Y_m(a_m) \perp\!\!\!\perp A_m \mid \mathbf{L}_m, E_m = 1$ for all m

At the outset, estimation of $\chi_m(P)$ may seem straightforward, for example on the basis of it's nonparametric influence function:

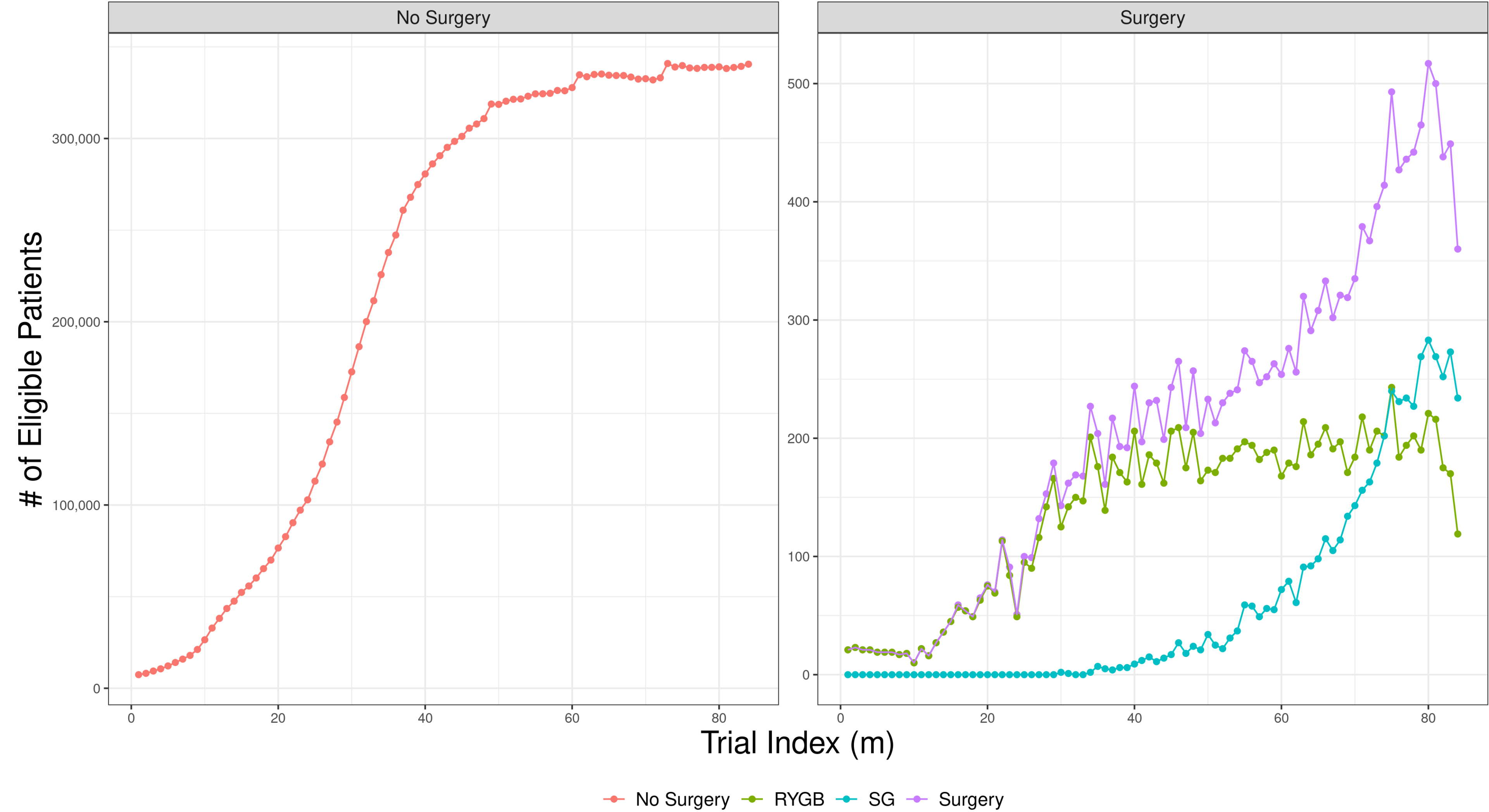
$$\dot{\chi}_m^*(O; P) = \frac{\mathbf{1}(E_m = 1)}{P(E_m = 1)} \left\{ \mu_m(1, \mathbf{L}_m) - \mu_m(0, \mathbf{L}_m) - \chi_m(P) + \left(\frac{A_m}{\pi_m(\mathbf{L}_m)} - \frac{1 - A_m}{1 - \pi_m(\mathbf{L}_m)} \right) \left(Y_m - \mu_m(A_m, \mathbf{L}_m) \right) \right\}$$

$$\mu_m(a_m, \mathbf{L}_m) = \mathbb{E}[Y_m \mid A_m = a_m, \mathbf{L}_m, E_m = 1]$$

$$\pi_m(\mathbf{L}_m) = P(A_m = 1 \mid \mathbf{L}_m)$$

But...how much data do we have at trial m ?

Trial Effective Sample Sizes



Marginal Structural Model + Projection Parameter Approach

- For a fixed m , insufficient for **ML-based** nuisance models
- Ultimately, our interest is in the **trend** in $\chi_m(P)$ across calendar time (m)
- Can we pool **across trials** without imposing strong parametric assumptions?

Marginal Structural Model + Projection Parameter Approach

- For a fixed m , insufficient for **ML-based** nuisance models
- Ultimately, our interest is in the **trend** in $\chi_m(P)$ across calendar time (m)
- Can we pool **across trials** without imposing strong parametric assumptions?
- Consider the marginal structural model (MSM)

$$\mathbb{E}[Y_m(a_m = 1) - Y_m(a_m = 0) \mid E_m = 1] \approx \psi(m; \beta)$$

- When MSM is saturated, model is correct
- Otherwise, we adopt **assumption-lean** projection approach:

$$\beta(P) = \operatorname{argmin}_{\beta \in \mathbb{R}^k} \sum_{m=1}^M w(m) P(E_m = 1) \left(\chi_m(P) - \psi(m; \beta) \right)^2$$

Benefits of Projection Parameter Approach

- Projection approach (Neugebauer & van Der Laan, 2007) is inherently model-agnostic i.e., valid even if MSM not correctly specified
 - Greater transparency about how information is shared across time
 - We combine with **pooled** but **flexible** nuisance models
- Directly feeds into model selection strategy for comparing candidate MSMs
 - Identify functional form of **how** treatment effects vary over calendar time
 - Including the possibility of a constant effect

Estimation Strategy

- Estimation/inference on trend still require estimates of trial-specific effects
 - Use pooled models for nuisance functions $\tilde{\mu}(A_m, \mathbf{L}_m, m)$ and $\tilde{\pi}(\mathbf{L}_m, m)$ estimated on pooled dataset

$$\mathcal{D} = \bigcup_{m=1}^M \left\{ (\mathbf{L}_{m,i}, A_{m,i}, Y_{m,i}, m) \right\}_{i:E_{m,i}=1}$$

- Use nonparametric/flexible machine learning models for pooled nuisance functions for influence-function based estimator

$$\hat{\chi}_m = \mathbb{P}_n \left[\frac{\mathbf{1}(E_m = 1)}{\mathbb{P}_n[E_m = 1]} \left\{ \hat{\mu}_m(1, \mathbf{L}_m) - \hat{\mu}_m(0, \mathbf{L}_m) - \left(\frac{A_m}{\hat{\pi}_m(\mathbf{L}_m)} - \frac{1 - A_m}{1 - \hat{\pi}_m(\mathbf{L}_m)} \right) \left(Y_m - \hat{\mu}_m(A_m, \mathbf{L}_m) \right) \right\} \right]$$

