

# **MEASURING HIV INCIDENCE: APPROACHES & CHALLENGES**

# VOCABULARY

HIV PREVALENCE

HIV INCIDENCE

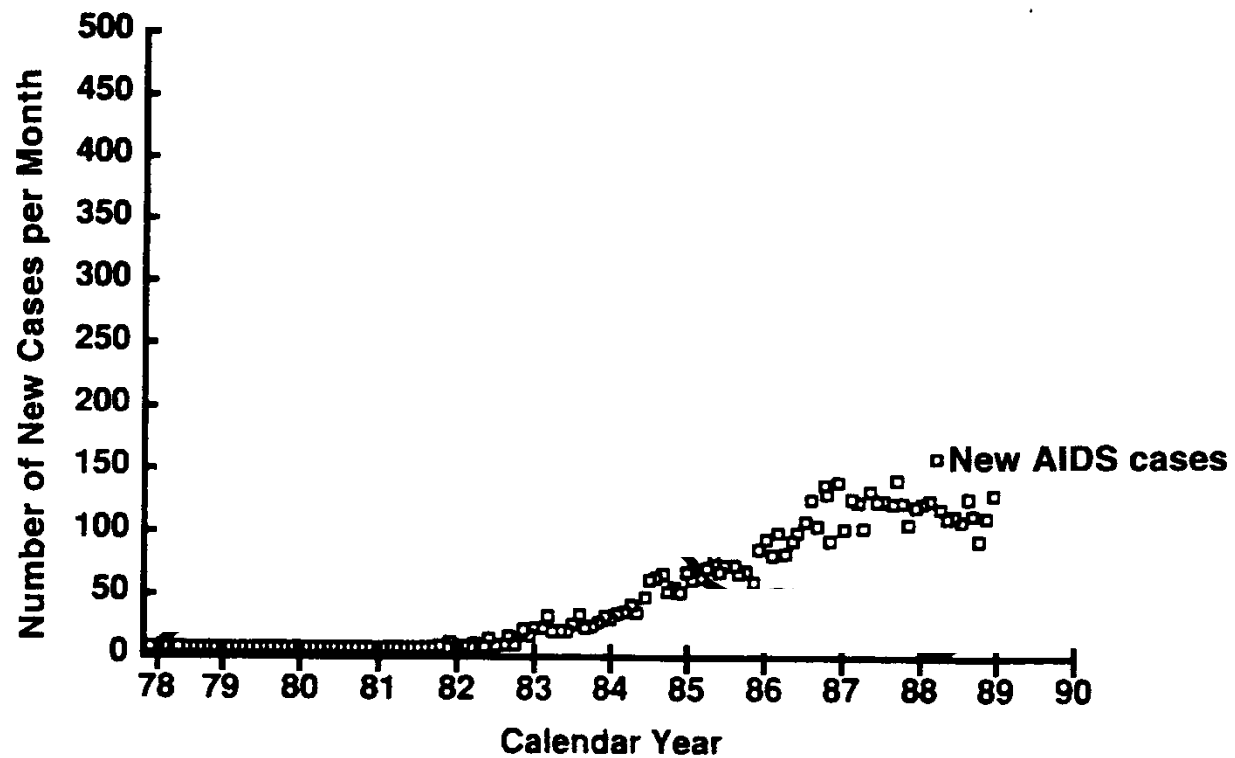


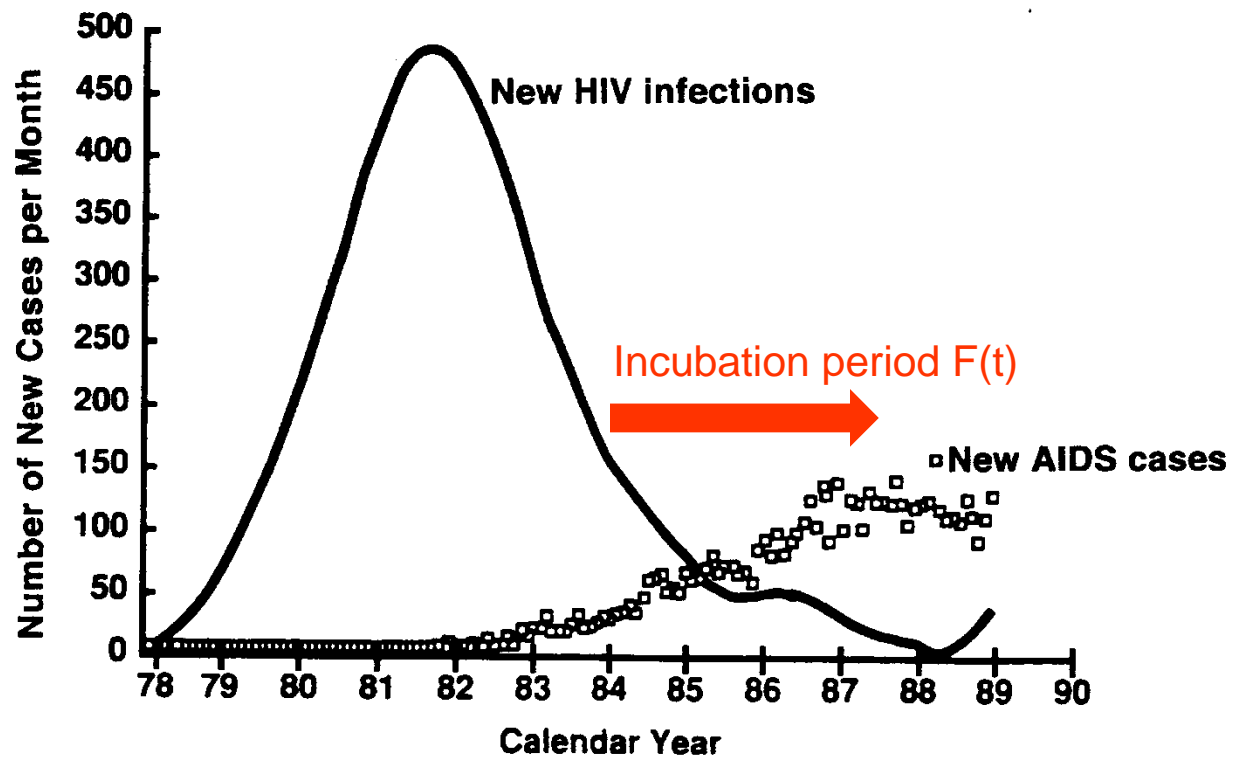
# U.S. Centers for Disease Control 2008

HIV PREVALENCE = 1.1 MILLION

HIV INCIDENCE = 56,000

