



# Application of Causal Inference Methods to Improve Treatment of HIV in Resource-Limited Settings

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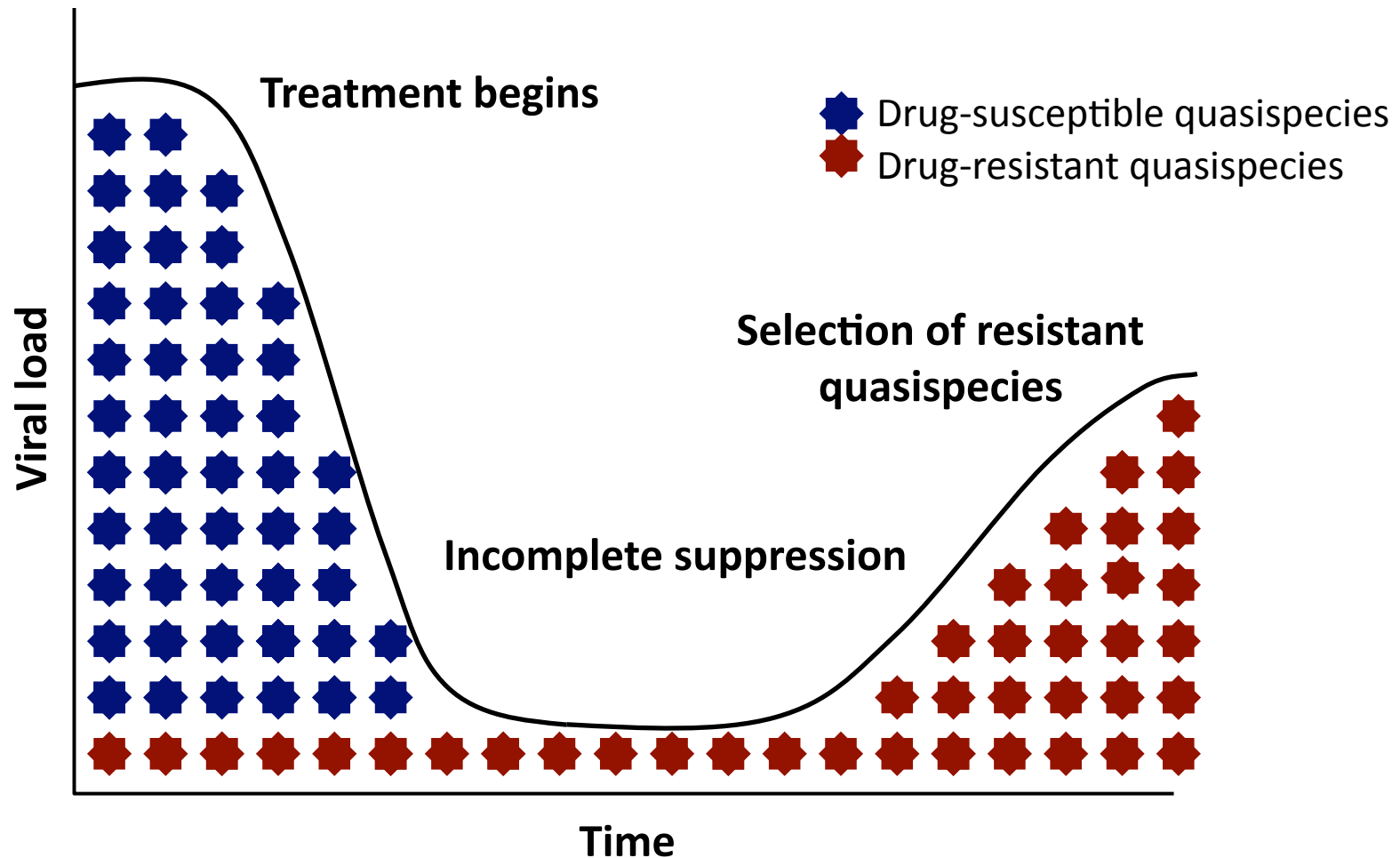
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# Outline

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1. What types of questions can causal inference methods help us to answer?
  - Example: Strategies for detecting and managing first line failure
2. When do standard analysis methods break down?
  - Time-dependent confounding of longitudinal treatments
3. What estimation tools are available?
  - Inverse Probability Weighting
  - Longitudinal G- computation
  - New methods coming...

# HIV frequently develops resistance to antiretroviral drugs



# Detecting and responding to virological failure

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- Once virological failure develops, change regimen immediately to prevent additional resistance and disease progression
- > Standard of care: Regular viral load monitoring
  - Detect failure and initiate change in regimen
- Viral loads are expensive and require significant infrastructure
  - Burden of HIV greatest in resource-limited settings where viral loads often not available

# When to switch? WHO 2010 Recommendations

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<b>Virological</b>	Plasma viral load above 5000 copies/ml (confirmation recommended)
<b>Immunological</b>	<ul style="list-style-type: none"><li>• Fall of CD4 count to pre-therapy baseline (or below); or</li><li>• 50% fall from the on-treatment peak value; or</li><li>• Persistent CD4 levels below 100 cells/mm</li></ul>
<b>Clinical</b>	New or recurrent WHO Stage 4 condition (not IRIS; +/- some Stage 3)

# Impact of Alternative WHO Failure Criteria?

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- Many studies: Immunologic criteria have poor sensitivity for detecting virological failure
  - Ex: Keiser (2009): Sensitivity 17% (7,32)
- ART-LINC (2009): Higher switch rates and switch at higher CD4 counts in programs that have VL monitoring
- **Use of CD4 vs. VL-based criteria will result in delayed switch for many**