Model Selection Among Candidate MSMs

- Let $\{\hat{\psi}_1, \dots, \hat{\psi}_K\}$ denote a set of K candidate MSMs

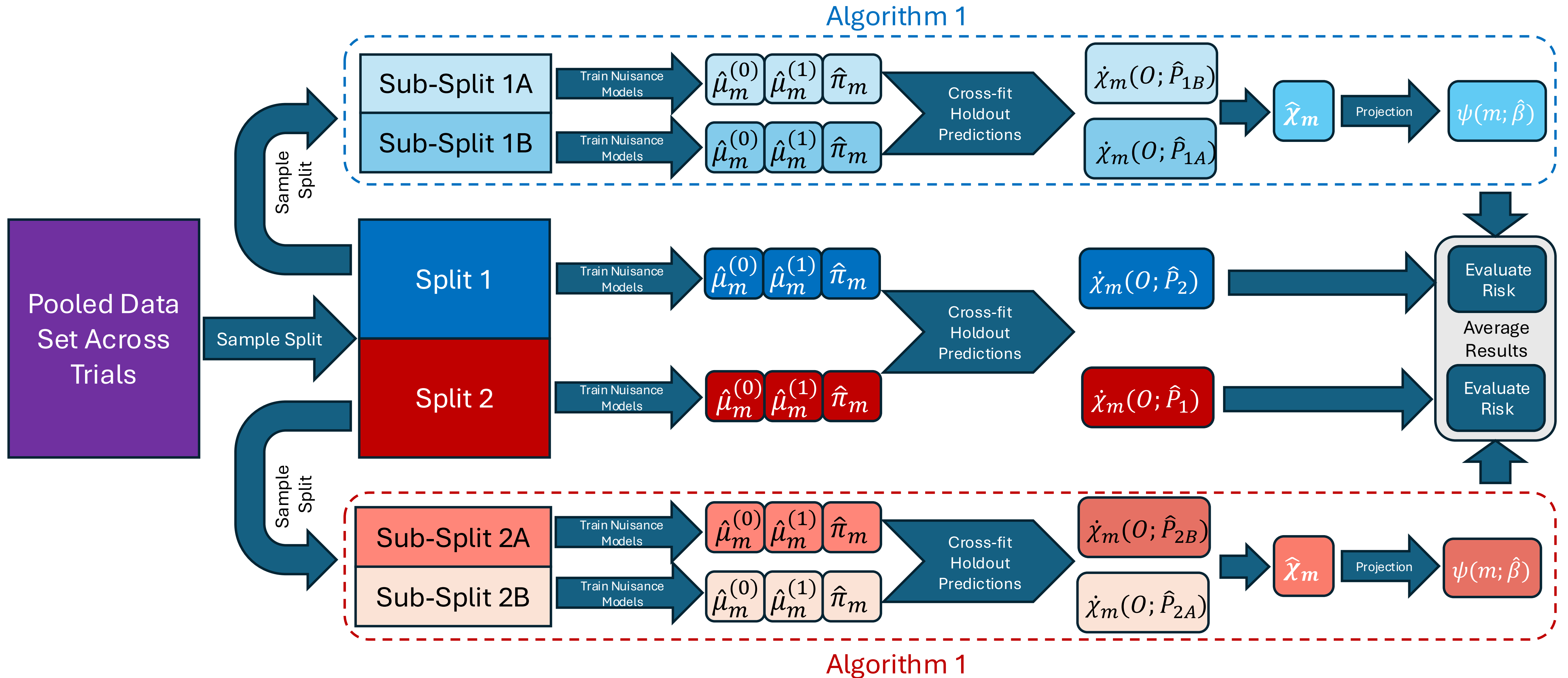
- Constant $\psi(m; \beta) = \beta$;
- Linear $\psi(m; \beta) = \beta_0 + \beta_1 m$;
- Higher order polynomials and splines

- Loss Function:

$$L(\hat{\psi}_k) = \mathbb{P}_n \left[\sum_{m=1}^M w(m) \{ \psi_k(m; \hat{\beta}_k)^2 - 2\psi_k(m; \hat{\beta}_k) \dot{\chi}_m(O; \hat{P}) \} \right]$$

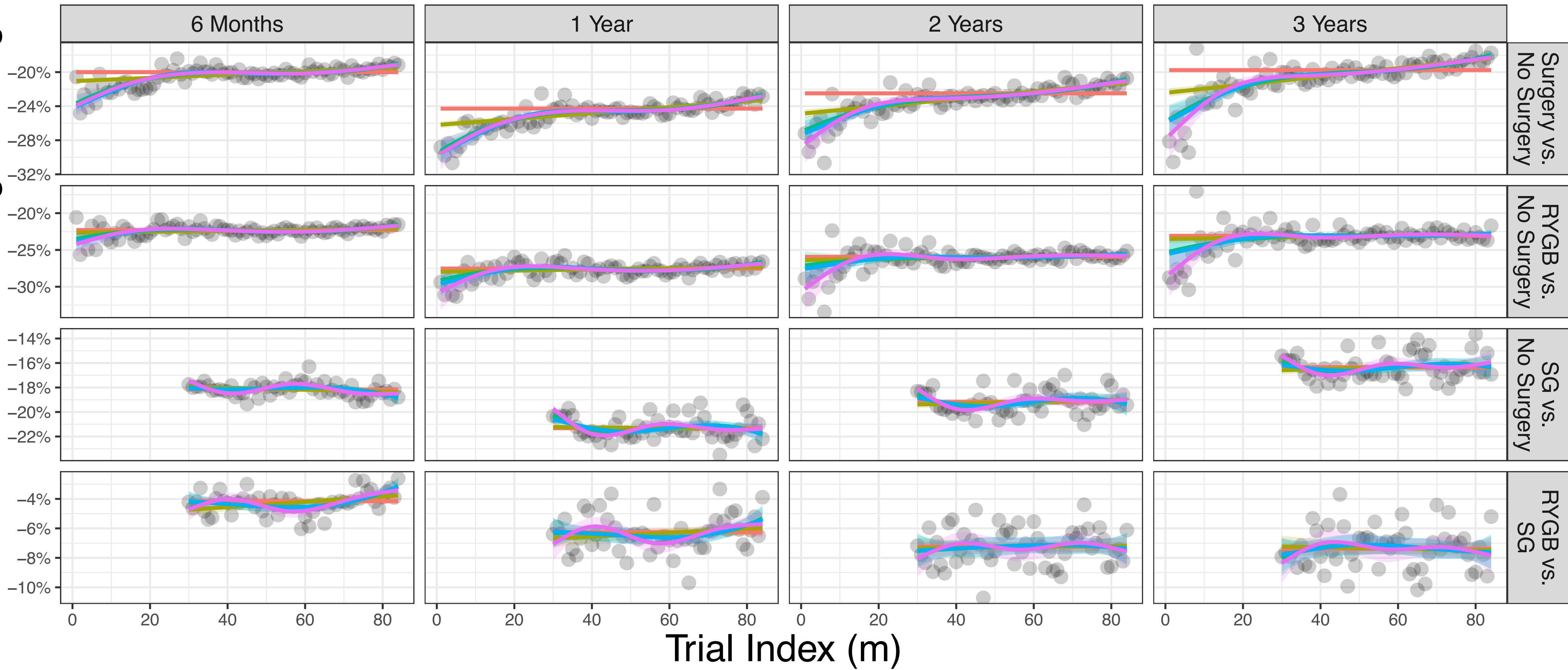
- Reasonable loss function to work with because it reflects estimating equation-based estimator of (pseudo)risk if one treats $\hat{\psi}_k$ as fixed