# HIV PREVALENCE





# BACK-CALCULATION

## DECONVOLUTION

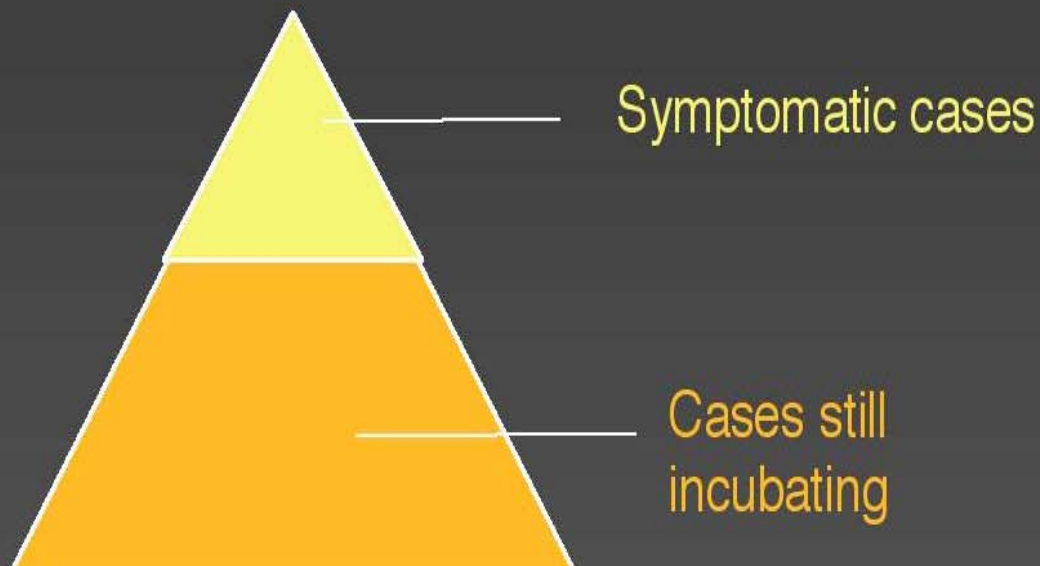
AIDS DX DATE = HIV INFECTION DATE + INCUBATION

$$E(y_i) = \int_0^{t_i} I(s) F(t_i - s) - F(t_{i-1} - s) ds$$

$$E(\mathbf{y}) = \boldsymbol{\beta} \mathbf{Z}$$

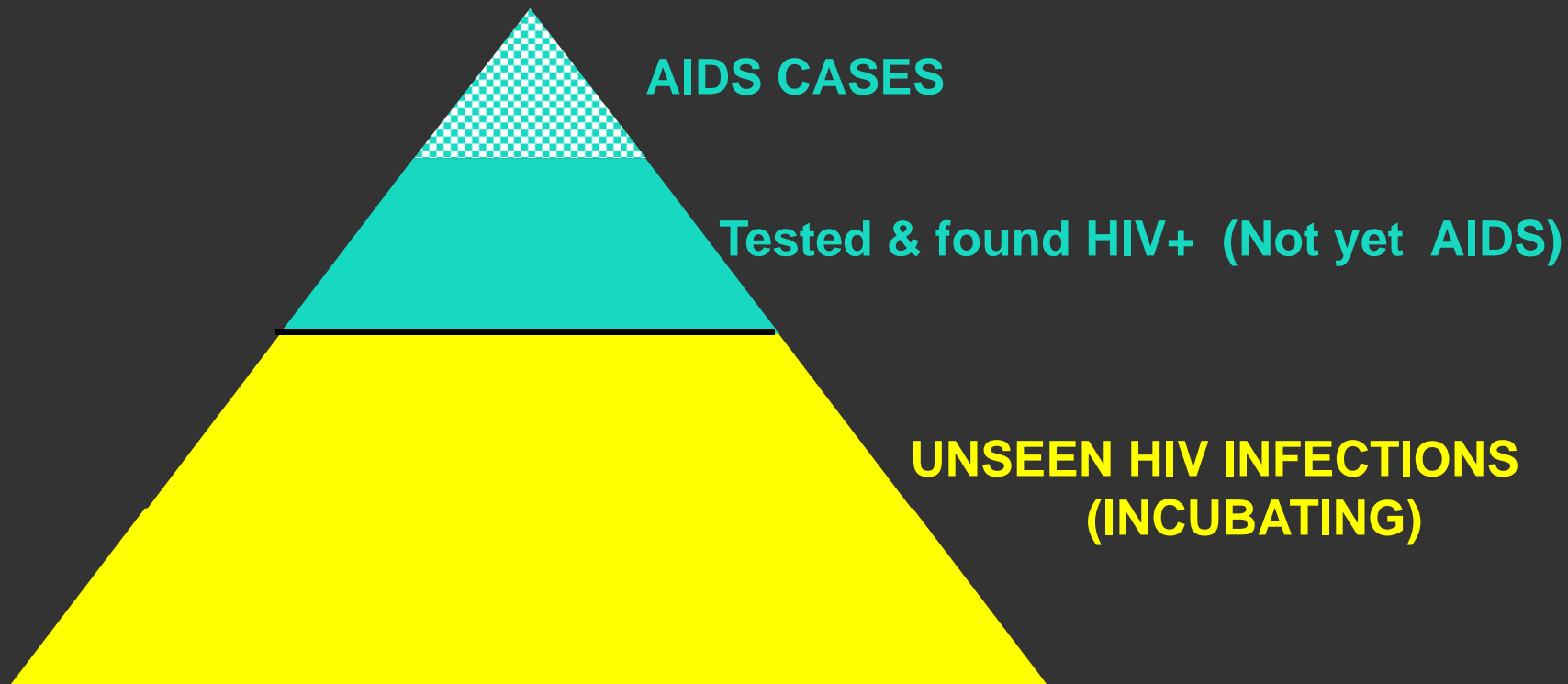
$$\text{var}(\mathbf{y}) = \sigma^2 E(\mathbf{y})$$

