

Double-Sampling Designs for Dropouts

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Outline

- Mbarra's Dropouts
- Inverse Prob. Weights/Horvitz Thompson
- Double Sampling for Dropouts
- An IPW Estimator and Variance
- Future Directions

PEPFAR

- Massive scale-up of ARV treatment
- 1.3 Million initiated ARV from 2004-2008
- Outcome results essential
- Required large-scale follow-up

Mbarara ISS

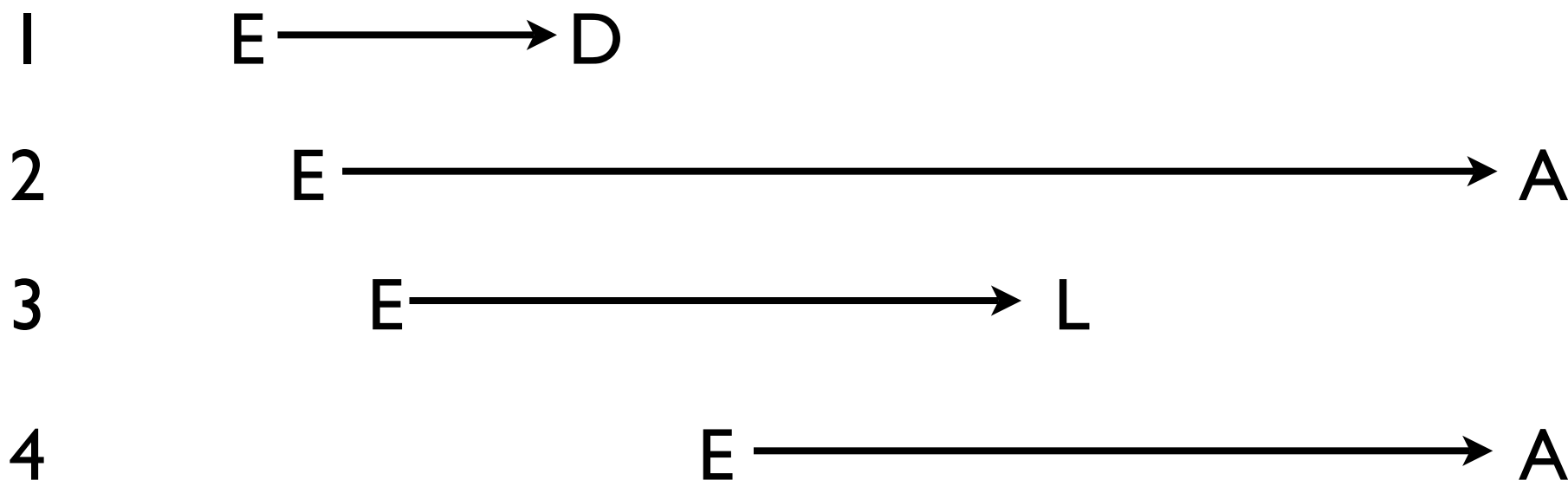
- Cohort of 3,340 HIV+ infected individuals
- Started on ARV from 1/04 to 7/07
- Followed through 7/1/07
- 56 died
2,530 alive as of 7/1/07
715 lost to FU prior to 7/1/07

Multiple Events

- Three possible events
 - Death
 - Dropout
 - Administrative Censoring
- Competing risks