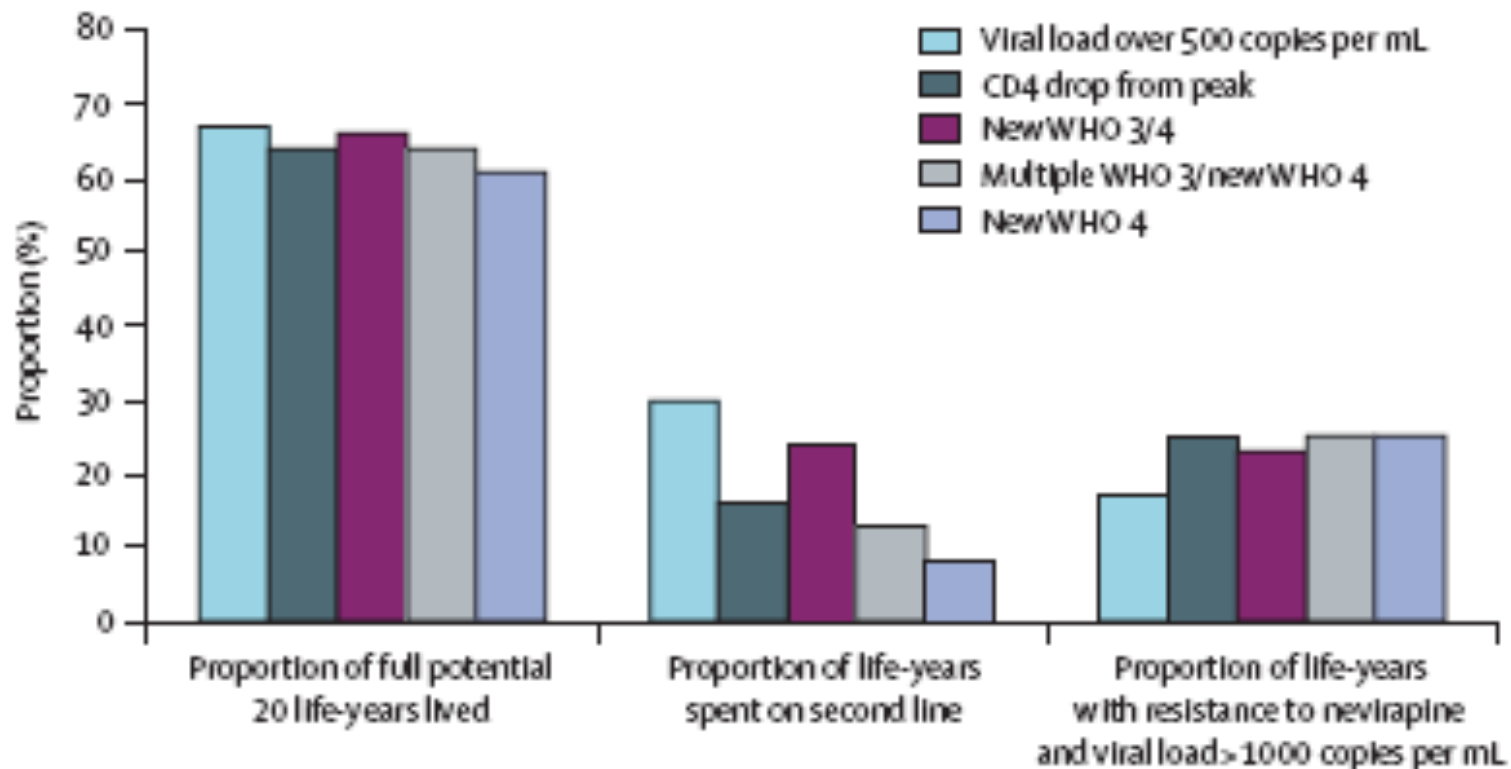
# Impact of Alternative WHO Failure Criteria?

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- Systematic review
  - Chang et al (Cochrane 2010)
- HBAC (Tororo, Uganda): RCT comparing clinical/CD4/VL
  - Published in abstract only (CROI 2008)...
- Three ongoing RCTs...
  - Zambia (Saag): routine vs. targeted VL
  - Thailand (Lallemant): CD4 vs. VL
  - Cameroon (Laurent): CD4/VL vs. clinical
  - Multi-country (PENPACT1): different VL thresholds in kids

# Mathematical model of alternative monitoring strategies in Africa

- Monitoring based on CD4 instead of viral load results in longer delay until treatment modification, but little long-term mortality difference



Phillips et. al. *Lancet* 2008, 20 year outcomes



# Impact of alternative monitoring strategies

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- Will delayed regimen modification after loss of viral suppression increase mortality? How much?
- Will use of CD4 (or clinical) rather than VL-based switching criteria increase mortality? How much?
- Which CD4 criteria will give the best patient outcomes?

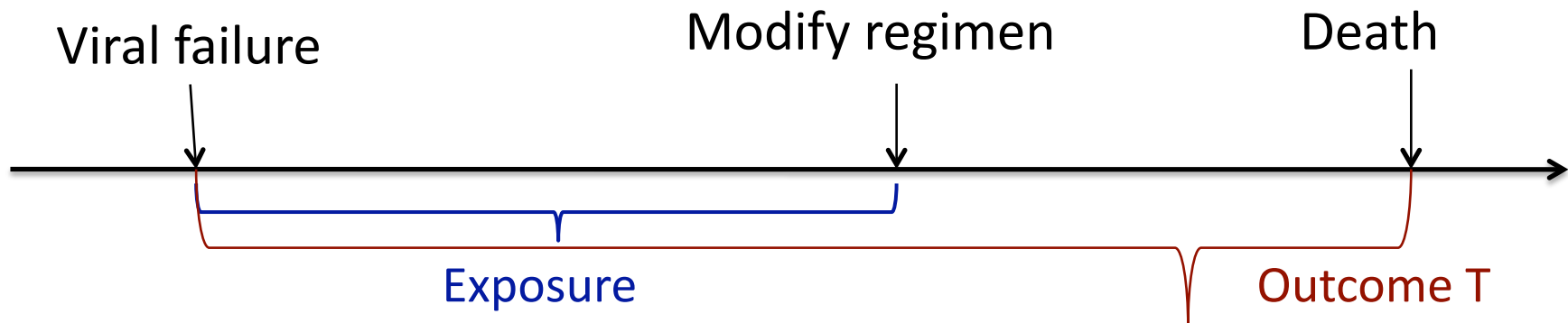
# Hypothetical Randomized Trial

- Will delayed regimen modification following viral failure increase long-term mortality?

Subjects enrolled at time of viral failure

Subjects remain on failing therapy for a randomly assigned duration

Follow until death



# Counterfactuals

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- $T_{\text{switch}}$  : counterfactual survival time under a specified switch time
  - Example:  $T_0$  = an individual's survival if she were to have switched immediately
- Ideally, observe the survival time of everyone in the population under each possible delay time
  - Compare survival under different delay times
- Example:  $E(T_{\text{switch at 0 mo}} - T_{\text{switch at 9 mo}})$ 
  - Difference in mean survival time if everyone in population had switched immediately versus after a 9 month delay

# Marginal Structural Models

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- Model on the counterfactual outcome
  - Ex. Model Discrete Hazard

$$P(T_{switch} = t | T_{switch} \geq t) = m(t, switch | \beta)$$

- One possible model:

$$\text{logit}(m(t, switch | \beta)) =$$

$$\beta_0 + \beta_1 \min(t, switch) + \beta_2 t + \beta_3 \min(t, switch) \times t$$