Complete Estimation and Selection Framework



Calendar Time–Specific Treatment Effect Estimates

Difference in Mean %–Weight Change



$\psi(m;\beta)$ — Constant — Linear — Cubic — Spline (2 Knots) — Spline (3 Knots)

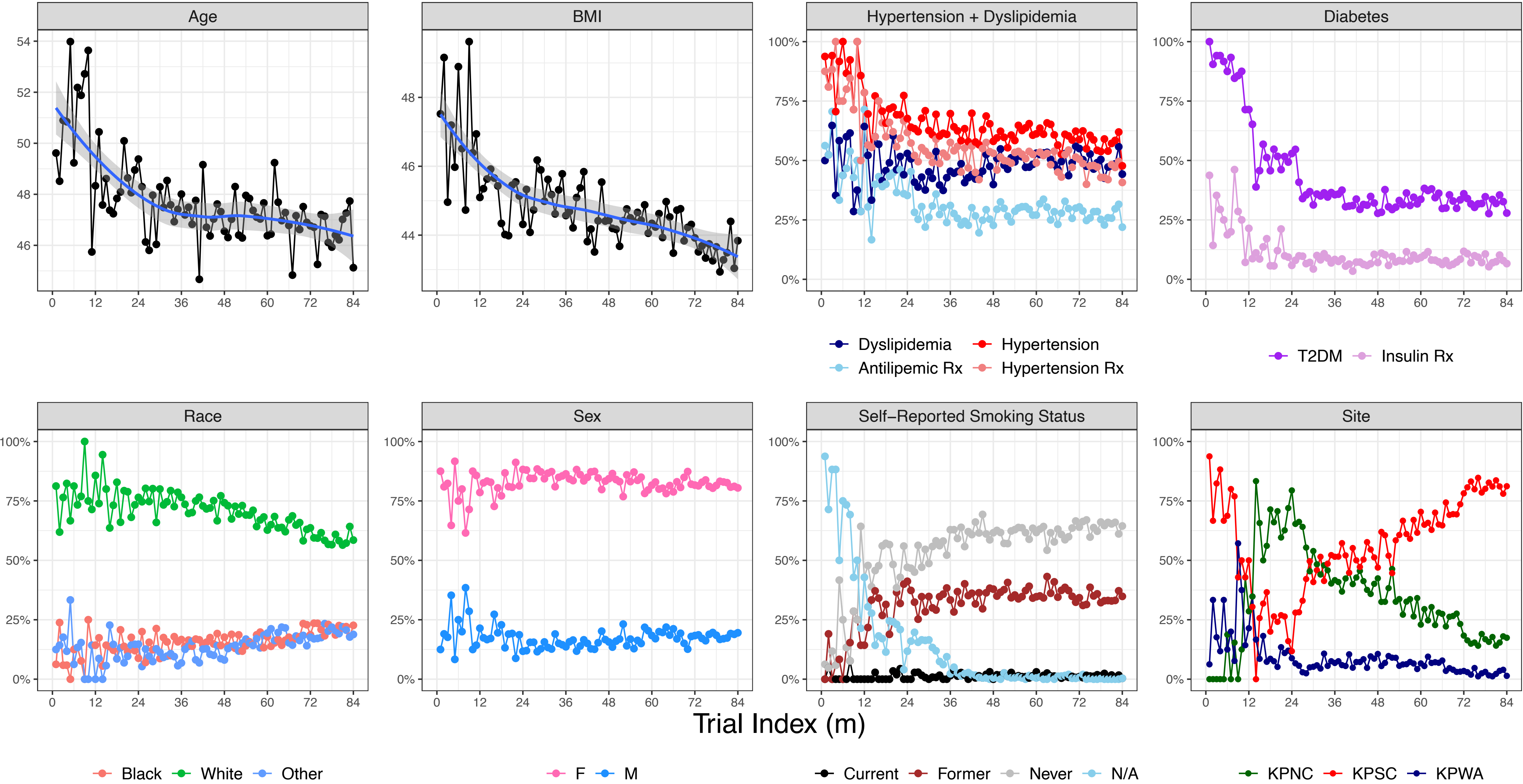
Results

Comparison	Outcome	Minimizing MSM	Selected MSM	95% Interval for θ
Surgery vs. No Surgery	6 Months	Spline (3 Knots)	Cubic	(0.684, 0.809)
	1 Year	Spline (3 Knots)	Spline (3 Knots)	(0.648, 0.767)
	2 Years	Spline (2 Knots)	Spline (2 Knots)	(0.644, 0.765)
	3 Years	Spline (3 Knots)	Spline (3 Knots)	(0.682, 0.777)
RYGB vs. No Surgery	6 Months	Cubic	Cubic	(0.668, 0.821)
	1 Year	Spline (3 Knots)	Spline (3 Knots)	(0.658, 0.795)
	2 Years	Spline (3 Knots)	Spline (3 Knots)	(0.647, 0.785)
	3 Years	Spline (3 Knots)	Spline (3 Knots)	(0.670, 0.787)
SG vs. No Surgery	6 Months	Spline (3 Knots)	Spline (3 Knots)	(0.971, 0.989)
	1 Year	Constant	Constant	(0.974, 0.991)
	2 Years	Linear	Constant	(0.980, 0.992)
	3 Years	Constant	Constant	(0.979, 0.992)
RYGB vs. SG	6 Months	Spline (3 Knots)	Cubic	(0.718, 0.807)
	1 Year	Constant	Constant	(0.670, 0.784)
	2 Years	Constant	Constant	(0.666, 0.784)
	3 Years	Constant	Constant	(0.688, 0.793)

Table 2: Summary of model selection results and 95% bootstrapped intervals for θ . The minimizing MSM denotes $\hat{\psi}^*$ minimizing $L(\hat{\psi})$, while the selected MSM denotes the simplest model within $c = 0.25$ weighted standard deviations of the minimizer.