Calendar Time Scale

Patient



2004

2005

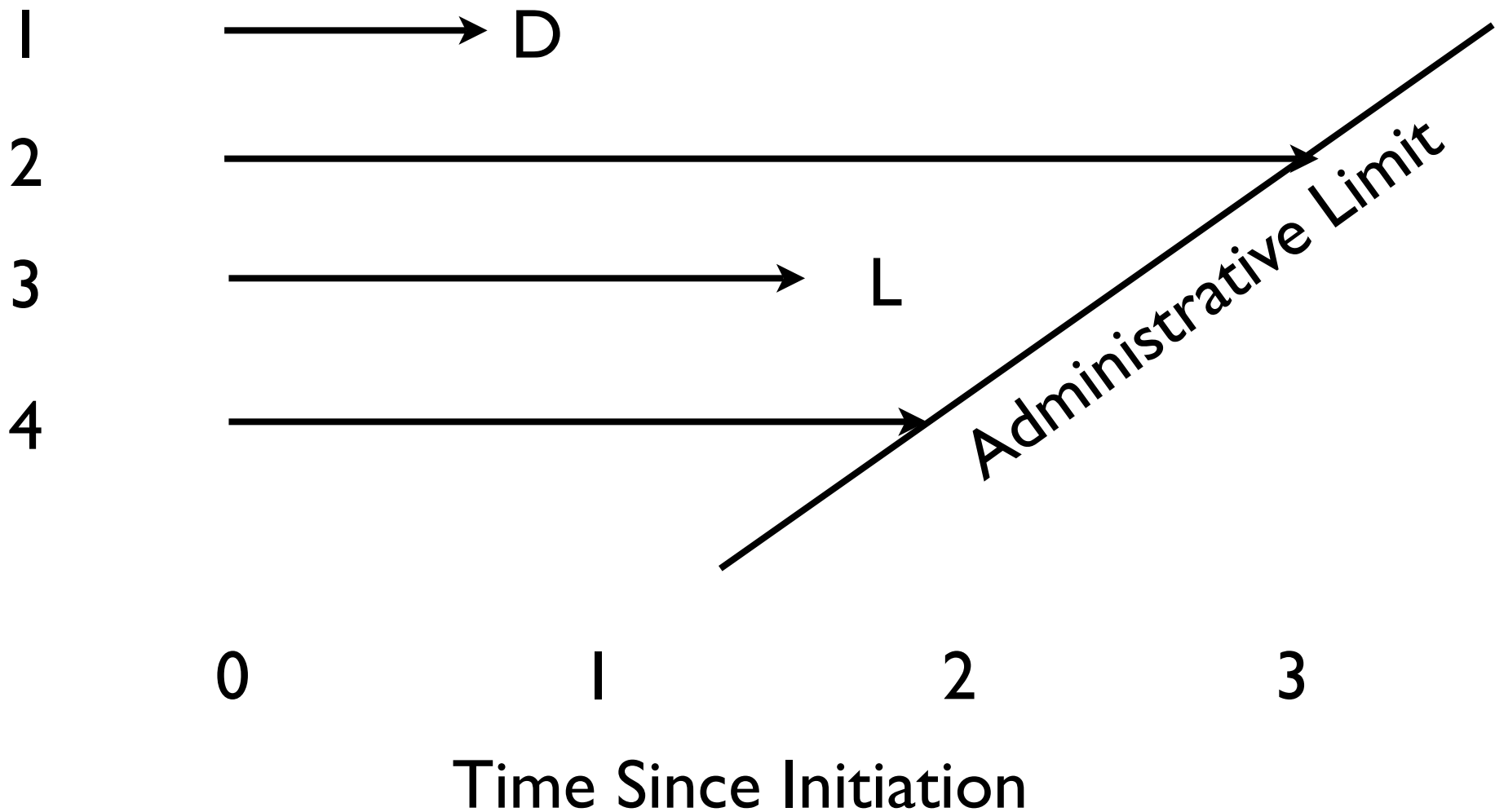
2006

2007

Date

Follow Up Time Scale

Patient



Notation

- C_i : time to administrative censoring
always known
- L_i : time to dropout
censored by C_i and T_i
- T_i : time to death
censored by C_i and L_i
- X_i : $\min(T_i, C_i)$, $\Delta_i = I(T_i \leq C_i)$
data in absence of dropouts
- $R^{obs}=0$ dropout, $R^{obs}=1$ non-dropout

Administrative Censoring

- Patients can only dropout if
 $L_i < \min(T_i, C_i)$
- Patients can only die if
 $D_i < \min(L_i, C_i)$
- Some people may have dropout later...