# BACK-CALCULATION: ICEBERG EFFECT



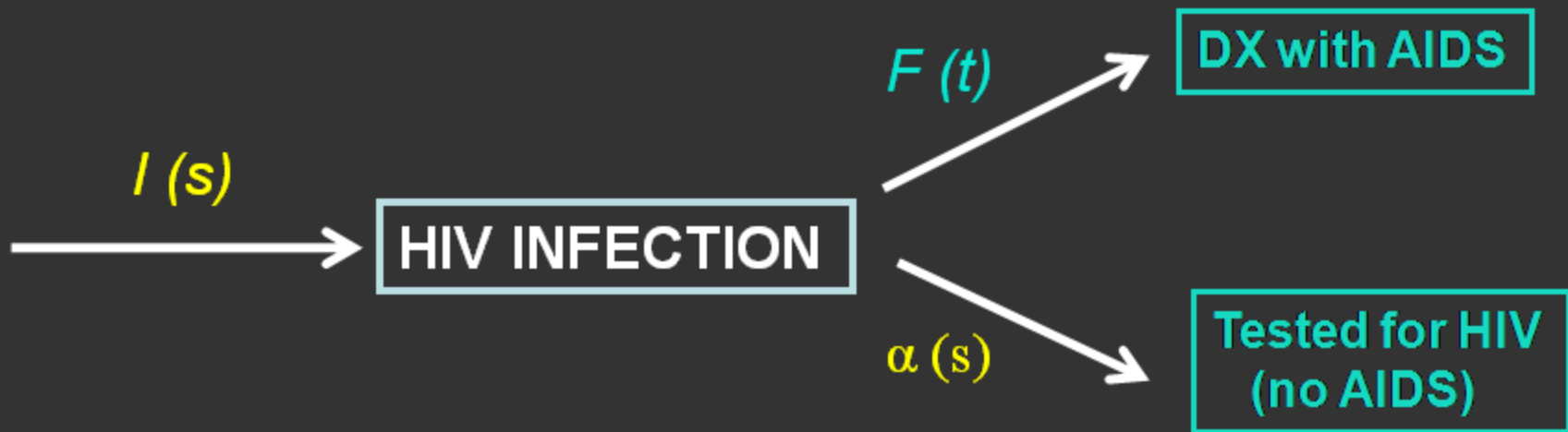


# BACK-CALCULATION: ICEBERG EFFECT



# EXTENDED BACK-CALCULATION

DECONVOLUTION  
Rhodes (2008), Hall et al (2008)



# US HIV PREVALENCE

**U.S. HIV PREVALENCE=1.1 million (JAMA, 2008)**

- STATISTICAL DECONVOLUTION
- UNCERTAINTIES
  - INCUBATION PERIOD
  - COMPLETENESS OF CASE REPORTING
  - PARAMETRIC MODEL FOR  $\alpha(s)$ ,  $I(s)$
  - RECENT INFECTIONS

# **CURRENT HIV INCIDENCE**

**MEASURES THE LEADING EDGE OF THE EPIDEMIC**

- **COHORT STUDY**
- **CROSS-SECTIONAL BIOMARKER APPROACH**

# CURRENT HIV INCIDENCE

## COHORT STUDY

$$\text{Incidence} = \frac{\text{incident infections}}{\text{person time}}$$

## ISSUES

Assembling cohort is difficult

Counseling may reduce HIV risk

Incidence is changing over time

Selection bias: who returns for follow-up?

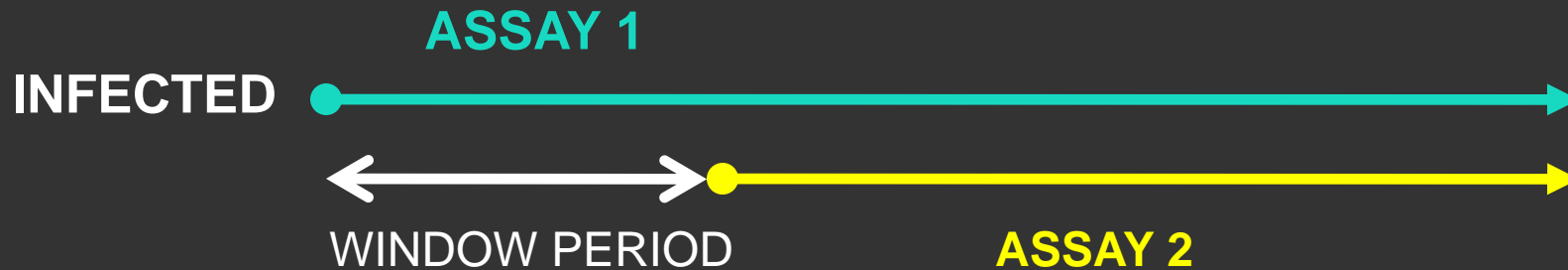
# CURRENT HIV INCIDENCE

## BIOMARKER APPROACH

- CROSS-SECTIONAL SAMPLE
- COLLECT BIOMARKERS OF RECENT INFECTION AT BASELINE
- SNAPSHOT APPROACH

# BIOMARKER APPROACH

# BIOMARKER APPROACH



$S(t)$  window period survival distrib.  
 $\mu$  = mean window period



# BIOMARKER APPROACH

$$\begin{aligned} P(\text{Window}) &= \int_0^{\infty} g(t) S(t) dt \\ &\approx g \int S(t) \\ &= g \mu \end{aligned}$$