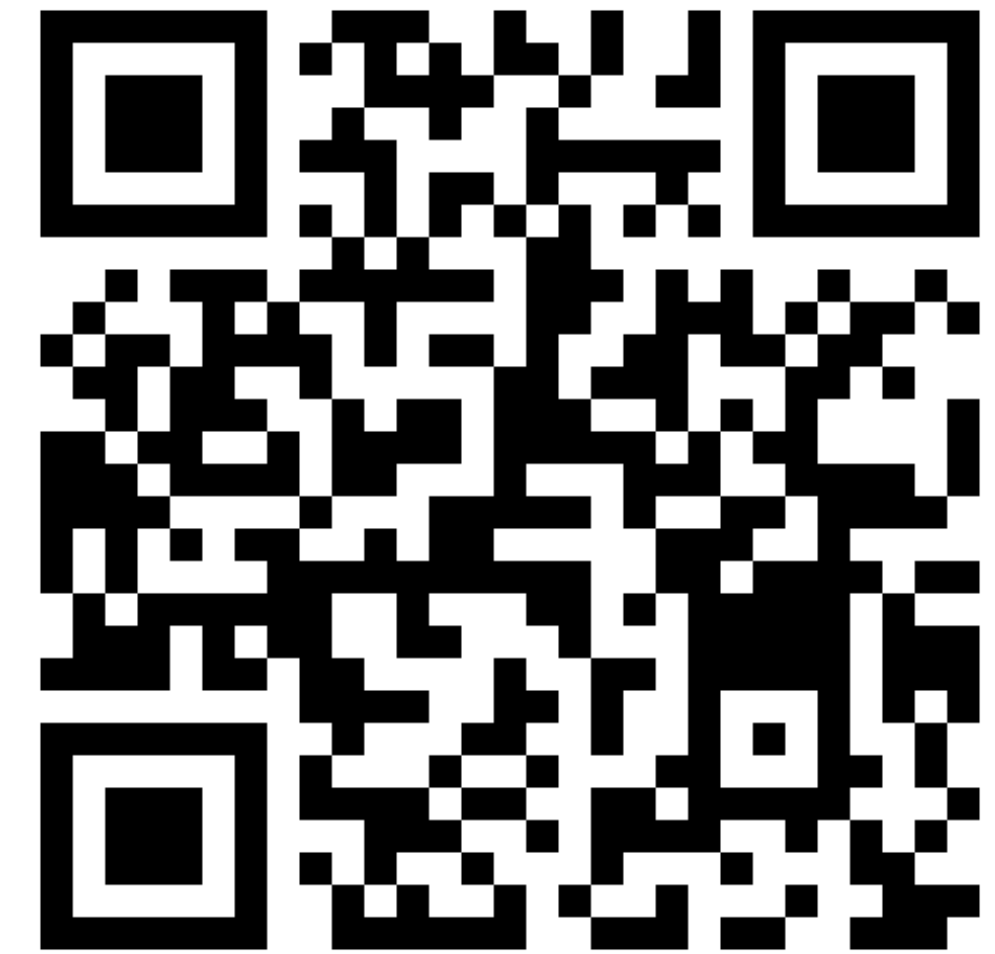
Distribution of Key Covariates in Study Population

Eligible Patients Undergoing Bariatric Surgery



Results Summary

- **Surgery vs. No-Surgery**
 - 3 Years: -27.4% at $m = 1$ vs. -18.3% at $m = 84$
[Change of 9.1%]
 - Constant effect: -19.8%
 - **Misses a lot of the story**



Paper

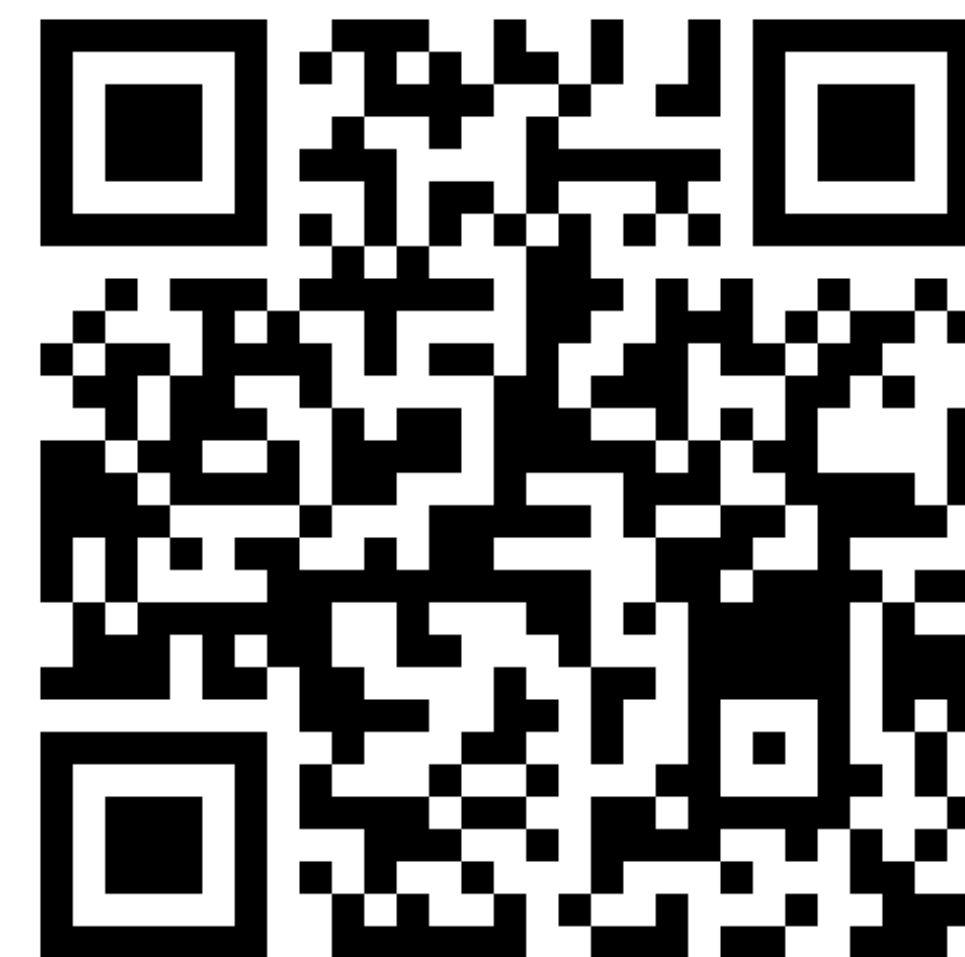


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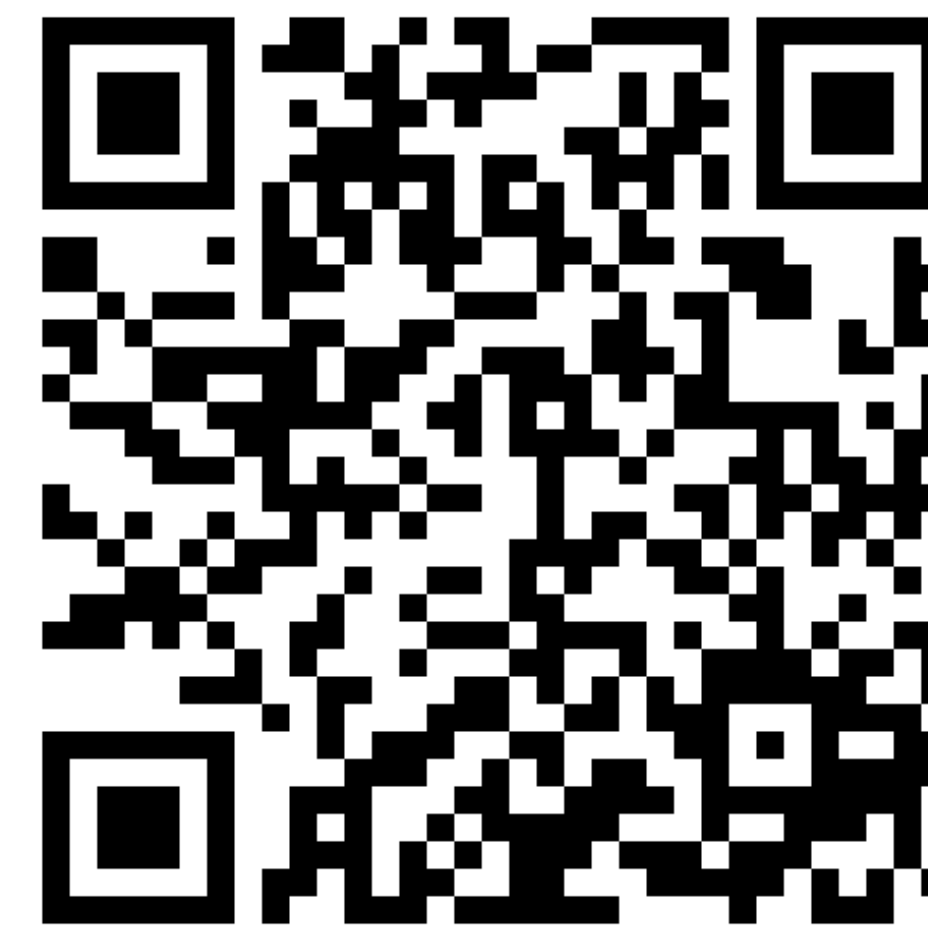
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- Constant effect: -19.8%
 - **Misses a lot of the story**
- Not surprising that surgery is “less effective” over time because distribution of procedures is changing
 - Good validation that our method is picking up clinically meaningful changes
 - Many papers use catch-all of “bariatric surgery” → **what that label means is changing**