Administrative Censoring

- (T, C) are independent
very standard assumption
violated if demographics change over time
- Can be relaxed to (T, C) independent
given a series of covariates
- Conditional on R^{obs} , (T_i, C_i) *NOT* indep
example of “collider” stratification
creates dependent censoring

Dependent Dropout

- (T,L) are likely correlated
- Dropout suggest ARV discontinuation
- Hastens death
- Not easily handled
- What about observing after dropout
how about sampling?

Sampling Plan

- 3,340 HIV+ initiated ART
- $R^{obs}=1$: $n_1=2,625$
 - 2,569 Alive and in FU as of 7/1/09
 - 56 Died in Follow-Up
- $R^{obs}=0$: $n_0=715$, $\tilde{n}_0=79$
 - 95 sought, 79 vital status ascertained

Advantages of Sampling

- Very flexible
- Sampling prob can vary by individual
using ancillary data
- Valid framework for dependent censoring
in a way that is model-independent
no need to specify $\text{cor}(T, L)$
will get information on this

Horvitz Thompson

- Have finite population of size n
- Want to estimate
$$\tilde{\mu} = n^{-1} \sum_{i=1}^n x_i$$
- $\xi_i = 1$ indicated if i th person sampled
- Sample with probability $E(\xi_i = 1) = \pi_i$

- HT estimator
$$\hat{\mu} = n^{-1} \sum_{i=1}^n \frac{\xi_i}{\pi_i} x_i$$

Variance

$$\begin{aligned} & \text{var}\{\sqrt{n}(\tilde{\mu} - \hat{\mu})\} \\ &= n^{-1} \sum_{i=1}^n \sum_{j=1}^n \left(\frac{\pi_{ij} - \pi_i \pi_j}{\pi_{ij} \pi_i \pi_j} \right) \xi_i \xi_j x_i x_j \end{aligned}$$

π_{ij} is the probability that i and j are selected
termed the second-order inclusion probability

Simple Sample

- Total population is n
- Sample with equal probability $\pi_i = \tilde{n}/n$
- Sample with replacement
chose $\tilde{n}$ from n (putting balls back in jar)
- Sample with quota
chose exactly $\tilde{n}$ different from n
- Sample without quota
chose i th person with probability $\tilde{n}/n$

Same estimate, different variances!!

Second-Order Weights

under equal probability schemes

	π_{ii}	π_{ij}
w/ replace	$(\tilde{n}/n)^2$	$(\tilde{n}/n)^2$
quota	$\tilde{n}/n$	$\tilde{n}(\tilde{n}-1)/n(n-1)$
no quota	$\tilde{n}/n$	$(\tilde{n}/n)^2$

Variance for Finite Population

under “quota sampling”

$$\text{var}\{\hat{\mu}\} = \frac{(\pi^{-1} - 1)\sigma^2}{n} \quad n: \text{size of total popn}$$

$$\text{var}\{\hat{\mu}\} = \frac{(1 - \pi)\sigma^2}{\tilde{n}} \quad \tilde{n}: \text{size of sampled popn}$$

For the standard sample mean, π is effectively 0

$$\text{var}\{\bar{x}\} = \frac{\sigma^2}{\tilde{n}}$$

Double Sampling

Neyman, 1938

- group of $i=1, \dots, n$ subjects
sampled from a population
(first level of sampling)
some additional data avail. $(z_1, \dots, z_n)$
- second level of data
richer data collection on subset
sampling determined by $(z_1, \dots, z_n)$
 $\text{pr}(\xi_i=1 | z_i) = \pi_i$

Example: Two-Stage Case Control Study

- Bladder cancer case/control study
- Exposure: metal working fluids (MWF)
- Phase I: Data collected including
Q: *“Have you ever worked in metal industry?”*
- Phase II: Detailed work history/exposure collected
- Metal workers oversampled in Phase II

Other Examples

- Two-phase genotyping studies
- Nested case-control studies
- Case-cohort studies
- All valid, comparatively efft designs
- With significant cost savings

