

SCI ANNUAL MEETING, 2026
SYMPOSIUM SESSION

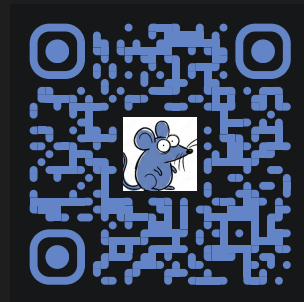
CATE Estimation in Complex Longitudinal Data

An Introduction

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🌐 <https://ainaimi.github.io/>



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Acknowledgements

- This (in progress) work benefitted from discussion with several colleagues:
 - Jennifer Kawwass, Austin Schirmer, and Heather Hipp at Emory Reproductive Center
 - Audrey Gaskins at Emory Epidemiology
 - Edward Kennedy at CMU
- Please feel free to report errors and/or send comments to me:
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Overview

An Applied Problem

A Starting Point with G-Estimation of SNMMs

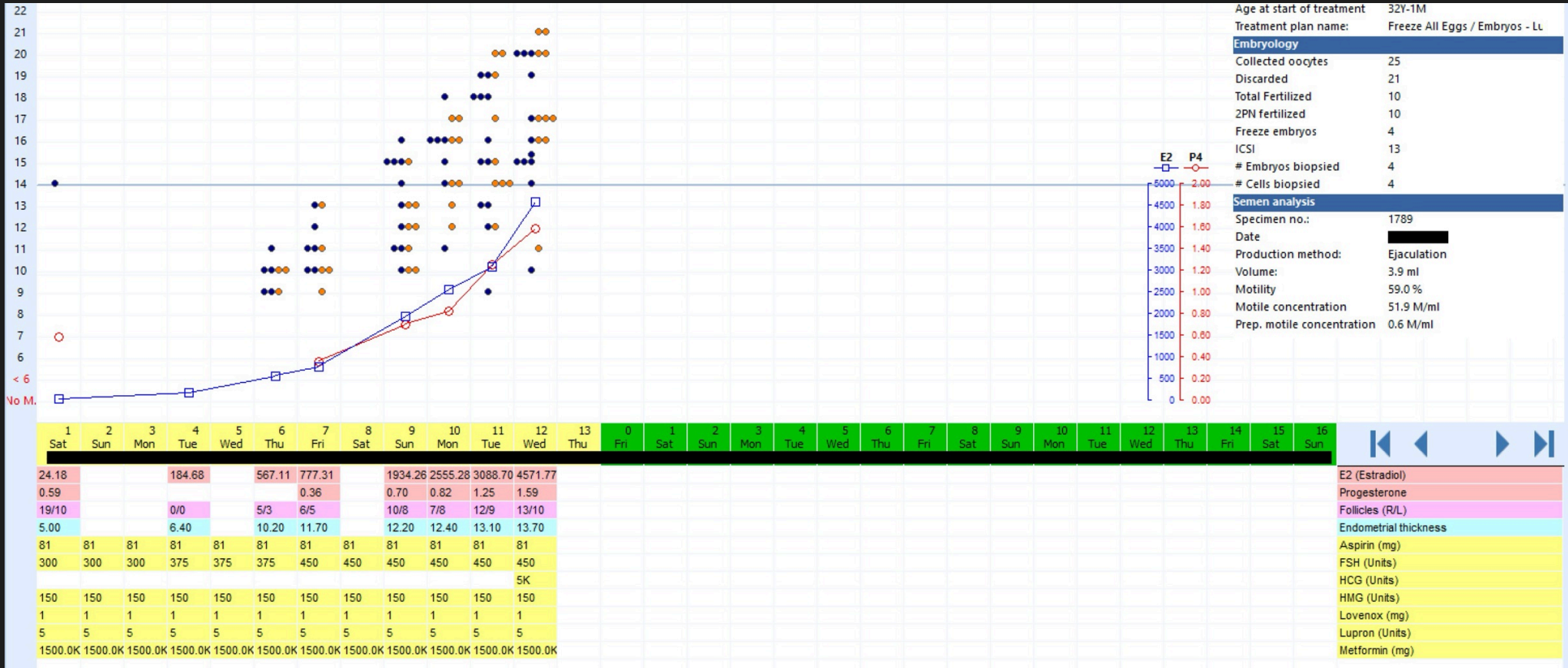
Three Approaches

Some Questions

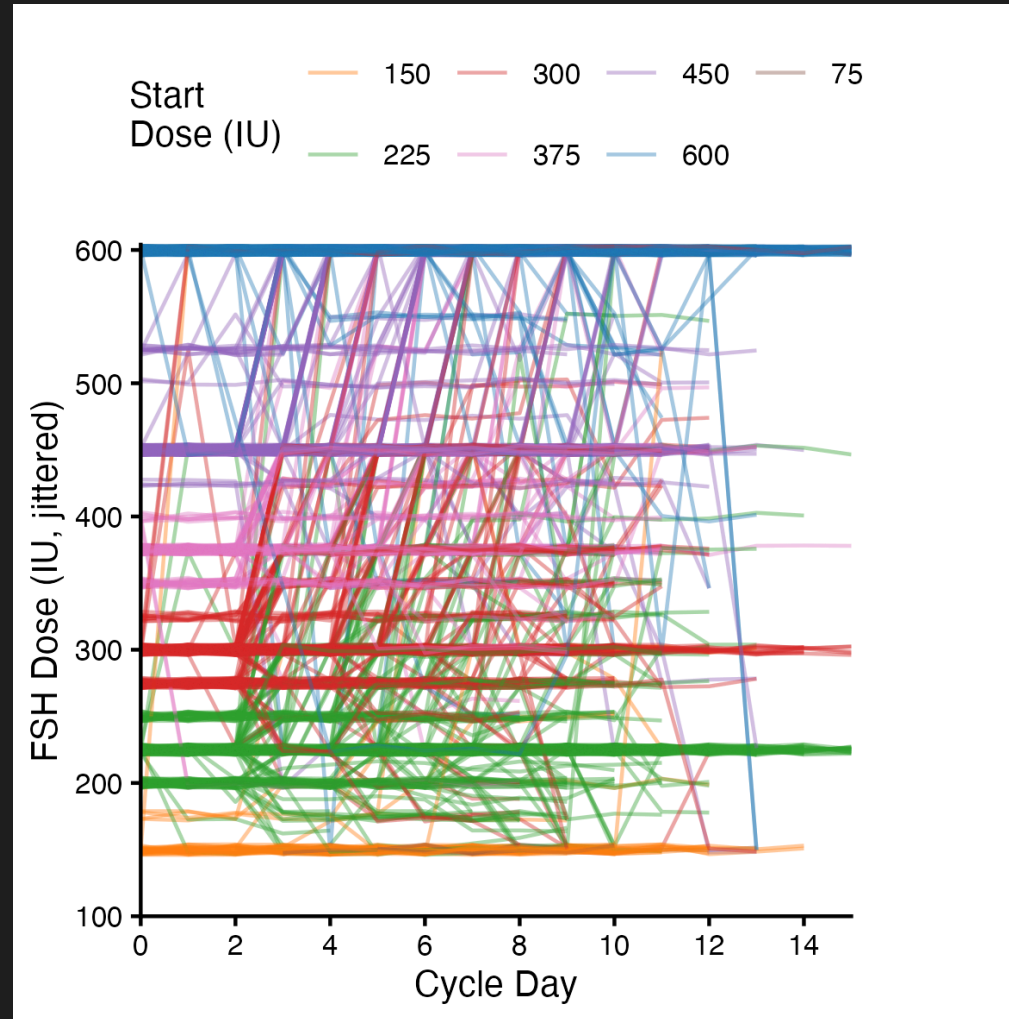
IVF: A Repeated, High-Stakes Treatment Decision

- Infertility affects roughly 15% of couples worldwide
- IVF: Daily gonadotropin injections to stimulate follicle growth + Trigger Shot
- Clinicians make dose decisions roughly every **2 days** based on:
 - Hormone Level Measures (e.g., Estradiol, Progesterone)
 - Follicle count and size (right + left ovary)
 - Baseline and Prior Daily Doses
 - Other
- But, in the US, fewer than 40% of IVF cycles result in live birth.