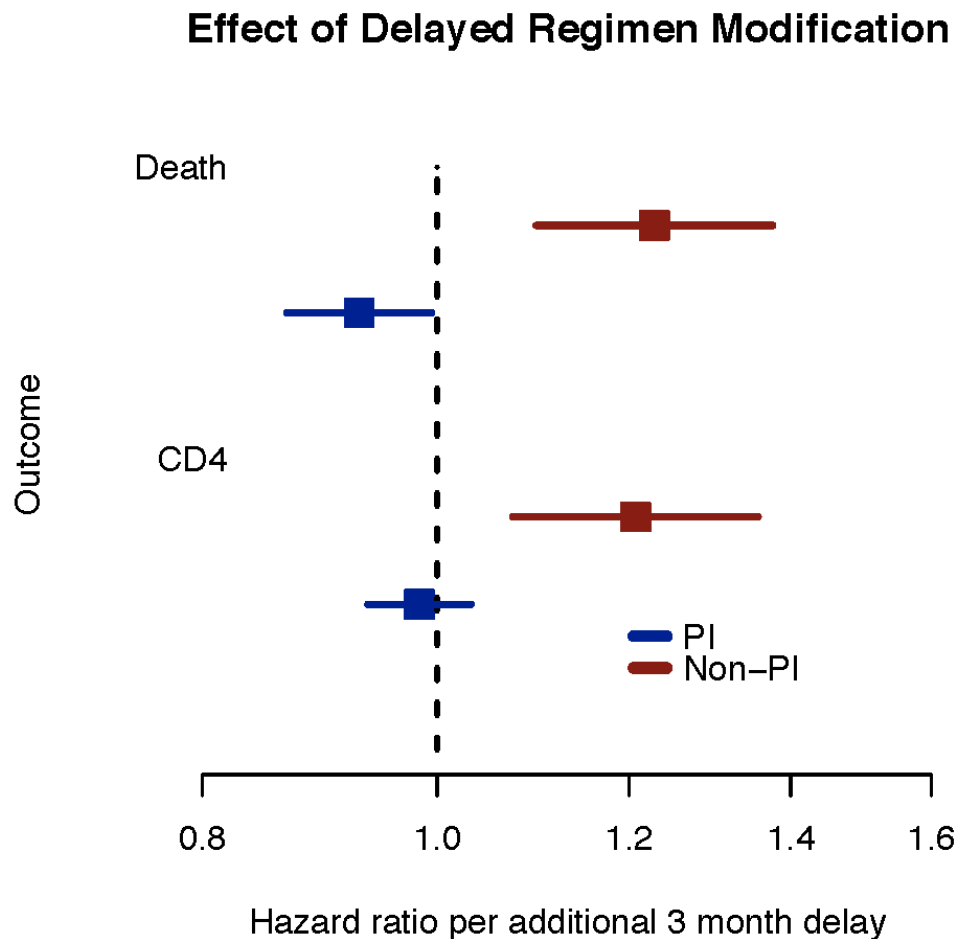
- For a given time  $t$ , linear summary (on log odds scale) of the cumulative effect of time on failing therapy on the discrete hazard of death at  $t$

# Prospective clinical cohort data

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- 2 cohorts of HIV-infected patients
  - Johns Hopkins ; Univ. North Carolina, Chapel Hill
- Eligible when experience virologic failure on HAART ( $t=0$ )
  - 982 subjects, 3414 person-years follow-up, 93 deaths
  - 742 failed a PI-based regimen; 240 fail a non-PI based regimen
- Primary outcome: All-cause mortality
  - Secondary outcome: immunological failure
- Exposure: Delay from virological failure to switch
- Covariates: Demographics, risk group, CD4/CD8, Viral Load, AIDS-related DX, ARV history

# Delayed modification following failure of a HAART regimen without a protease inhibitor increases mortality



# How much does this tell us about the importance of viral load monitoring in Africa?

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- Subjects from North America
  - Different health care systems, patient populations....
- A historical cohort, not representative of current first line failures
  - Included failures between 1996 and 2006
  - Roughly 50% of sample exposed to ARVs prior to 1996
    - Not powered to exclude these individuals
- Does not directly address the question: What impact will lack of viral load monitoring have on mortality?