Back to Mbarara

Double Sample for Dropouts

- $R^{\text{obs}}=1$: observed death/admin censoring
 - data already completely observed
 - sampling fraction 1.00
- $R^{\text{obs}}=0$: observed dropouts
 - n_0 dropouts
 - sample $\tilde{n}_0$ for vital ascertainment
 - sampling fraction $\tilde{n}_0/n_0 = \pi$

Observed Data

- (X_i, Δ_i) if $\xi_i = 1$
- ignored otherwise
- $\xi = 1$ if $R^{\text{obs}} = 1$ or $R^{\text{obs}} = 0$ & sampled
- $\text{pr}(\text{sampled} \mid R^{\text{obs}} = 0) = \pi$
- Data is MAR conditional on R^{obs}

Frangakis and Rubin

- Double sampling estimate of survivor function
- Ignore data if $\xi=0$
if dropout & not dble sampled, drop data
- Constructed survivor estimate
- Showed that (T,C) not independent conditional on the value of R^{obs}
- Estimate hazard and transform to survival

Frangakis Rubin Estimator

$$\hat{\Lambda}(t) := \sum_{g=0,1} \int_0^t \hat{w}_g(u) d\hat{\Lambda}_g^{\text{crd}}(u),$$

where

$$\hat{w}_g(t) := \frac{\hat{\pi}_g(t) \hat{p}_g}{\sum_{g'=0,1} \hat{\pi}_{g'}(t) \hat{p}_{g'}}, \quad \hat{\pi}_g(t) := Y_g(t)/n_g,$$

consistent with a Gaussian limiting distribution

Their representation

- Cumulative hazard is weighted sum
- Of crude hazards in 2 groups
- Weight varies with time
- Not an intuitive representation
why does it work?

Double-Sampling

- $\Lambda(t)$ True cumulative hazard function
- $\tilde{\Lambda}(t)$ Nelson-Aalen estimator
if complete data avail on cohort
- $\hat{\Lambda}(t)$ The FR estimator
based on dble-sampled data

More Notation

- $N(t) = I(X \leq t, \Delta = 1)$
- $Y(t) = I(X \geq t)$
- H_1 : set of people with $R^{\text{obs}} = 1$
has size $n_1, \tilde{n}_1$ sampled ($n_1 = \tilde{n}_1$)
 i in H_1 $\pi_i = 1$
- H_0 : set of people with $R^{\text{obs}} = 0$
has size $n_0, \tilde{n}_0$ sampled
 i in H_0 : $\pi_i = \tilde{n}_0/n_0$

Complete Data on Cohort

$$\bar{N}_g(t) = n_g^{-1} \sum_{i \in \mathcal{H}_g} N_i(t)$$

$$\bar{Y}_g(t) = n_g^{-1} \sum_{i \in \mathcal{H}_g} Y_i(t)$$

$$\bar{N}(t) = \sum_g \frac{n_g}{n} \bar{N}_g(t)$$

$$\bar{Y}(t) = \sum_g \frac{n_g}{n} \bar{Y}_g(t)$$

Other Representation

$$\Lambda(t) = \int_0^t \frac{\mathcal{N}(du)}{\mathcal{Y}(u)}$$

$$\text{as } n \rightarrow \infty$$

$$\bar{N}(t) \rightarrow \mathcal{N}(t)$$

$$\bar{Y}(t) \rightarrow \mathcal{Y}(t)$$

$$\tilde{\Lambda}(t) = \int_0^t \frac{\bar{N}(du)}{\bar{Y}(u)}$$

Horvitz Thompson

$$\hat{N}_g(t) = \tilde{n}_g^{-1} \sum_{i \in \mathcal{H}_g} \frac{\xi_i}{\pi_i} N_i(t)$$

$$\hat{Y}_g(t) = \tilde{n}_g^{-1} \sum_{i \in \mathcal{H}_g} \frac{\xi_i}{\pi_i} Y_i(t)$$