$\mu$ = mean window period

$g$ =pdf of window entry times

# BIOMARKER APPROACH

PROPORTION IN THE WINDOW = INCIDENCE  $\times \mu$

$$\hat{I} = \frac{x}{N\mu}$$

X= # in window : assay 1 +, assay 2 -

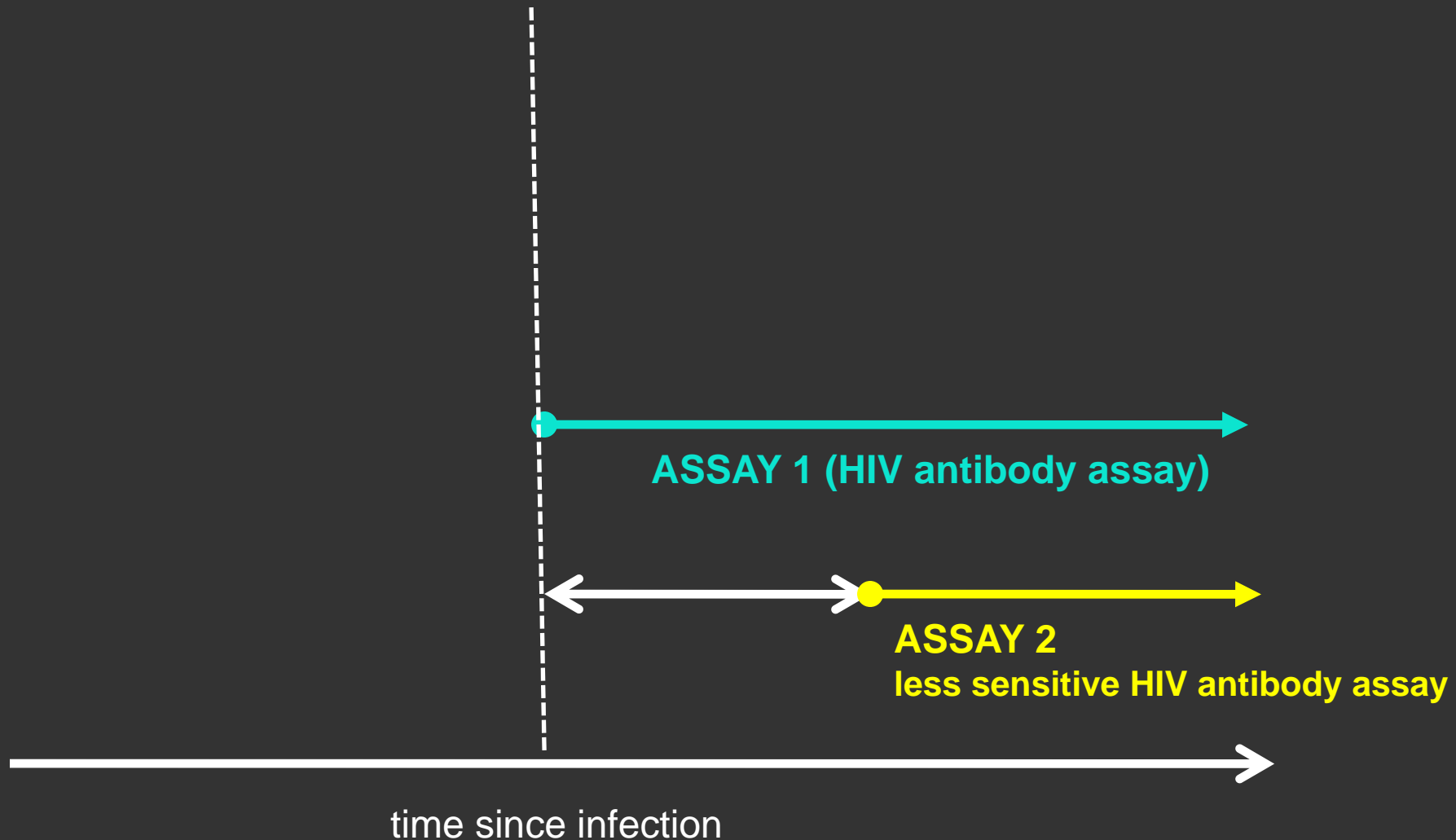
N= # assay 2 -

- CROSS-SECTIONAL SAMPLE

- NO FOLLOW-UP!

- SNAPSHOT ESTIMATOR

# ANTIBODY BIOMARKERS





# BIOMARKERS FOR RECENT INFECTION

DETUNED ASSAY\*  $\mu=129$  days

BED ASSAY\*  $\mu=156$  days

\*Need to assay only those HIV antibody +

# NEW U.S ESTIMATE OF HIV INCIDENCE

**56,300 ANNUAL NEW HIV INFECTIONS IN U.S.  
*JAMA* 2008**

- BASED ON CROSS-SECTIONAL BIOMARKER APPROACH (BED ASSAY)
- ESTIMATE PROBABILITY PERSON GETS HIV TEST
- 19 EXTRAPOLATED TO 50 STATES

# BIOMARKER APPROACH

U.S.  
EUROPE  
INDIA  
THAILAND  
CARIBBEAN  
AFRICA



# CONTROVERSY

## CDC

“The BED biomarker assay is the preferred approach for calculating HIV incidence in the U.S.”