Paper

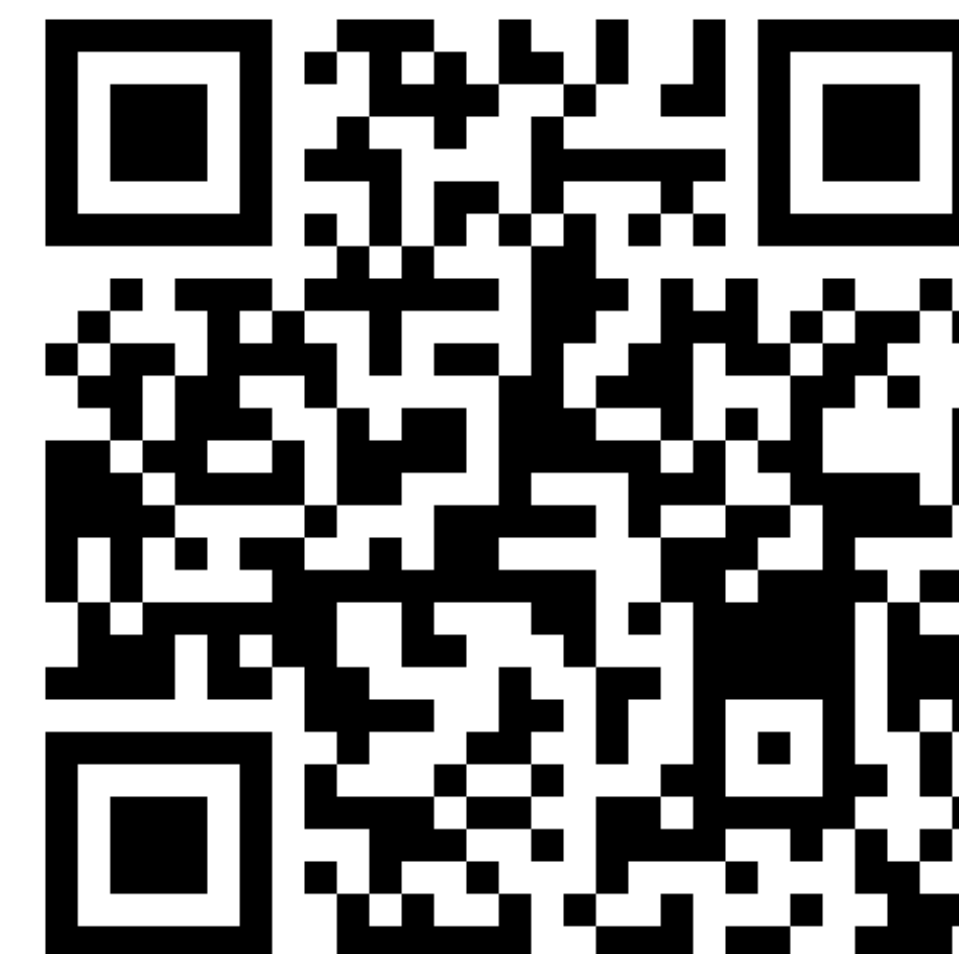


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- Not surprising that surgery is “less effective” over time because distribution of procedures is changing
 - Good validation that our method is picking up clinically meaningful changes
 - Many papers use catch-all of “bariatric surgery” → **what that label means is changing**
- Tools for disentangling changes in treatment effect from changes in population receiving treatment → see paper



Paper



GitHub

Appendix



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Estimation Strategy

- As long as our candidate MSM is twice differentiable in β , then under A1-A3, $\beta(P)$ is identified as the solution to the estimating equation

$$0 = \sum_{m=1}^M w(m) P(E_m = 1) \nabla_{\beta} \psi(m; \beta) [\chi_m(P) - \psi(m; \beta)]$$

- Furthermore, the influence function of $\beta(P)$ (up to proportionality) is given by

$$\dot{\beta}^*(O; P) \propto \dot{\beta}^{\dagger}(O; P) = \sum_{m=1}^M P(E_m = 1) w(m) \nabla_{\beta} \psi(m; \beta) |_{\beta=\beta(P)} \left(\dot{\chi}_m^{\dagger}(O; P) - \psi(m; \beta(P)) \right)$$

$$\dot{\chi}_m^{\dagger}(O; P) = \mu_m(1, \mathbf{L}_m) - \mu_m(0, \mathbf{L}_m) + \left(\frac{A_m}{\pi_m(\mathbf{L}_m)} - \frac{1 - A_m}{1 - \pi_m(\mathbf{L}_m)} \right) \left(Y_m - \mu_m(A_m, \mathbf{L}_m) \right)$$

- Motivates estimator which solves $\mathbb{P}_n[\dot{\beta}^{\dagger}(O; \hat{P})] = 0$, that is

$$0 = \sum_{m=1}^M w(m) \mathbb{P}_n(E_m = 1) \nabla_{\beta} \psi(m; \beta) |_{\beta=\hat{\beta}} \left\{ \boxed{\hat{\chi}_m} - \psi(m; \hat{\beta}) \right\}$$

A Cross-Trial Contrast to Remove Distribution Shift

$$\chi_{j,m}(P) = \int_{\mathcal{L}} \mathbb{E} \left[Y_m(a_m = 1) - Y_m(a_m = 0) \mid L_m = \ell, g(L_m) = 1 \right] dP_{L_j|E_j=1}(\ell \mid E_j = 1)$$

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- Difference in mean counterfactual outcomes at time m

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- Re-weight (standardize) the covariate distribution among eligible population at time m to match the covariate distribution of eligible population at time j

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- Difference in mean counterfactual outcomes at time m
- Re-weight (standardize) the covariate distribution among eligible population at time m to match the covariate distribution of eligible population at time j
- Critical to the definition of $\chi_{j,m}(P)$ is a common population with which to standardize treatment effects to across calendar time
- $m_1 \neq m_2$ differences between $\chi_{j,m_1}(P)$ and $\chi_{j,m_2}(P)$ not be attributable to covariate shift
 - Underlying population on which the causal contrast is defined is the same
 - Will exploit this to quantify the role of effect modification over time