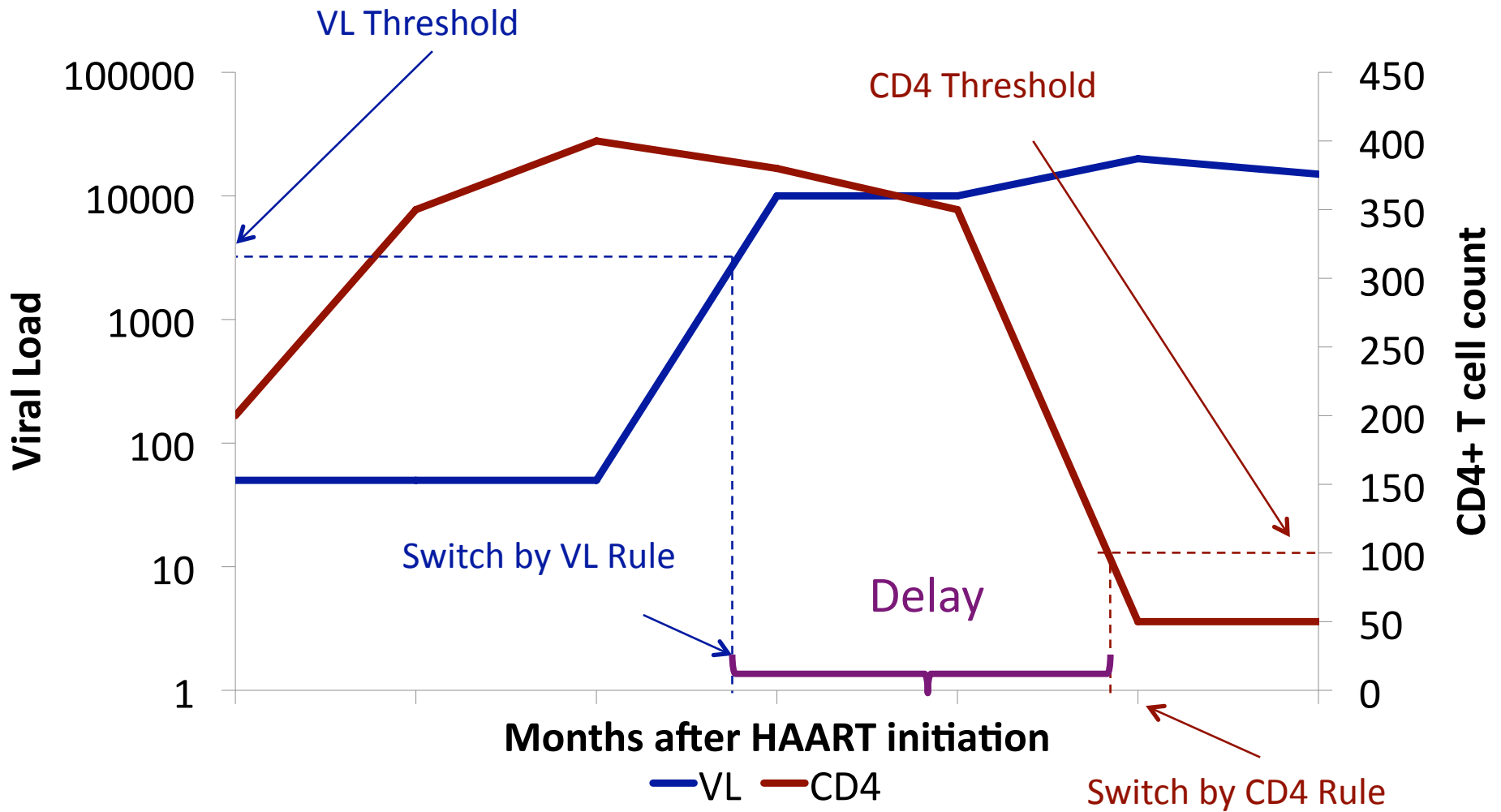
# Refining the Question

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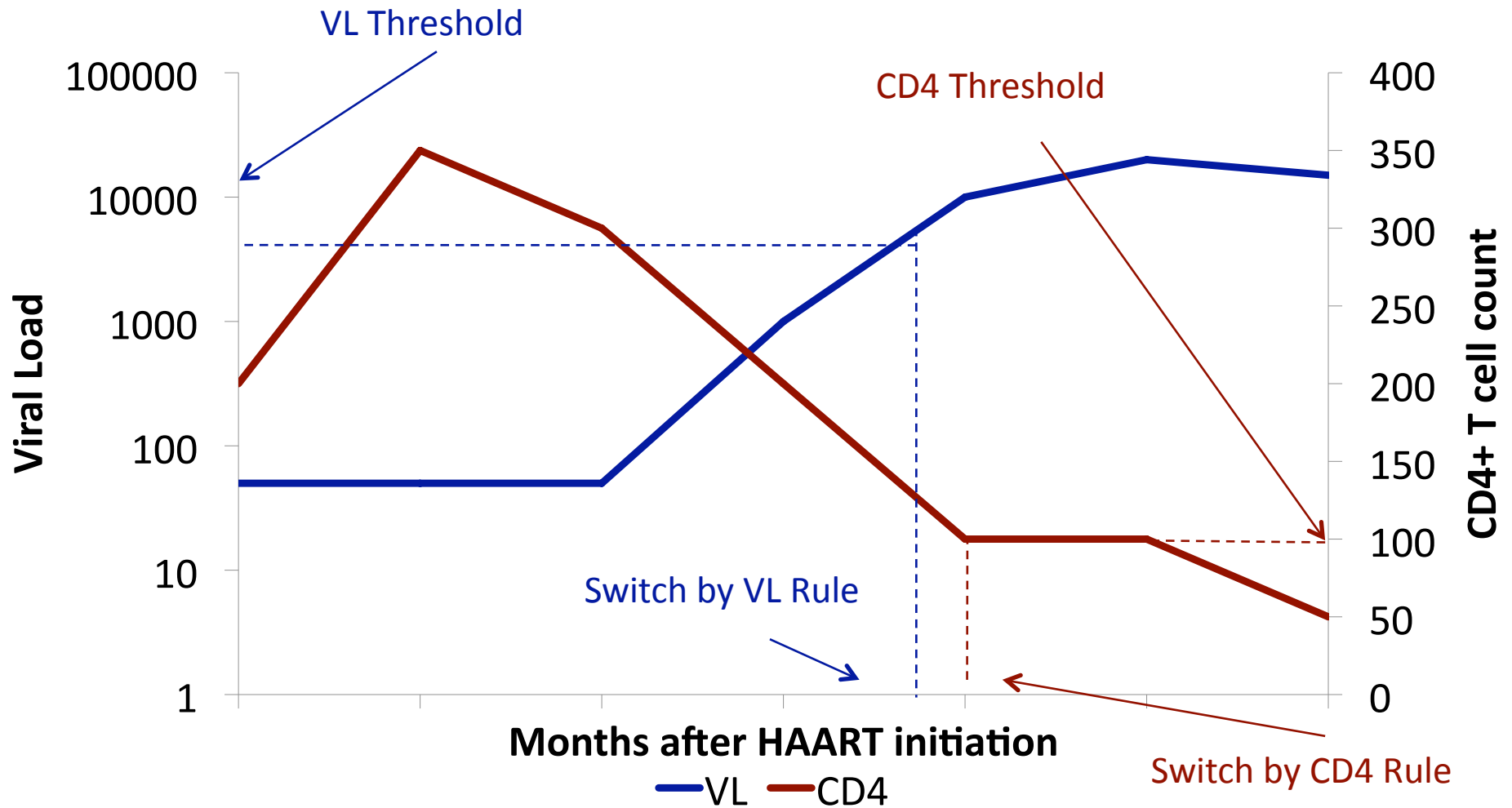
- Say we find that delayed modification after viral failure does increase mortality
- Substantial data suggest that use of CD4 counts rather viral loads will increase the average delay time between viral failure and switch
- Does this imply that use of a CD4-based switching strategy will increase mortality?



# Subject #1: Delayed switch after failure; CD4 elevated for much of the delay time



# Subject #2: Minimally delayed switch after failure



# Refining the Question

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- Say we find that delayed modification after viral failure does increase mortality
- Substantial data suggest that use of CD4 counts rather viral loads will increase the average delay time between viral failure and switch
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## Not necessarily....

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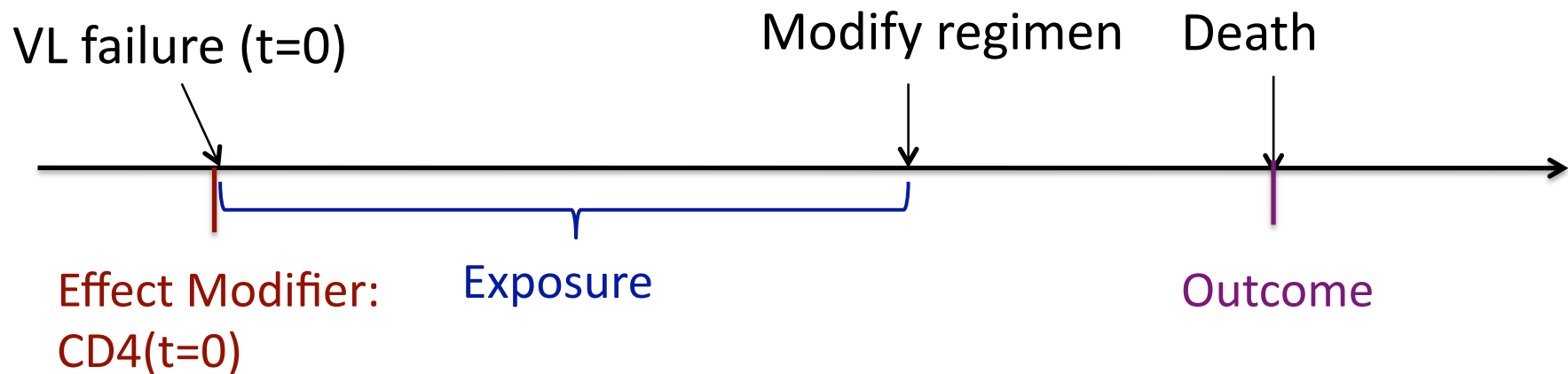
- Subject 1 meets CD4 failure criteria months after virologically failing
  - Substantial delay in modification if CD4 versus viral load strategy is used for patients like this
- Subject 2 meets CD4 failure criteria shortly after virologically failing
  - Minimal delay in modification if CD4 versus viral load strategy is used for patients like this
- *Is delay harmful for subject 2, or just for subject 1?*