$$\hat{N}(t) = \sum_g \frac{n_g}{n} \hat{N}_g(t)$$

$$\hat{Y}(t) = \sum_g \frac{n_g}{n} \hat{Y}_g(t)$$

Approximations

$$\hat{N}_1(t) = \bar{N}_1(t)$$

$$\hat{Y}_1(t) = \bar{Y}_1(t)$$

$$\hat{N}_0(t) \longrightarrow \bar{N}_0(t)$$

$$\hat{Y}_0(t) \longrightarrow \bar{Y}_0(t)$$

A Natural Estimator

$$\hat{\Lambda}(t) = \int_0^t \frac{\hat{N}(du)}{\hat{Y}(u)}$$

identical to FR estimator and to

$$\hat{\Lambda}(t) = \int_0^t \frac{\sum_i \frac{\xi_i}{\pi_i} N_i(du)}{\sum_i \frac{\xi_i}{\pi_i} Y_i(u)}$$

and has a IPW representation

**But what about
variance?**

FR Variance

$$V_{\{p_1\}}(s, t) + \sum_{g=0,1} V_{\{\pi_g\}}(s, t) + V_{\{\Lambda_g\}}(s, t) + V_{\{\pi_g, \Lambda_g\}}(s, t) \\ + V_{\{\pi_g, \Lambda_g\}}(t, s), \quad (\text{A.2})$$

where

$$V_{\{p_1\}}(s, t) := p_1 p_0 \int_0^t d\Lambda_{\{p_1\}}^{\text{crd}}(u) \int_0^s d\Lambda_{\{p_1\}}^{\text{crd}}(u^*)$$

and, for $k_0 = 1/\{p_0 p^{(S)}\}$, $k_1 = 1/p_1$, and $g = 0, 1$,

$$V_{\{\pi_g\}}(s, t) := k_g \int_0^t \int_0^s [\pi_g\{\max(u, u^*)\} - \pi_g(u)\pi_g(u^*)] \\ \times d\Lambda_{\{\pi_g\}}^{\text{crd}}(u^*) d\Lambda_{\{\pi_g\}}^{\text{crd}}(u),$$

$$V_{\{\Lambda_g\}}(s, t) := k_g \int_0^{\min(s,t)} \frac{\{w_g(u)\}^2}{\pi_g(u)} d\Lambda_g^{\text{crd}}(u)$$

and

$$V_{\{\pi_g, \Lambda_g\}}(s, t) \\ := -k_g \int_0^{\min(s,t)} \int_u^s \frac{\pi_g(u^*)}{\pi_g(u)} w_g(u) d\Lambda_{\{\pi_g\}}^{\text{crd}}(u^*) d\Lambda_g^{\text{crd}}(u).$$

Two-Part Variance

$$\begin{aligned}\sqrt{n}\{\hat{\Lambda}(t) - \Lambda(t)\} &= \\ \sqrt{n}\{\hat{\Lambda}(t) - \tilde{\Lambda}(t)\} &+ \sqrt{n}\{\tilde{\Lambda}(t) - \Lambda(t)\}\end{aligned}$$

last two terms are independent

$$\sigma^2(t) = \text{var}\{\hat{\Lambda}(t) - \Lambda(t)\}$$

$$\sigma_1^2(t) = \text{var}\{\hat{\Lambda}(t) - \tilde{\Lambda}(t)\}$$

$$\sigma_2^2(t) = \text{var}\{\tilde{\Lambda}(t) - \Lambda(t)\}$$

$$\sigma^2(t) = \sigma_1^2(t) + \sigma_2^2(t)$$

Variance Decomposition

$\sigma^2(t)$: Total Variance

$\sigma_2^2(t)$: Variance if full cohort observed

Usual variance for Nelson-Aalen estimate

Easily estimated

$\sigma_1^2(t)$: Variance due to double sampling

HT type variance

Estimation more complicated

Nelson-Aalen Variance

$$\sigma_2^2(t) = \int_0^t \frac{\Lambda(du)}{\mathcal{Y}(u)}$$

$$\hat{\sigma}_2^2(t) = \int_0^t \frac{\hat{\Lambda}(du)}{\hat{Y}(u)}$$

$$= \frac{n_0}{n} \int_0^t \frac{\hat{N}_0(du)}{\hat{Y}^2(u)} + \frac{n_1}{n} \int_0^t \frac{\hat{N}_1(du)}{\hat{Y}^2(u)}$$