Summarizing Effect Modification

$$\sigma_m^2 = \frac{1}{M-1} \sum_{j=1}^M \left(\chi_m(P) - \chi_{m,j}(P) \right)^2$$

$$\gamma_m^2 = \frac{1}{M-1} \sum_{j=1}^M \left(\chi_m(P) - \chi_{j,m}(P) \right)^2$$

- σ_m^2 : How much variation in treatment effects had trial m population received treatment at other points in time (**Change treatment time for fixed population**)
- γ_m^2 : characterizes the variability around $\chi_m(P)$ across all trial-eligible populations, had each been treated at time m (**Change population for fixed treatment time**)

$$\theta = \frac{1}{M} \sum_{m=1}^M \theta_m = \frac{1}{M} \sum_{m=1}^M \frac{\sigma_m^2}{\sigma_m^2 + \gamma_m^2}$$

- Close to 0: variation driven by differences in population (effect modification)
- Close to 1: variation not driven by differences in population (treatment efficacy)

Illustration of $\hat{\chi}_{j,m}$ for Select Comparisons

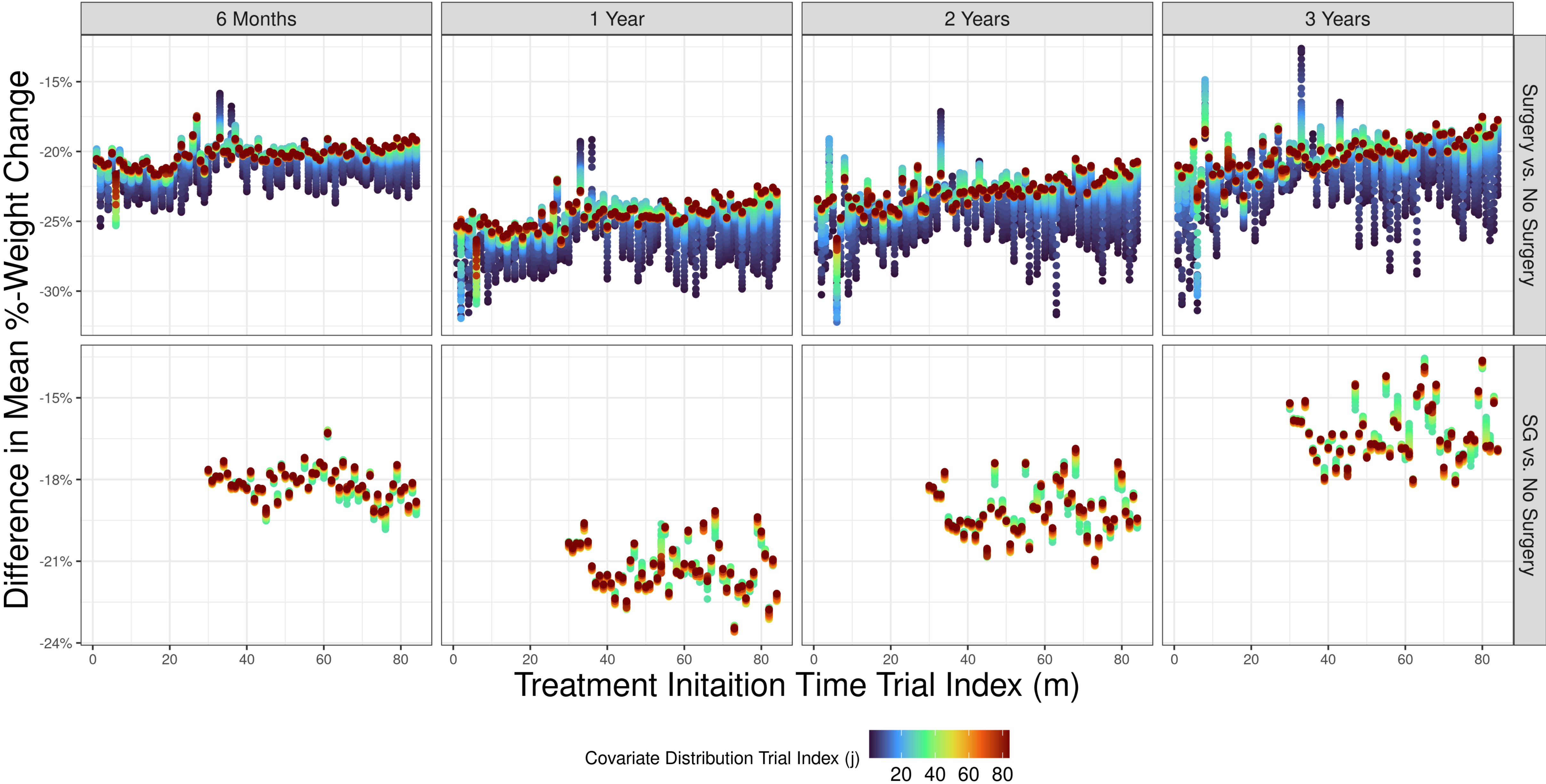


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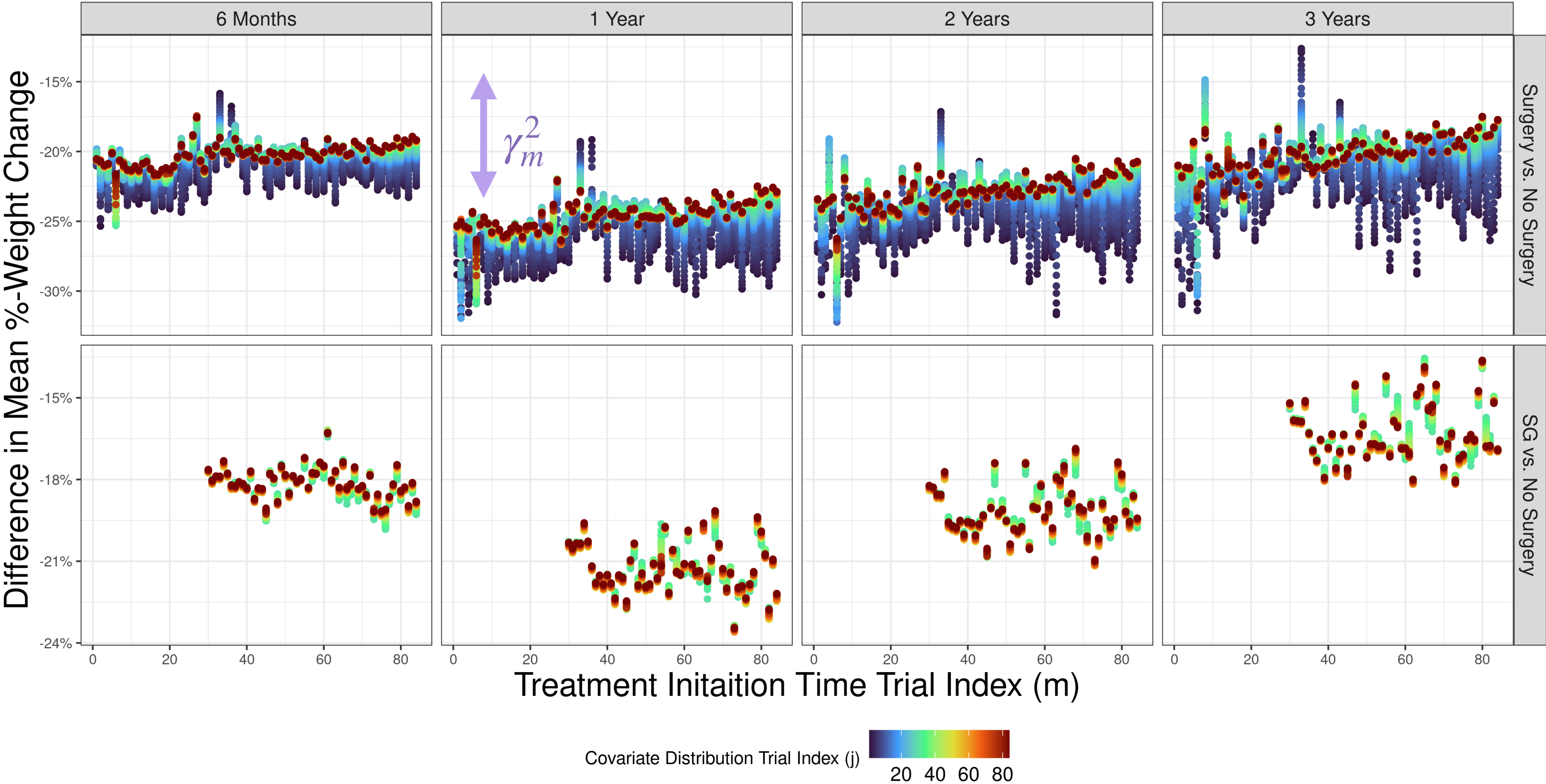
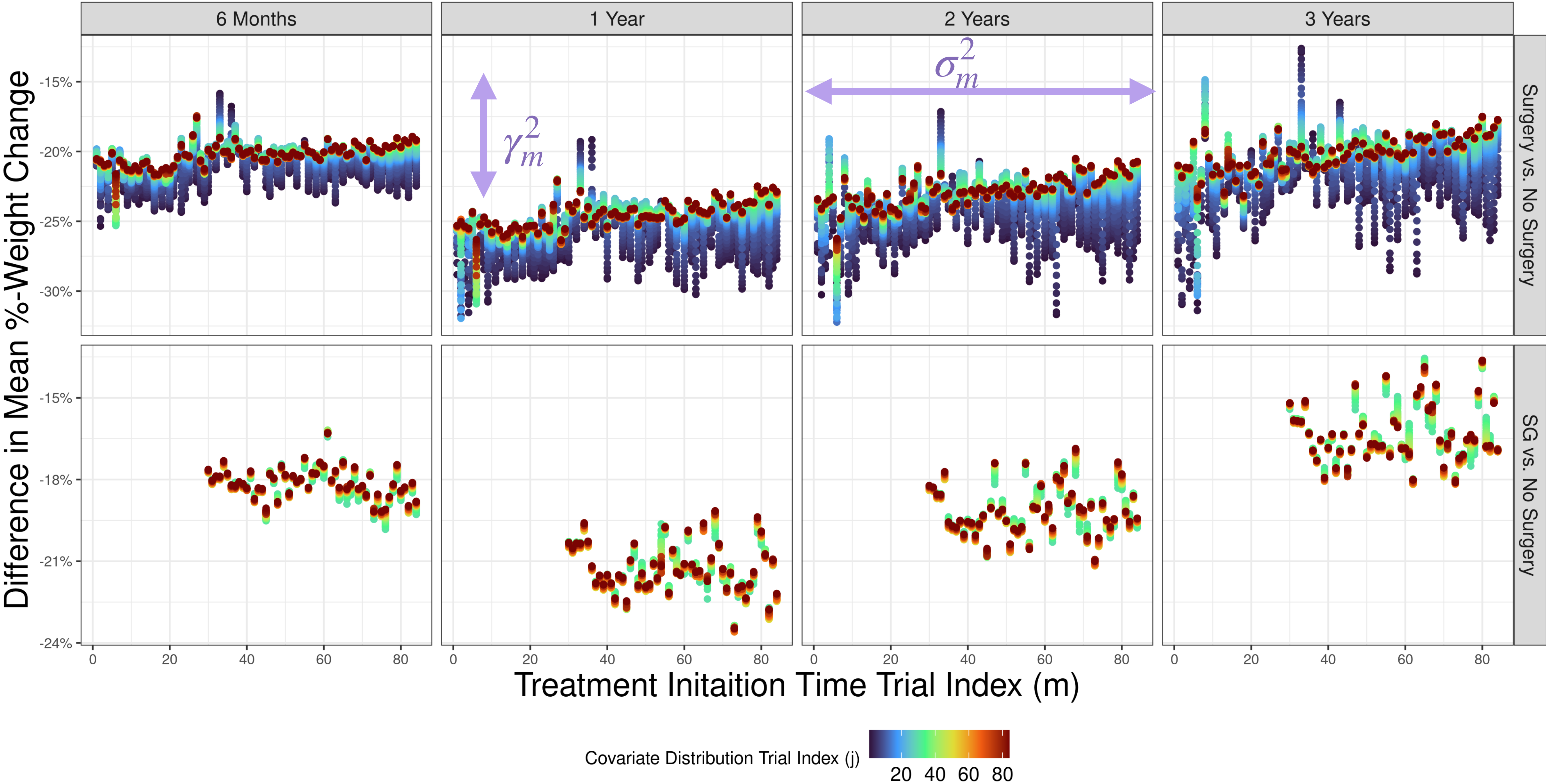


Illustration of $\hat{\chi}_{j,m}$ for Select Comparisons



Estimation of Cross-Trial Effects