## UNAIDS

“Does not recommend the BED assay for determining incidence”



# ZIMBABWE ZVITAMBO STUDY OF NEW MOTHERS

BIOMARKER ESTIMATE  $\approx 2 \times$  HIGHER THAN COHORT  
Hargrove (2008)

- Follow-up bias?
- “False recents”  
McWalter and Welte (2009)  
Wang and Lagakos (2009)





# BIOMARKER APPROACH: THEORY

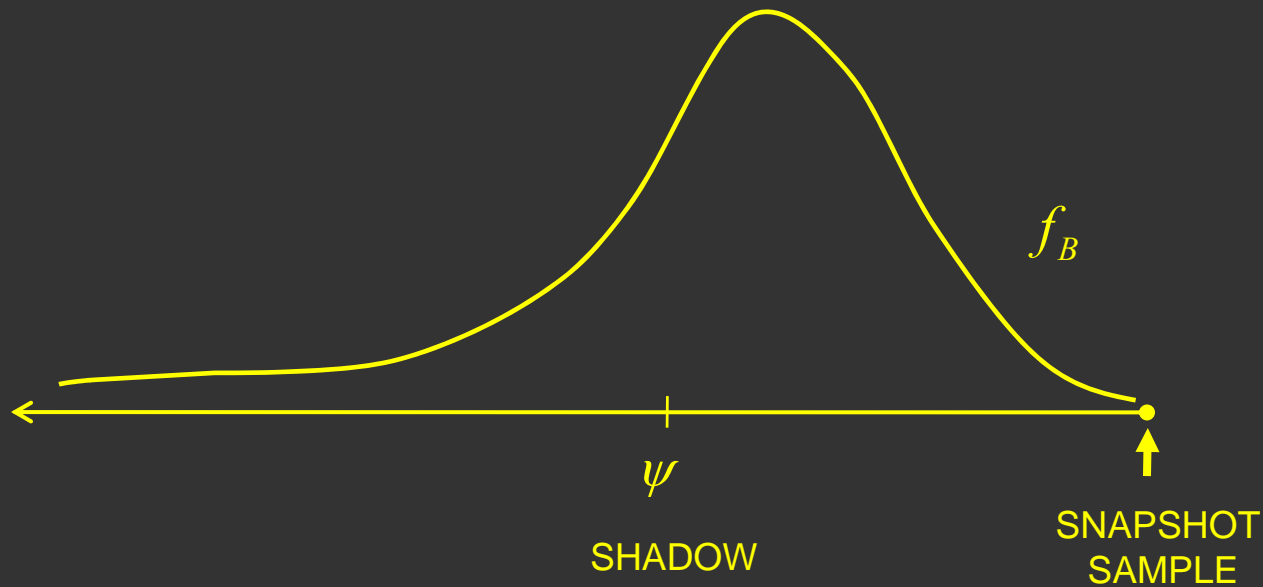
INCIDENCE NOT CONSTANT ?

$$\hat{I} \rightarrow \bar{I} \approx \int_0^{\infty} I(t) f_B(t) dt$$

$$f_B(t) = \frac{S(t)}{\mu} = \text{backward recurrence density}$$

$$\psi = \int t f_B(t) dt = \int t \frac{S(t)}{\mu} dt$$

$$\bar{I} \approx I(\psi) + \frac{I''(\psi)V}{2}$$



SHADOW  $\psi$  = mean of backward recurrence

$$\bar{I} \approx I(\psi)$$

$$\psi = \int t f_B(t) dt = \int t \frac{S(t)}{\mu} dt = \mu(1 + CV^2)$$

# SHADOWS

SHADOW DEPENDS ON MEAN AND CV OF WINDOW

Mean window (yrs)	<u>SHADOW (years)</u>			
	Coefficient of variation			
	<u>.50</u>	<u>1.0</u>	<u>2.0</u>	<u>3.0</u>
.25	.16	.25	.63	1.25
.50	.31	.50	1.25	2.50
1.00	.63	1.00	2.50	5.00
1.50	.94	1.50	3.75	7.50



# ACCURACY ?

- BIAS & VARIANCE
- ACCURACY OF ASSAY FOR HIV INCIDENCE:  
MEAN WINDOW ( $\mu$ )  
SHADOW (CV,  $\mu$ )
- BIG  $\mu$  OR SMALL  $\mu$ ?  
TRADEOFF : BIAS VS VARIANCE

# ISSUES WITH BED/DETUNED ASSAY METHOD

- **ELITE CONTROLLERS**
  - LONG TAILS OF  $S(t)$
  - LOW VIRAL LOADS

Brookmeyer, “On the Statistical Accuracy of Biomarker Assays for HIV Incidence,”  
J. AIDS, 2010



# ISSUE

## ELITE CONTROLLERS: MIXTURE MODEL

- FAST PROGRESSORS THROUGH WINDOW =  $\gamma$
- SLOW PROGRESSORS THROUGH WINDOW =  $1 - \gamma$   
(e.g. elite controllers)

$$S(t) = \gamma S_1(t) + (1 - \gamma) S_2(t)$$

$$shadow = \frac{\mu_1^2 (1 + CV_1^2) \gamma + \mu_2^2 (1 + CV_2^2) (1 - \gamma)}{2(\mu_1 \gamma + \mu_2 (1 - \gamma))}$$