$$= \int_0^t \frac{\hat{N}(du)}{\hat{Y}^2(u)}$$

Double-Sample Variance

long HT-based variance arguments lead to

$$\begin{aligned}\hat{\sigma}_1^2(t) &= (\pi^{-1} - 1) \frac{n_0}{n} \int_0^t \frac{\hat{N}_0(du)}{\hat{Y}^2(u)} \\ &= n^{-1} \sum_{i: R^{obs}=0} \int_0^t \frac{\xi_i}{\pi_i} \frac{(\pi_i^{-1} - 1) N_i(du)}{\tilde{Y}^2(u)}. \\ &= n^{-1} \sum_{i=1}^n \int_0^t \frac{\xi_i}{\pi_i} \frac{(\pi_i^{-1} - 1) N_i(du)}{\hat{Y}^2(u)}\end{aligned}$$

The total variance

$$\begin{aligned}\hat{\sigma}^2(t) &= \hat{\sigma}_1^2(t) + \hat{\sigma}_2^2(t) \\ &= n^{-1} \sum_{i=1}^n \int_0^t \frac{\xi_i}{\pi_i} \frac{N_i(du)}{\hat{Y}^2(u)} + \\ &\quad n^{-1} \sum_{i=1}^n \int_0^t \frac{\xi_i}{\pi_i} \frac{(\pi_i^{-1} - 1)N_i(du)}{\hat{Y}^2(u)} \\ &= n^{-1} \sum_{i=1}^n \int_0^t \frac{\xi_i}{\pi_i} \frac{\pi_i^{-1} N_i(du)}{\hat{Y}^2(u)}\end{aligned}$$

Source of Variance

$R^{\text{obs}}=0$

$R^{\text{obs}}=1$

$\hat{\sigma}_1^2(t)$	$(\pi^{-1} - 1) \frac{n_0}{n} \int_0^t \frac{\hat{N}_0(du)}{\hat{Y}^2(u)}$	0
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$\hat{\sigma}_2^2(t)$	$\frac{n_0}{n} \int_0^t \frac{\hat{N}_0(du)}{\hat{Y}^2(u)}$	$\frac{n_1}{n} \int_0^t \frac{\hat{N}_1(du)}{\hat{Y}^2(u)}$
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$\hat{\sigma}^2(t)$	$\pi^{-1} \frac{n_0}{n} \int_0^t \frac{\hat{N}_0(du)}{\hat{Y}^2(u)}$	$\frac{n_1}{n} \int_0^t \frac{\hat{N}_1(du)}{\hat{Y}^2(u)}$
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Some observations

- $R^{\text{obs}}=1$: no contribution to double-sample variance
- $R^{\text{obs}}=0$: ratio of NA variance to FR variance
 $= 1/\pi = n_0/\tilde{n}_0$
just the ratio of sample size in $R^{\text{obs}}=0$
between DS data and full data
- Intuitive look at the variance