- **Cross-Trial Overlap:** $p(\mathcal{L} \mid E_j = 1) > 0 \implies p(\mathcal{L} \mid E_m = 1) > 0$

$$\chi_{j,m}(P) = \mathbb{E}[\mu_m(1, \mathbf{L}_j) - \mu_m(0, \mathbf{L}_j) \mid E_j = 1]$$

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$$\begin{aligned} \dot{\chi}_{j,m}^*(O; P) &= \frac{\mathbf{1}(E_j = 1)}{P(E_j = 1)} \left\{ \mu_m(1, \mathbf{L}_j) - \mu_m(0, \mathbf{L}_j) - \chi_{j,m}(P) \right\} \\ &\quad + \frac{\mathbf{1}(E_m = 1)}{P(E_m = 1)} \xi_{j,m}(\mathbf{L}_m) \left(\frac{A_m}{\pi_m(\mathbf{L}_m)} - \frac{1 - A_m}{1 - \pi_m(\mathbf{L}_m)} \right) \left(Y_m - \mu_m(A_m, \mathbf{L}_m) \right) \end{aligned}$$

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- AIPW residual from eligible patients in trial m
- Transport ratio required to “make trial m distribution look like that of trial j ”

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- Transport ratio required to “make trial m distribution look like that of trial j ”

$$\xi_{j,m}(\ell) = \frac{p(\mathbf{L}_j = \ell \mid E_j = 1)}{p(\mathbf{L}_m = \ell \mid E_m = 1)} = \frac{P(E_m = 1)P(T = j \mid \mathbf{L} = \ell, E = 1)}{P(E_j = 1)P(T = m \mid \mathbf{L} = \ell, E = 1)}$$