# Defining the Target Parameter in Terms of a Series of Hypothetical Randomized Trials

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- Hypothesis: The effect of delayed regimen modification differs depending on a subject's current CD4 count

At  $t=0$ , randomly assign delay conditional on (within strata of) current CD4 count

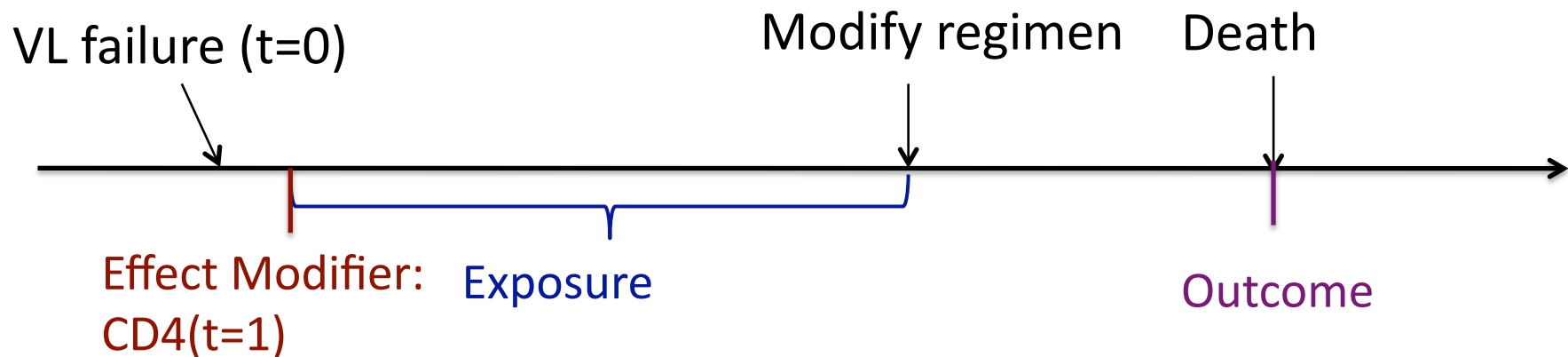


# Defining the Target Parameter in Terms of a Series of Hypothetical Randomized Trials

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- Hypothesis: The effect of delayed regimen modification differs depending on a subject's current CD4 count

At  $t=1$ , randomly assign delay time to subjects who have not yet switched conditional on current CD4 count

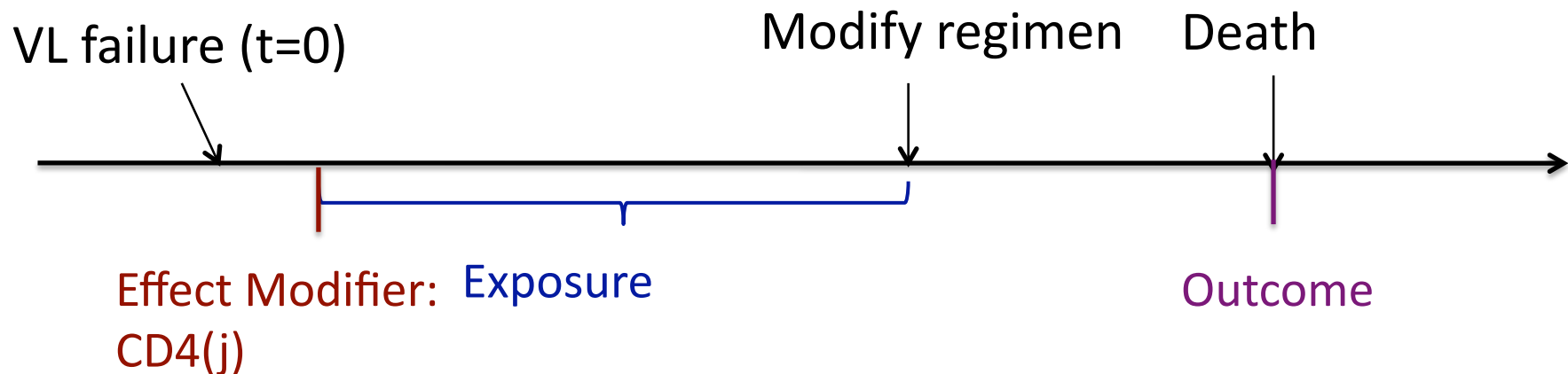


# Defining the Target Parameter in Terms of a Series of Hypothetical Randomized Trials

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- Series of trials, beginning at successive times  $j$ 
  - Effect of additional delay until regimen modification on future outcome
  - Modification of this effect by CD4 count at time  $j$

At time  $j$ , randomly assign delay time to subjects who have not yet switched conditional current CD4 count



# History-Adjusted Marginal Structural Models (HA-MSM)

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- Generalization of standard MSM
  - Different parameter of interest
    - Indexed by interventions beginning at different time points
  - HA-MSM assume a standard MSM beginning at each time point
    - For counterfactuals indexed by treatment after time  $j$
    - Conditional on some subset of the observed history up till time  $j$
- HA-MSM allow us to assume common parameters across time points



## Defining the Target Parameter Using a History-Adjusted Marginal Structural Model

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- Ex: Model the counterfactual probability of survival at least 3 years among subjects who have not yet switched therapy as a function of future switch time and current CD4 count

$$\begin{aligned} E(Y_{Orig(j)=1,switch}(j+3) | Orig(j) = 1, CD4(j)) = \\ \beta_0 + \beta_1 \min(j+3, switch) + \beta_2 CD4(j) \\ + \beta_3 \min(j+3, switch) \times CD4(j) \end{aligned}$$

# Defining the Target Parameter Using a History-Adjusted Marginal Structural Model

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- Allows us to test the hypotheses
  1. Effect of delayed regimen modification following virological failure differs depending on a subject's CD4 count during failure
  2. Delayed regimen modification remains harmful for subjects who maintain CD4 counts about the WHO switching threshold
    - These are the subjects who will actually be subjected to delays
    - Evidence for importance of VL monitoring

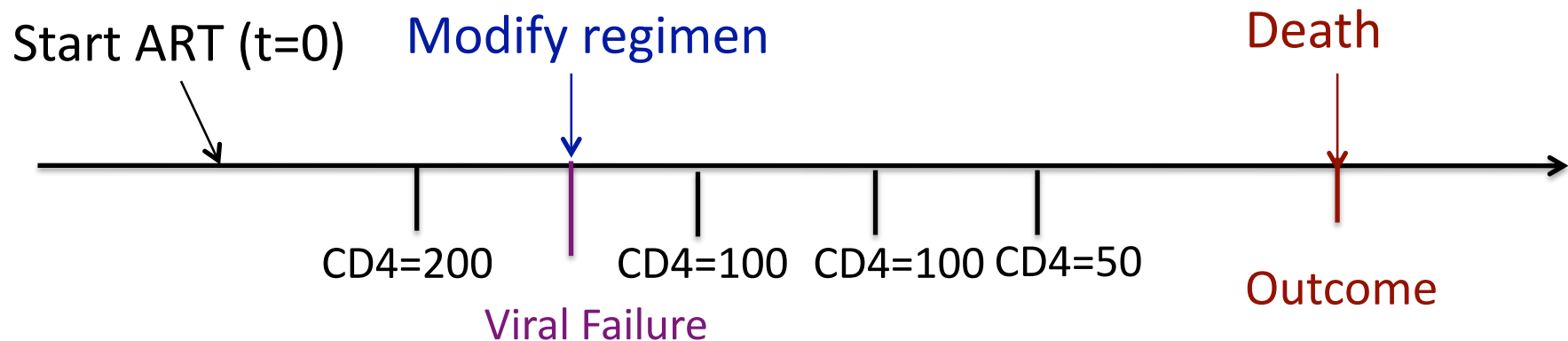
# Impact of alternative monitoring strategies