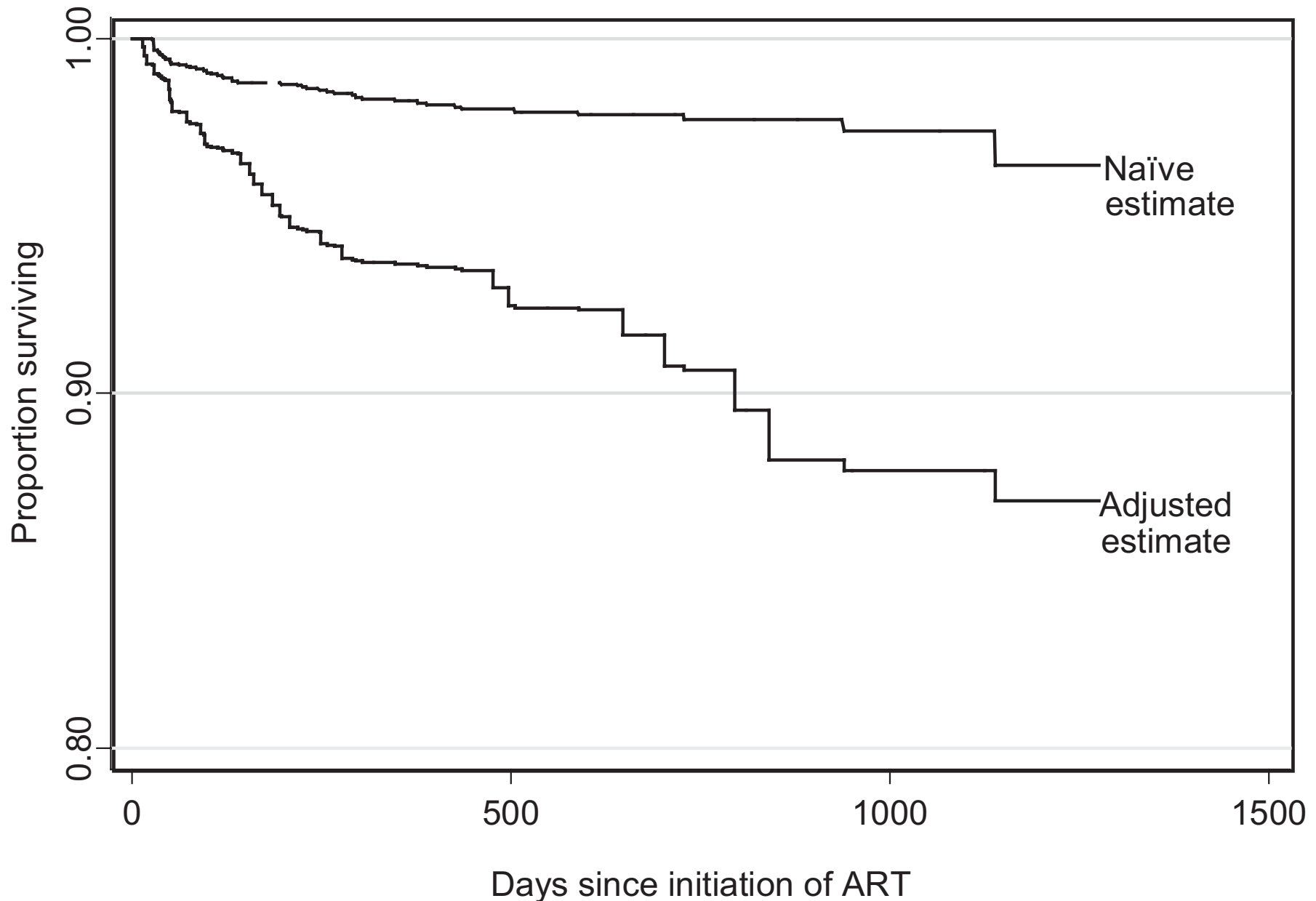
Variance Estimate

- Easily computed
- Demystifies the form
- Facilitates sample size calculations
look at effect of various sample fractions
- Performs great in simulations

Data Example

- Cohort of 3,340 HIV+ infected individuals
- $R^{\text{obs}}=1$: $n_1=2,625$ (56 died)
- $R^{\text{obs}}=0$: $n_0=715$, $\tilde{n}_0=79$ (26 died)
- $\pi = 1/9.18 = 0.109$
- Rate of death is about 16 times higher
in dropouts compared to non-dropouts