Practical Guidance: How and Why Effects Vary over Calendar Time

Projection Model: Constant ($\psi(m, \beta) = \beta$)			
	Fail to Reject $H_0(\theta \leq \delta)$	Fail to Reject H_0 ($\theta \geq 1 - \delta$)	Reject H_0 ($\theta \in (\delta, 1 - \delta)$)
Calendar Time Varying Effect (Study Population)	✗	✗	✗
Calendar Time Varying Effect (Fixed Population)	✗	✓*	✓*
Reason(s) for Variation	No Variation	*Variation not clinically meaningful	*Variation not clinically meaningful
Action	Report common effect	Report common effect	Report common effect
Projection Model: Not Constant ($\psi(m, \beta) \neq \beta$)			
	Fail to Reject $H_0(\theta \leq \delta)$	Fail to Reject H_0 ($\theta \geq 1 - \delta$)	Reject H_0 ($\theta \in (\delta, 1 - \delta)$)
Calendar Time Varying Effect (Study Population)	✓	✓	✓
Calendar Time Varying Effect (Fixed Population)	✗	✓	✓
Reason(s) for Variation	Covariate shift in effect modifiers	Possible changes in treatment efficacy	Covariate shift in effect modifiers, possible changes in treatment efficacy
Action	Acknowledge changes across time driven by changes in underlying populations Consider standardization to fixed population and reporting common effect in that population	Report calendar time-varying effect	Report calendar time-varying effect Consider standardization to fixed population to further study changes in treatment efficacy

Hypothesis Testing for θ

$$H_0 : \theta \in \{0,1\} \text{ vs. } H_1 : \theta \in (0,1)$$

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- **Challenge 1:** Test is at boundary point(s)
 - Asymptotic distribution of θ under some complex ratio of mixtures of chi-squared distributions
 - **Solution 1:** bootstrap

Hypothesis Testing for θ

$$H_0 : \theta \in \{0,1\} \text{ vs. } H_1 : \theta \in (0,1)$$

- **Challenge 1:** Test is at boundary point(s)
 - Asymptotic distribution of θ under some complex ratio of mixtures of chi-squared distributions
 - **Solution 1:** bootstrap
- **Challenge 2:** Can't do a full nonparametric bootstrap
 - Computationally too intensive due to sample splitting, re-fitting pooled models
 - # of predictions required in cross-trial effects is on the order of $M \times |\mathcal{D}|$
 - **Solution 2:** Asymptotic version of parametric bootstrap
 - $S \in \mathbb{R}^{M \times M}$ standardization matrix with entries $\chi_{j,m}(P)$
 - $\sqrt{n}(S - \hat{S}) \xrightarrow{d} \mathcal{N}(0, \Sigma)$ where $\Sigma \in \mathbb{R}^{M^2 \times M^2}$ is the covariance matrix of cross-trial influence function contributions
 - Resample $S^{(1)}, \dots, S^{(B)} \sim \mathcal{N}(\hat{S}, \hat{\Sigma}_n)$ and compute $\hat{\theta}^{(1)}, \dots, \hat{\theta}^{(B)}$ for B bootstrap replicates

Hypothesis Testing for θ

- **Challenge 3:** Bootstrapped values of $\hat{\theta}^{(b)}$ will not be 0, 1 in practice
 - Would always reject H_0
- **Solution 3:** Modify test away from boundary by some small margin δ

$$H_0 : \theta \in [0, \delta] \cup [1 - \delta, 1] \text{ vs. } H_1 : \theta \in (\delta, 1 - \delta)$$

- One can interpret δ to the fraction of variability in entries in $\mathcal{S}$ for which true treatment change is non-negligible in the eyes of the analyst

Why $\hat{\theta} = 1$ May Not Imply Treatment Efficacy is Changing

- Under assumptions/conditions in this work, changes in $\chi_m(P)$ over time:
 - Underlying treatment efficacy is changing
 - Covariate shift in effect modifier
- Under violations of these assumptions, $\hat{\theta} > 0$ even when efficacy is unchanged
 - Including m as a covariate in non-parametric pooled models can be a proxy for unmeasured confounders whose distribution varies across time
 - Difference in effect estimates could reflect **both** difference in treatment efficacy and differential bias introduced by unmeasured confounders
 - Similar w/ model misspecification and residual confounding
- If worried about residual confounding in pooled models influencing estimate of trend
 - Set c parameter in MSM selection procedure to be greater
 - Requires changes in $\hat{\chi}_m$ over time must be greater in order for a non-constant MSM to be selected

$L(\hat{\psi}_k)$ for Candidate MSMs

Comparison	Outcome	Constant	Linear	Cubic	Spline (2 Knots)	Spline (3 Knots)
Surgery vs. No Surgery	6 Months	-3.518 (12.91)	-3.525 (3.47)	-3.527 (0.01)	-3.527 (0.36)	-3.527 (0.00)
	1 Year	-5.251 (23.79)	-5.267 (6.97)	-5.273 (0.37)	-5.273 (0.61)	-5.274 (0.00)
	2 Years	-4.552 (26.05)	-4.573 (4.55)	-4.577 (0.38)	-4.577 (0.00)	-4.576 (0.98)
	3 Years	-3.625 (44.40)	-3.659 (16.01)	-3.675 (2.04)	-3.675 (1.99)	-3.678 (0.00)
RYGB vs. No Surgery	6 Months	-4.222 (1.81)	-4.223 (0.78)	-4.223 (0.00)	-4.223 (0.14)	-4.223 (0.13)
	1 Year	-6.474 (4.71)	-6.476 (3.10)	-6.478 (0.75)	-6.479 (0.32)	-6.479 (0.00)
	2 Years	-5.791 (7.84)	-5.792 (6.60)	-5.794 (5.32)	-5.796 (3.87)	-5.800 (0.00)
	3 Years	-4.645 (10.70)	-4.647 (8.71)	-4.655 (3.14)	-4.656 (2.07)	-4.659 (0.00)
SG vs. No Surgery	6 Months	-1.813 (0.42)	-1.813 (0.56)	-1.814 (0.30)	-1.814 (0.29)	-1.814 (0.00)
	1 Year	-2.490 (0.00)	-2.486 (6.16)	-2.486 (5.88)	-2.486 (6.08)	-2.486 (5.39)
	2 Years	-2.020 (0.01)	-2.020 (0.00)	-2.020 (0.41)	-2.020 (0.37)	-2.020 (0.05)
	3 Years	-1.466 (0.00)	-1.463 (3.58)	-1.464 (2.41)	-1.464 (2.57)	-1.465 (1.92)
RYGB vs. SG	6 Months	-0.101 (0.47)	-0.101 (0.27)	-0.101 (0.08)	-0.101 (0.09)	-0.102 (0.00)
	1 Year	-0.220 (0.00)	-0.217 (2.75)	-0.217 (2.87)	-0.217 (2.71)	-0.217 (2.53)
	2 Years	-0.288 (0.00)	-0.283 (3.73)	-0.281 (6.00)	-0.281 (6.08)	-0.281 (5.87)
	3 Years	-0.298 (0.00)	-0.297 (1.00)	-0.296 (1.87)	-0.296 (1.85)	-0.297 (1.16)

Table S1: Values of loss function $L(\hat{\psi}_k)$ for each candidate MSM. Cells in bold denote the MSM $\hat{\psi}^*$ which minimizes $L(\hat{\psi}_k)$ in each setting. Values in parenthesis denote the number of weighted standard deviations, ϵ , by which $L(\hat{\psi}_k)$ exceeds $L(\hat{\psi}^*)$.