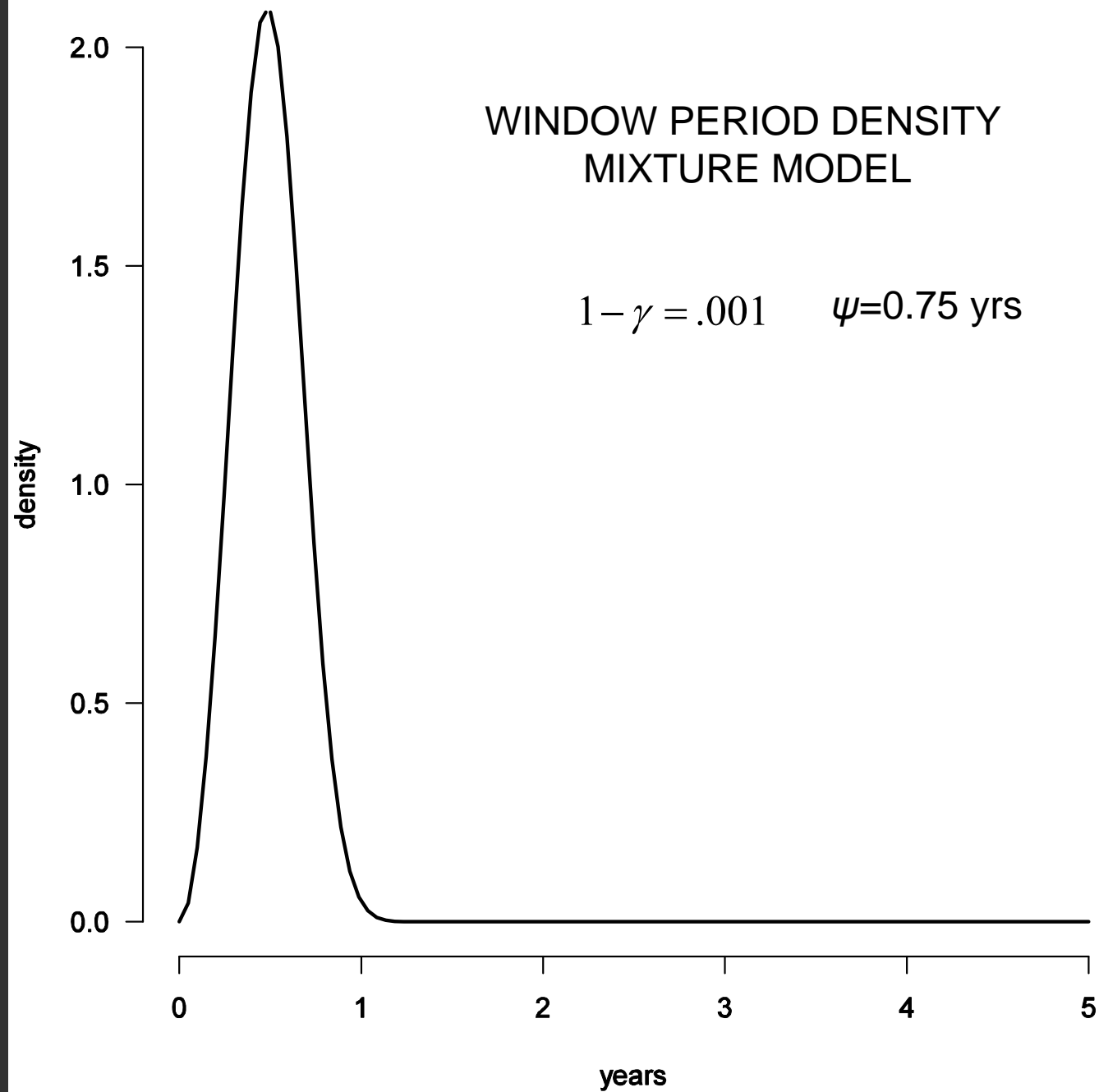
# MIXTURE MODEL: ELITE CONTROLLERS

$$S(t) = \gamma S_1(t) + (1-\gamma) S_2(t)$$

$\mu_1 = 0.5$  yrs  
weibull  $cv_1 = .36$

$\mu_2 = 20$  yrs  
weibull  $cv_2 = .50$

$1-\gamma$	$\mu$ yrs	Shadow yrs
.000	0.50	0.28
.001	0.52	0.75
.005	0.60	2.33
.010	0.70	3.81
.015	0.79	4.91
.020	0.89	5.77