- Failure is expensive for the patient, at \$15K to \$30K per cycle

A Patient Example



Treatment Variation



What Do We Want to Know?

- Person-Day Specific Effects:

For a specific patient with a dose and covariate history H through day k , what would happen to their outcome(s) if we changed the dose on day k alone?

- Person Specific Optimal Regimes:

For a specific patient with a specific dose and covariate history H through day k , what changes to their dose on day k would optimize the outcome(s) of interest?

- Clinic Specific Policies:

For {new patients, returning patients with unexpected prior failures} can/should there be clinic specific time-varying dose protocols that optimize outcomes?

How Can We Answer These Questions?

Robins (1994) introduced the **blip function** encoded in a SNM, which formalizes the person-day CATE

For example, if we let:

- $Y^{\bar{a}_k, \underline{a}_{k+1}^*}$ represent the end-of-cycle oocyte retrieval count under
 - a intervention history up to day k , denoted \bar{a}_k
 - a referent dose future from day $k + 1$, denoted \underline{a}_{k+1}^*

$$\gamma_k(h_k; \psi) = E\left[Y^{\bar{a}_k, \underline{a}_{k+1}^*} - Y^{\bar{a}_{k-1}, \underline{a}_k^*} \mid H_k = h_k \right]$$

where $h_k = (\bar{Z}_k = \bar{z}_k, \bar{A}_{k-1} = \bar{a}_{k-1})$ is the full patient history through day k .