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- Will delayed regimen modification after loss of viral suppression increase mortality? How much?
- Will use of CD4 (or clinical) rather than VL-based switching criteria increase mortality? How much?
- Which CD4 criteria will give the best patient outcomes?

# Hypothetical Randomized Trial

- At time of ART initiation, randomly assign patients to VL versus CD4 monitoring
  - Rule=1 (assigned to VL arm)
- Stay on first line regimen until meet failure criteria for your assigned strategy
- Measure Outcome

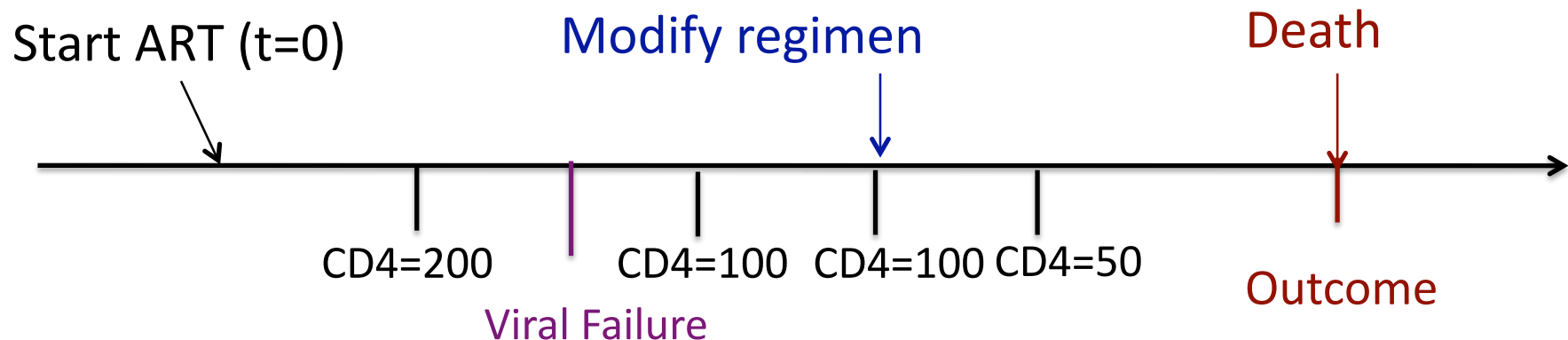


Exposure=What strategy are you assigned to? (“Rule”)

Ex: Rule=1 (assigned to VL arm)

# Hypothetical Randomized Trial

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Exposure=What strategy are you assigned to? (“Rule”)

Ex: Rule=0 (assigned to CD4 arm)

# Dynamic versus static regimes

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- A different type of intervention
- Rather than assigning a fixed treatment, assign treatment according to a strategy or rule
  - What treatment a given subject gets depends on that subject's covariate values
  - Ex. VL trajectory, CD4 trajectory...
- This is an example of a **dynamic regime** question
  - Different counterfactuals
  - Different estimation methods

# Dynamic Counterfactuals

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- $T_{\text{rule}}$  : counterfactual survival time under specific rule for deciding when to switch
  - Ex.  $T_{\text{rule}=1}$  = a subject's survival time if he had followed a VL-based switching rule
  - Ideally, compare survival distribution under VL versus CD4 switching rules
- Example:  $E(T_{\text{rule}=1} - T_{\text{rule}=0})$ 
  - Difference in mean survival time if whole population had followed a VL-based versus CD4-based switching strategy

# Dynamic Marginal Structural Models

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- Model on counterfactuals indexed by dynamic rules
  - Ex: Model for Discrete Hazard

$$P(T_{rule} = t | T_{rule} \geq t) = m(t, rule | \beta)$$

- One possible model:

$$\text{logit}(m(t, rule | \beta)) =$$

$$\beta_0 + \beta_1 rule + \beta_2 t + \beta_3 rule \times t$$



# Impact of alternative monitoring strategies

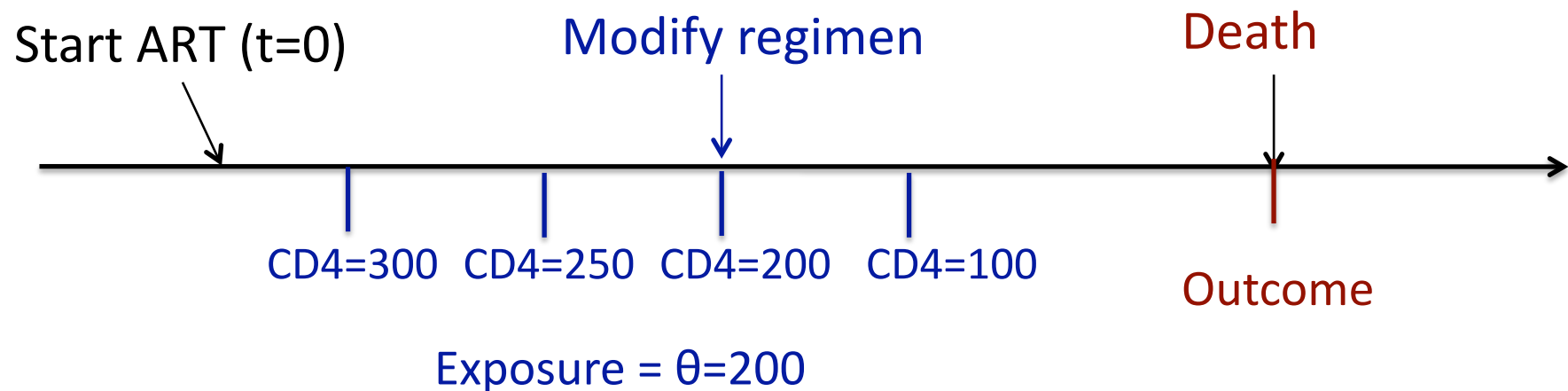
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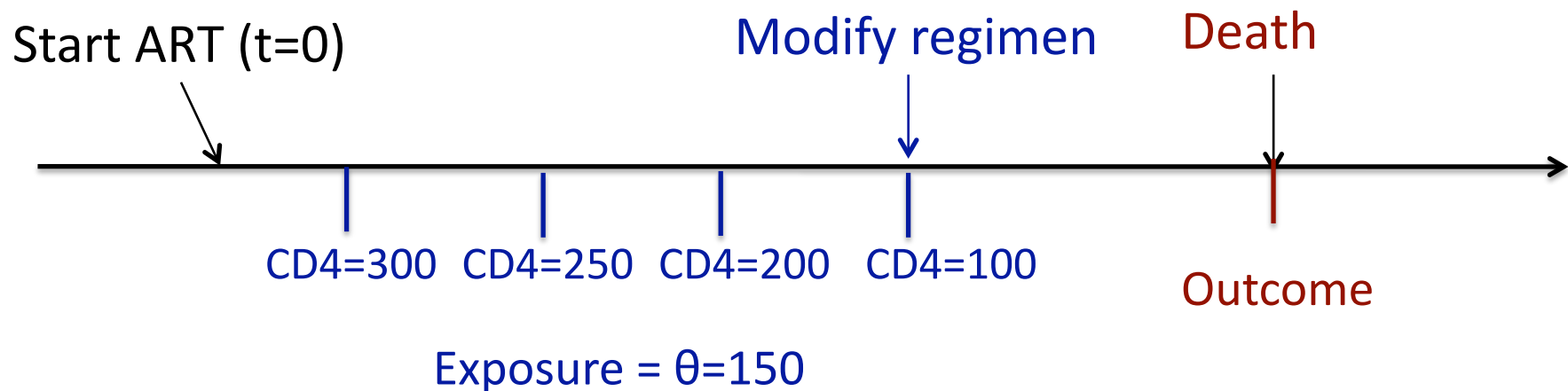
- At time of ART initiation, randomly assign CD4 modification threshold  $\theta$
- Stay on first line regimen until CD4 count meets this criteria, then modify immediately (or within some allowed window)
- Measure Outcome