density

2.0  
1.5  
1.0  
0.5  
0.0

0

1

2

3

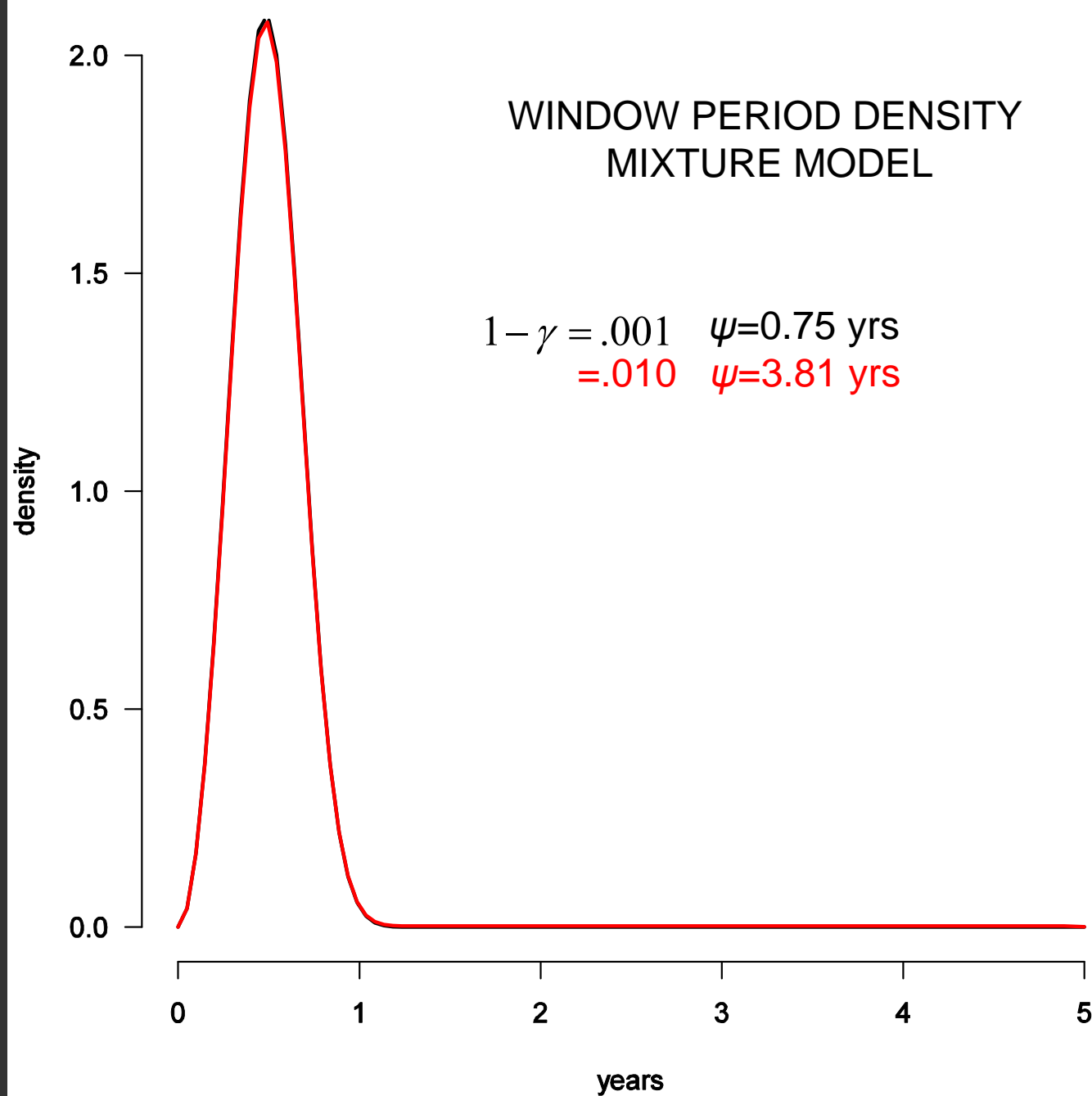
4

5

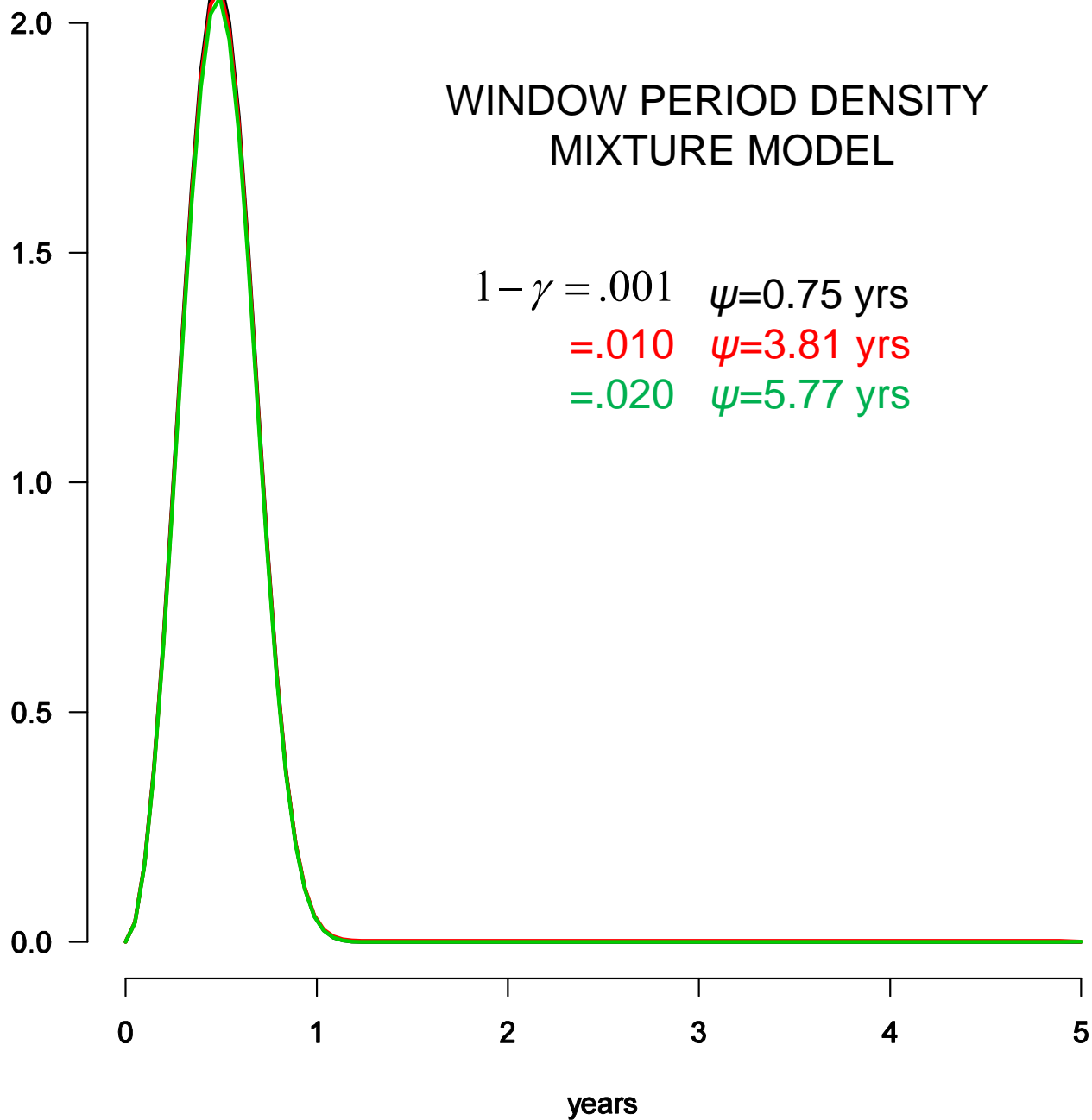
years

# WINDOW PERIOD DENSITY MIXTURE MODEL

$1 - \gamma = .001$     $\psi = 0.75$  yrs  
                   $= .010$     $\psi = 3.81$  yrs



density





# ZIMBABWE CONTROVERSY

- Biomarker incidence higher than cohort estimate in study of new mothers
- A shadow of 2+ years is produced with only 0.5 % elite controllers.
- HIV Incidence :  
Pre-partum vs post partum period  
Counseling/behavior change