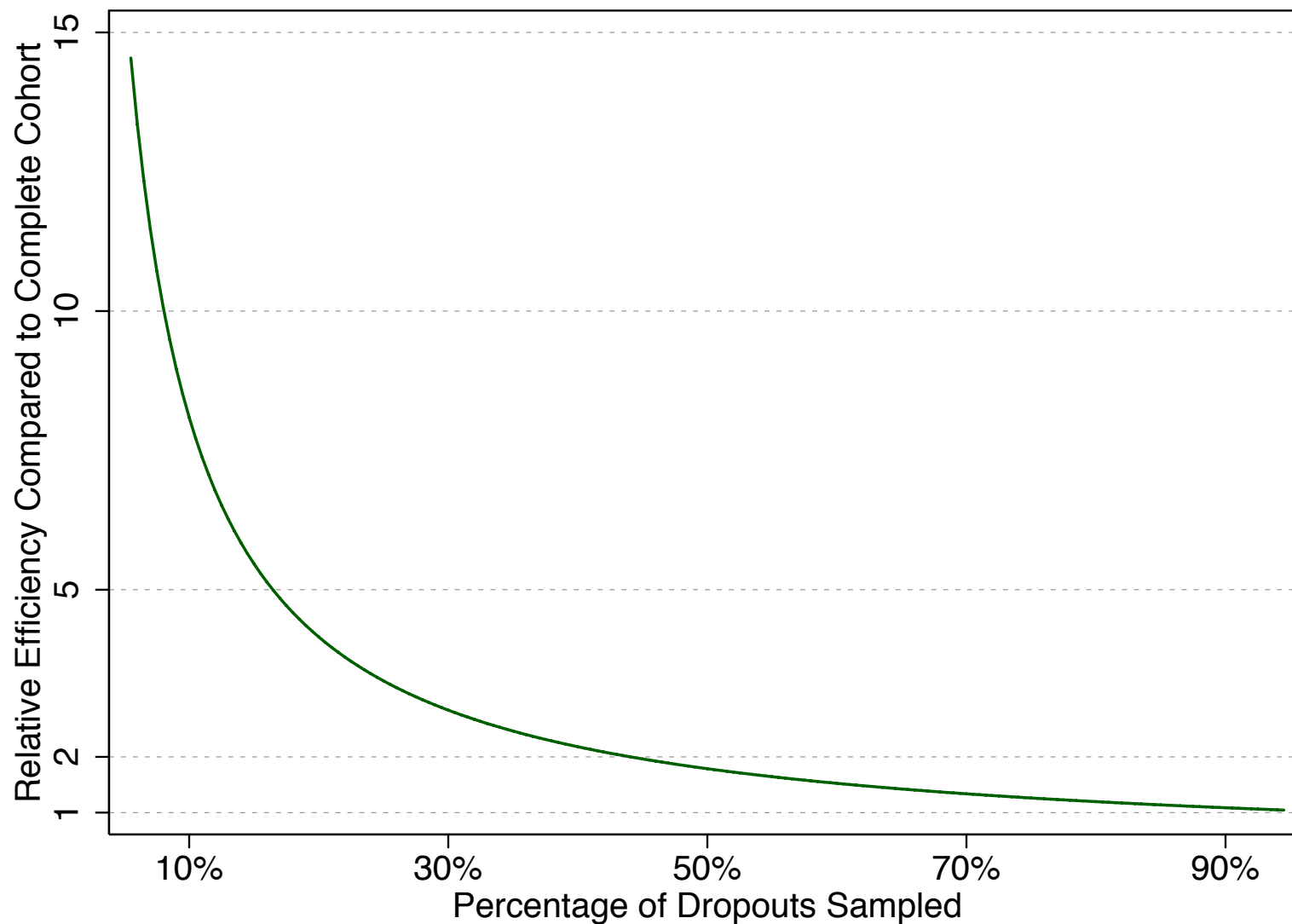
FR and Naïve Survival



Relative Efficiency

- Trade-off between sampling fractions
- What is efficiency of sampling ρ dropouts compared to all dropouts
- Can be consistently estimated
- Based on Mbarara data

Relative Efficiency



Future Directions

- Apply insights from survey statistics
 - formulae and approximations
 - post-stratification, calibration
auxiliary variables => more efficiency
- Look at using non-sample dropout data

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