# Hypothetical Randomized Trial

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- Stay on first line regimen until CD4 count meets this criteria, then modify immediately (or within some allowed window)
- Measure Outcome



## Dynamic marginal structural models (2)

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- What is the “best” threshold?
- Ex. Model the probability of survival for at least 7 years as a function of modification threshold  $\theta$  and baseline CD4 count

$$P(T_{rule_\theta} \geq 7) = m(\theta|\beta)$$

- One possible model:

$$\text{logit}(m(t, \theta|\beta)) = \beta_0 + \beta_1\theta + \beta_2\theta^2$$

- Solve for  $\theta_{opt}$  that maximizes probability of surviving at least 7 years....

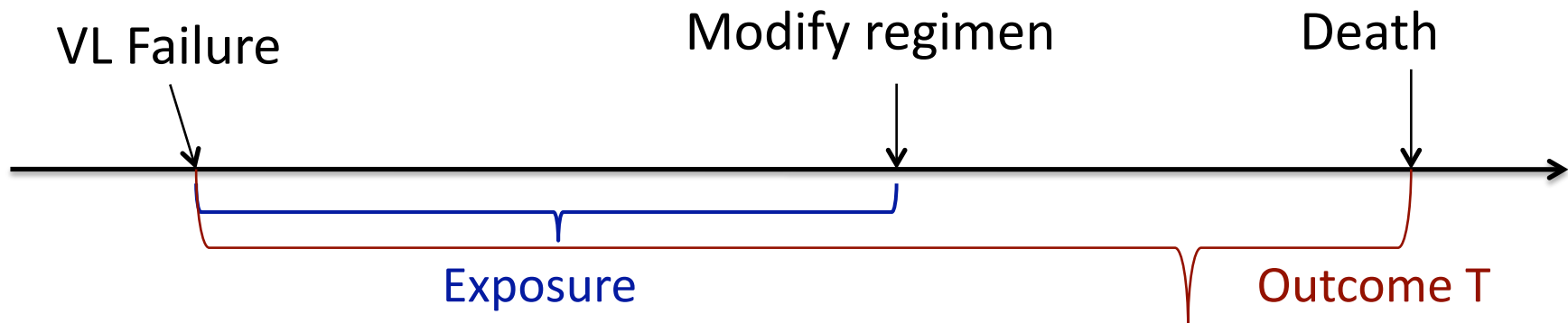
# Outline

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1. What types of questions can causal inference methods help us to answer?
2. When do standard analysis methods break down?
  - Time-dependent confounding of longitudinal treatments
3. What estimation tools are available?
  - Inverse Probability Weighting
  - Longitudinal G- computation
  - New methods coming...

# For Illustration, focus on first question

- Will delayed regimen modification after loss of viral suppression increase mortality? How much?



# Simple Confounding

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- Patients who are sicker (eg have lower CD4 counts/OD) at time of VL failure are more likely to modify immediately
  - Clinicians less likely to delay second line
  - These patients also more likely to die
  - Unadjusted analysis will underestimate harm from delayed switch
- Could control for this confounding by adjusting for baseline CD4 count (and other prognostic markers)

# Time-dependent confounding

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- Confounding by covariates after baseline
  - A subject whose CD4 counts decline rapidly following failure is more likely to modify therapy
  - A subject with rapidly declining CD4 has a worse prognosis
- Subjects who modify early are likely to be sicker than those who modify late
  - Even after controlling for baseline differences between the two groups
- Associations adjusted only for baseline covariates will still tend to underestimate the harm of delayed modification