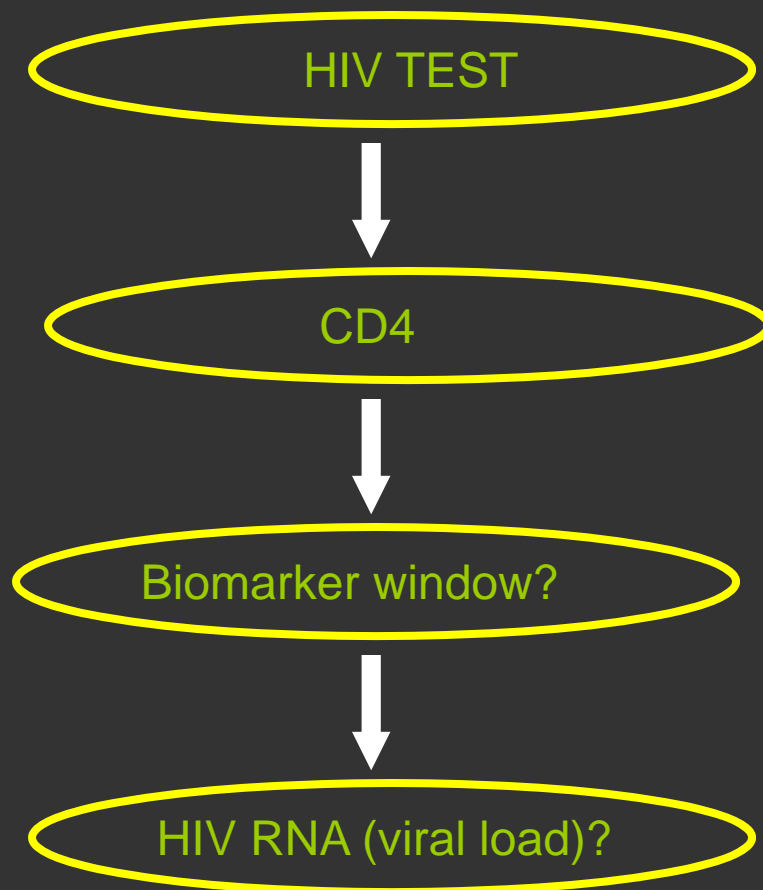
TINY ALMOST IMPERCEPTIBLE  
DIFFERENCES IN TAIL BEHAVIOR  
CAN HAVE HUGE EFFECTS !



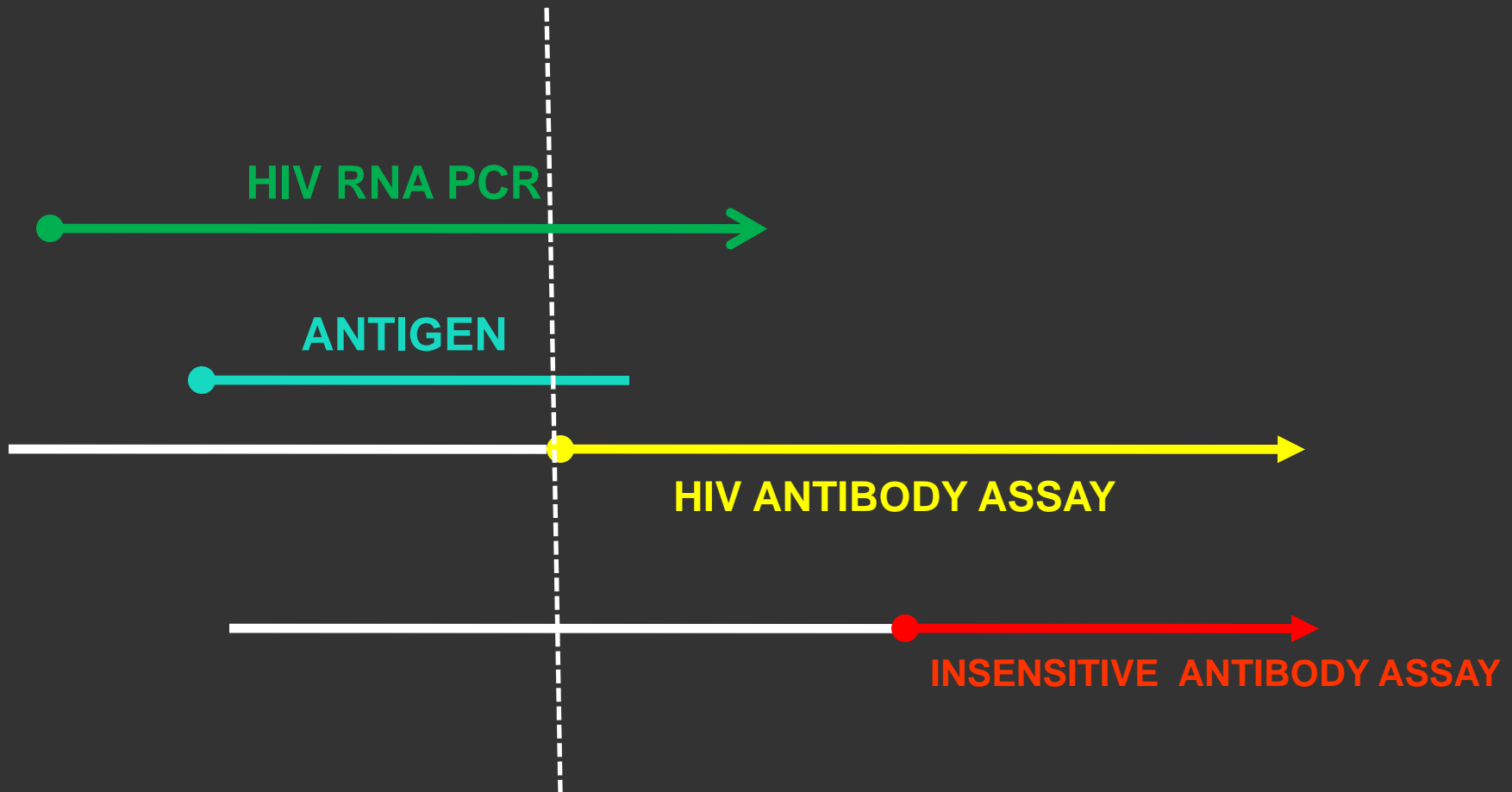
# SOME SUGGESTIONS

# TRIM THE TAILS

WEED OUT ELITE CONTROLLERS, AIDS CASES, ON ART



# RNA PCR /ANTIBODY BIOMARKER





# PERSPECTIVES

- STATISTICAL CONTRIBUTIONS TO MEASURING HIV EPIDEMIC
- INCORPORATE BIOLOGY INTO MODELS
- SMALL CHANGES IN TAILS CAN HAVE BIG EFFECTS!
- ITS NOT ONLY ABOUT A SINGLE NUMBER: TRENDS