For example, on the third ($k = 2$) day of COS, we can specify a linear SNMM for a "blip" increment as:

$$\gamma_2(h_2; \psi) = (\psi_1 + \psi_2 z_0 + \psi_3 z_1 + \psi_4 z_2 + \psi_5 a_1 + \psi_6 a_0)(a_2 - a_2^*)$$

(but...)

G-Estimation

Robins also introduced g estimation, a moment based estimating function for ψ .

Find $\hat{\psi}$ such that the "blip-free" outcome

$$Y^{\text{free}}(\psi) = Y - \sum_k \gamma_k(H_k; \psi)$$

is mean independent of treatment residual ($A_k - E[A_k | H_k]$), given history.

At each k , solve:

$$\sum_i [d^*(A_{ik}, H_{ik}) - \hat{E}\{d^*(A_k, H_{ik}) | H_{ik}\}] [Y_i^{\text{free}}(\psi) - \hat{E}\{Y_i^{\text{free}}(\psi) | H_{ik}\}] = \mathbf{0}$$

where $d^*(A_{ik}, H_{ik})$ is a p -dimensional function of current treatment and history.

Problem 1: It's Complicated

For decades, authors (e.g., Stijn Vansteelandt) have been advocating for more use in applied settings, but the concepts are genuinely demanding:

- SNMs and blip functions are very unfamiliar objects
- The construction of Y^{free} , and the choice of d^* require understanding before implementation
- Estimating equations are not standard for many applied researchers

Software situation remains poor:

- `gesttools` in R: no longer on CRAN and appears non-operational
- To my knowledge, no well-maintained, general-purpose package exists in R or Python
- Practitioners who want to use these methods largely have to code from scratch

Problem 2: Fully Conditional Structural Model

G-estimation requires **correct specification of the SNMM**:

- Must correctly model covariates that modify the effect
- The "double robustness" property is not like AIPW or TMLE of the ATE.
- There is some interesting work on interpreting ψ in misspecified SMMs, but not clear:
 - How this generalizes to time-varying settings
 - How to convey these "projection" parameter interpretations to clinical colleagues

What We'll Discuss Today

DR Learner (Kennedy): a broadly applicable "Swiss army knife"

- Doubly-robust pseudo-outcome regression, extended to the longitudinal setting
- No structural model required; any ML method in the second stage

Extension of the R-Learner (Syrgkanis): closely related to g-estimation

- Retains the Neyman-orthogonal estimating equation logic of SNMMs
- Incorporates ML flexibility directly into the structural model for the blip

Causal Excursion Effects (Chen): in micro-randomized trials

- Known randomization probabilities eliminate time-varying confounding
- Weighted estimating equation approach yields direct inference on day-specific CATEs

Panel Discussion (Scharfstein): tying it all together

- Lead the panel discussion on theoretical and practical challenges

5:00
App Store

Home Details

Invited Session

CATE Estimation in Complex Longitudinal Data

Wed, May 13
8:30 AM - 10:00 AM

Location: Salon G-J

50 Attending
50 New

0 Session Polls

2 Likes
2 New


0 Comments


0 Questions Asked

Overview

Session Co-Chair: Ashley Naimi, Emory University Edward Kennedy, Carnegie Mellon University Methods for conditional average...
[See more](#)

Speaker

 Ashley Naimi
Professor
Emory University

 Edward Kennedy

Some Questions

How Hard Is This to Implement?

DR Learner: very accessible

- Two-stage regression; second stage uses any ML or parametric model
- We often implement in time-fixed settings with the Cross-Validated Super Learner
- Same ease in longitudinal extensions?

DML with G-Estimation: moderate complexity

- Very closely related to g Estimation proper
- Can it be implemented via a residual-on-residual type approach under a nonparametric structural model?

CEE / MRT methods: designed for accessibility

- Weighted GEE-type estimating equations
- R package `MRTAnalysis` available
- How do these translate to observational data?

Robustness: What Can Go Wrong?

"Double robustness" can mean different things in different settings. How "robust" is each approach?

What happens if:

- One or Both models misspecified?
- Near positivity violations?
- How "robust" is variance estimation
- Can we even estimate the variance of the CATE functional?

But also:

How amenable are these to:

- measurement error correction methods?
- sensitivity / bias analyses for unmeasured confounding?
- Nonparametric bounding of LCATE effects?

Machine Learning for Nuisance Functions?

When fitting nuisance functions, can ML/nonparametric methods be used?

- What happens if we use very slowly convergent models (random forests?)
- Does longitudinal data affect how we might do cross-fitting?
- Does cross-fitting matter more, less, or the same for longitudinal CATEs?

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