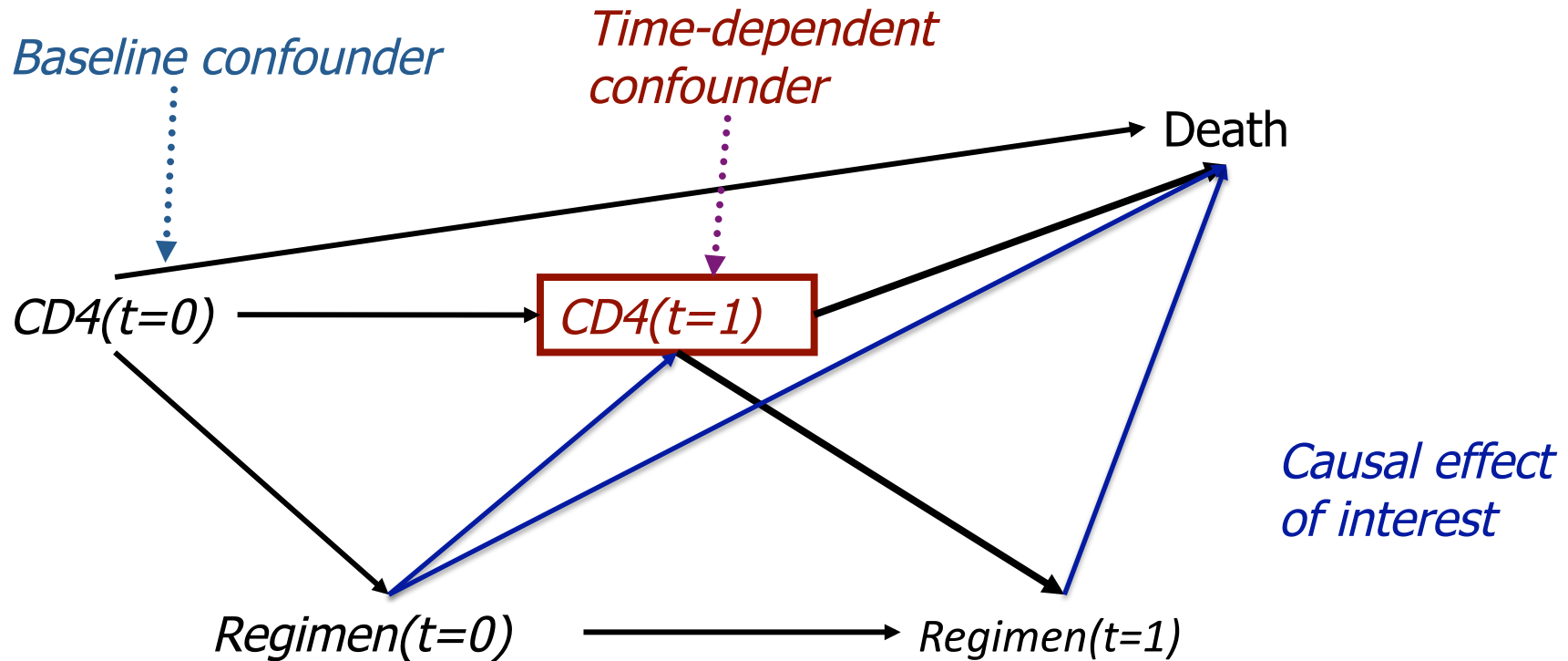


# Limitations of Standard Approaches

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- Why can't we just adjust for these post-baseline differences between subjects who modify early and those who modify late?
  - Ex. Regress vital status on delay time, baseline CD4, and time-updated CD4?
- Post baseline confounders may be affected by the exposure!
  - Part of the causal pathway we are interested in
  - Adjusting for them will bias our estimates

# Time-dependent confounding



- Can't control for  $CD4(t=1)$  in standard analyses: on causal pathway!

# Outline

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# Inverse probability of treatment weighting (IPTW)

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- Confounding can be viewed as a problem of biased sampling
- Because the decision when to switch is not randomized, sicker patients are over-represented in the group that switches earlier
- Goal of IPTW: reweight the sample to remove these imbalances

# Inverse probability of treatment weighting (IPTW)

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- IPTW: Creates a re-weighted dataset in which the exposure is randomized
  - Up-weight individuals whose observed exposure is rare given their covariates
  - Count the experience of rare individuals more than once to make up for people like them who are missing
- Ex. Sicker patients that delay modification get bigger weights

# Implementation of IPTW (1)

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1. Model the probability of modifying therapy (exposure) given covariate history among subjects who remain on their failing therapy
  - E.g. Using multivariable logistic regression
2. Use this model to calculate predicted probability of each subject having his observed exposure history (switch time)

## Implementation of IPTW (2)

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3. Weight=  $1 / \text{predicted probability of having observed switch time, given observed covariate history}$ 
  - Can also use stabilized weights
4. Fit a weighted regression of survival on time spent on failing therapy
  - Using specified MSM
  - Ex. Pooled logistic regression to estimate discrete hazard ratios

# Inverse weighting is just one possible way to adjust for time-dependent confounders...

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- IPTW: Control for confounding based on estimate of the probability of treatment over time given the past
- G-computation: Control for confounding based on estimate of the distribution of time-dependent confounders (and survival) over time given the past



# Longitudinal G-computation (1)

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1. Model the distribution of each covariate at each time point given past covariates and past treatment
  - Ex. Using series of multivariable pooled logistic and/or linear regression models
2. Model hazard of death at each time point given past covariates and past treatment
  - Ex. Using multivariable pooled logistic regression model

## Longitudinal G-computation (2)

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3. Use these models to simulate distribution of survival times under various delay times
  - Monte Carlo Simulation
  - Generate predicted survival times for each subject under each possible switch time
  
4. Using simulated data, fit a regression of survival on switch time
  - Using specified MSM, such as pooled logistic model

# Model Specification

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- Consistency of IPTW estimator depends on consistently estimating how treatment assignment depends on confounders
- Consistency of G-computation estimator depends on consistently estimating how covariates (and outcome) depend on the past
- Getting the specification of models to estimate either can be very challenging
- Data-adaptive algorithms
  - Cross-validation to choose best bias-variance tradeoff
  - Example: Superlearner

# Key Assumptions (1)

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## 1. No unmeasured confounders

- Key determinants of survival that also affect the decision when to modify are measured
  - Plausible?
    - Unrecorded comorbidities
    - Adherence
  - May be reasonable to argue that residual confounding expected to be negative
- > Estimate of harm due to delayed switch conservative...

## Key Assumptions (2)

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2. Adequate variability in exposure
  - “Experimental Treatment Assignment/Positivity”
  - Need variability in delay time
  - Beyond this- need that covariates do not deterministically predict switch
    - Ex. If subjects with low CD4 always switch, no information about experience of subjects with low CD4 that do not switch

# Estimator Comparison: IPTW

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- Pros
  - Well-established for studying the effect of longitudinal treatments
    - Track record in HIV
  - Easy to implement
- Cons
  - Inefficient (high variability)
  - Substantial finite sample bias when some covariate/treatment combinations are rare

# Estimator Comparison: G-computation

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- Pro
  - More efficient
    - Parametric MLE-based estimation
- Cons
  - Harder to implement (but doable)
  - Requires modeling dependence of all time-varying covariates and survival on the past
    - Getting model specification right can be a challenge!
    - With African cohort data, limited number of longitudinal covariates and time points

# A Third Option...

## Double Robust Estimators

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1. Augmented- IPTW
2. Targeted Maximum Likelihood estimation (TMLE)
  - More robust and efficient
  - TMLE software for point treatment effects available
  - TMLE software for longitudinal effects: under development...pilot coming...



# Current Research

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- Implementation of above with
  - SA-leDEA, some East Africa cohorts
  - CNICs (North America)
- Application of TMLE to longitudinal data/  
dynamic regime estimation
  - Comparison with standard IPTW approaches
- Prediction of viral failure using MEMs data
  - Machine-learning for high dimensional prediction

# Current Research

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- Analysis of RCT data
  - Methods described can be used to estimate improved “as treated” effect
  - Secondary analysis to look at additional questions than those initially targeted by the trial
    - Ex. DART: RCT comparing clinical vs. CD4 arms
    - How would impact of CD4 monitoring have differed if CD4 monitored at different frequency?
- Design and analysis of studies to investigate community based interventions
  - CRTs/ Combination prevention programs in Africa

# References: NPSEM/Counterfactuals/ G-computation formula

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# References: Standard and HA- MSM

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# References: Dynamic MSM

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# References: Data-adaptive Estimation

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