

Reducing Decision Errors in the Paired Comparison of the Diagnostic Accuracy of Continuous Screening Tests

Brandy M. Ringham,¹ Todd A. Alonzo,² John T. Brinton,¹
Aarti Munjal,¹ Keith E. Muller,³ Deborah H. Glueck¹

¹Department of Biostatistics and Informatics, University of Colorado Denver

²Department of Preventive Medicine, University of Southern California

³Department of Health Outcomes and Policy, University of Florida

Acknowledgements

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Outline

Science

- Case Study
- Cancer Screening Trial Design
- Cancer Screening Analysis

Statistics

- Bias Correction Algorithm
- Evaluation Studies
- Oral Cancer Screening Demonstration

Oral Cancer Screening Case Study

VISIBLE LIGHT



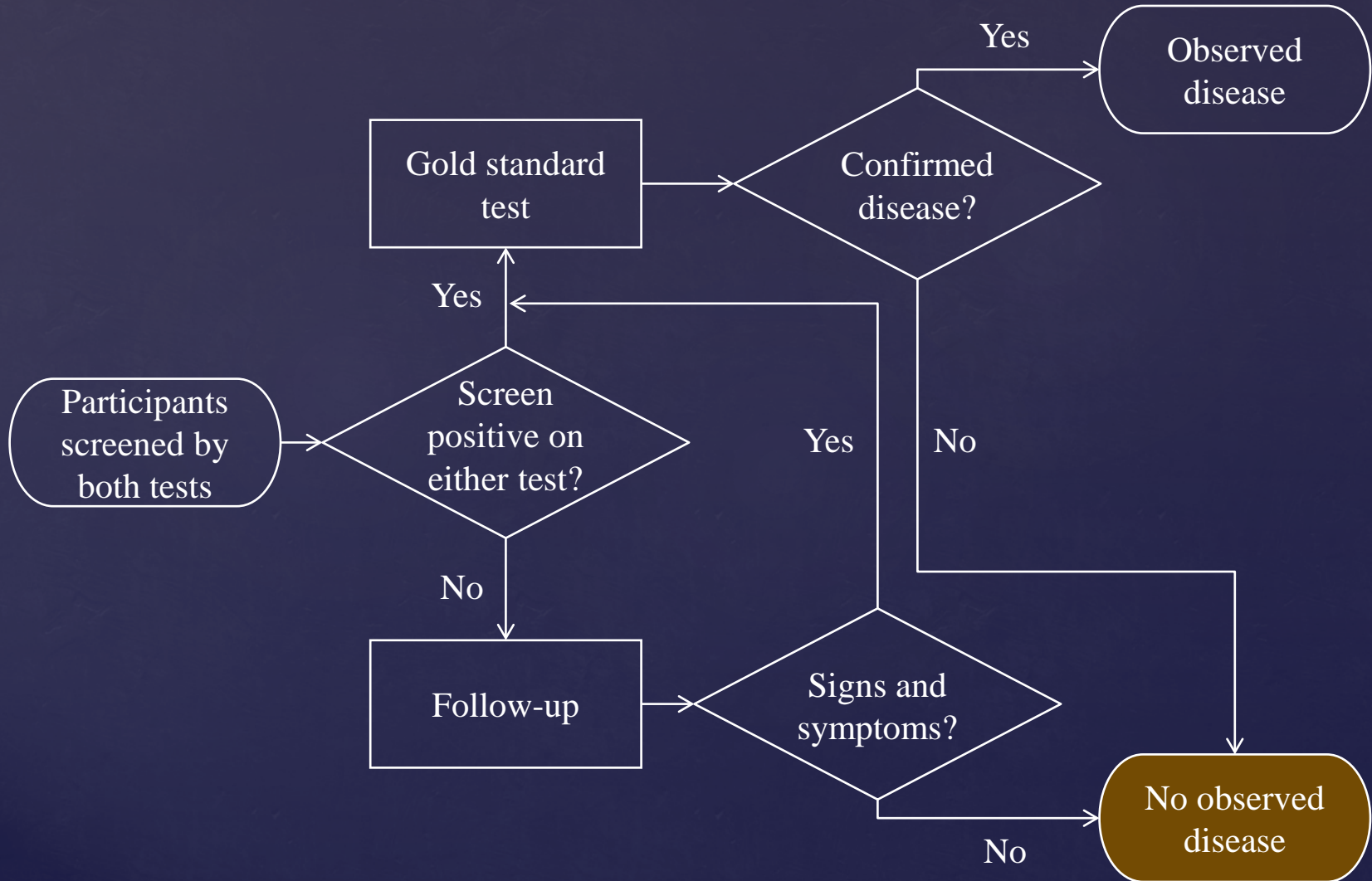
No visible lesion

AUTOFLUORESCENCE

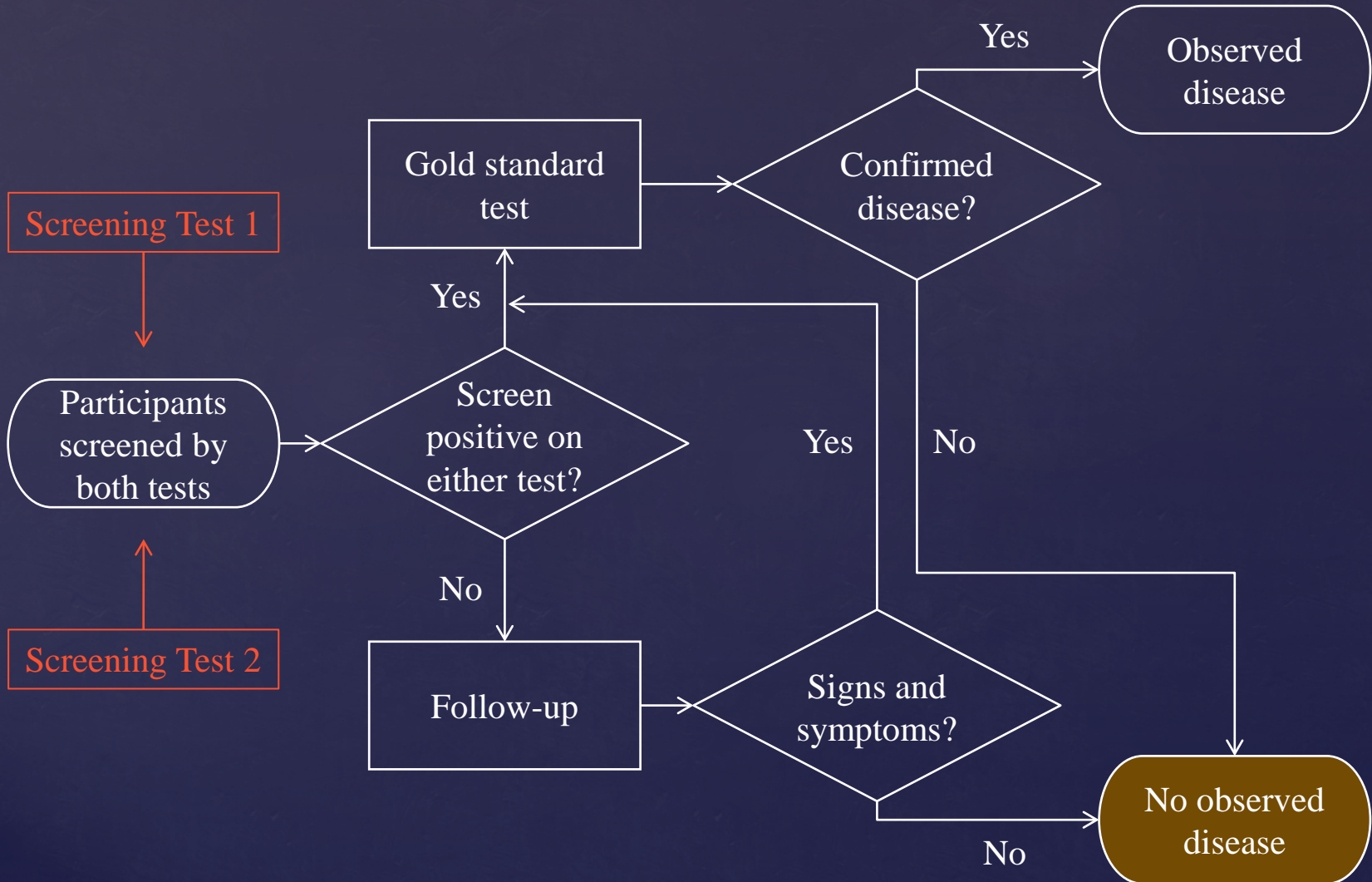


Dark region confirmed to be carcinoma in situ

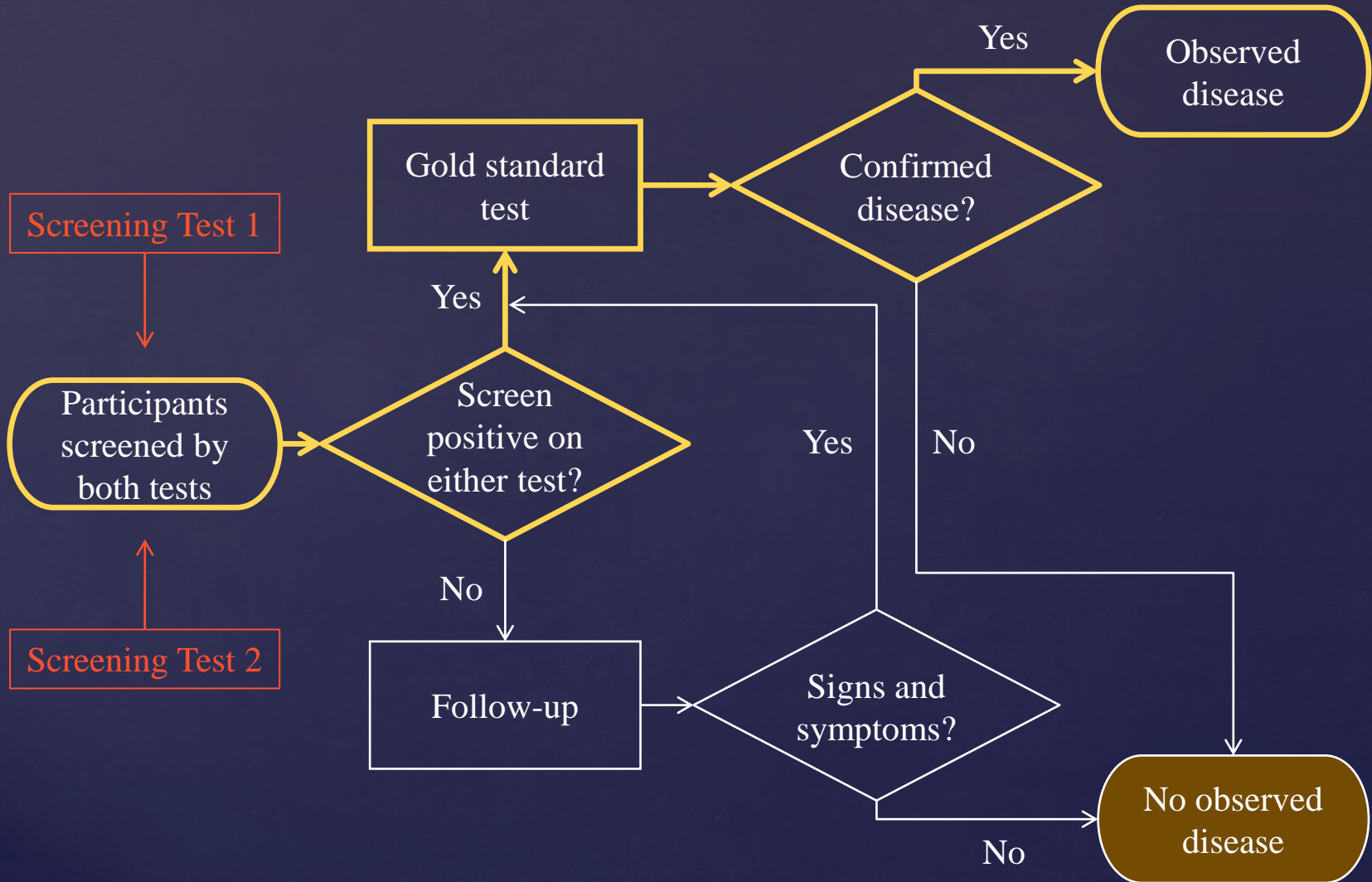
Paired Cancer Screening Trial



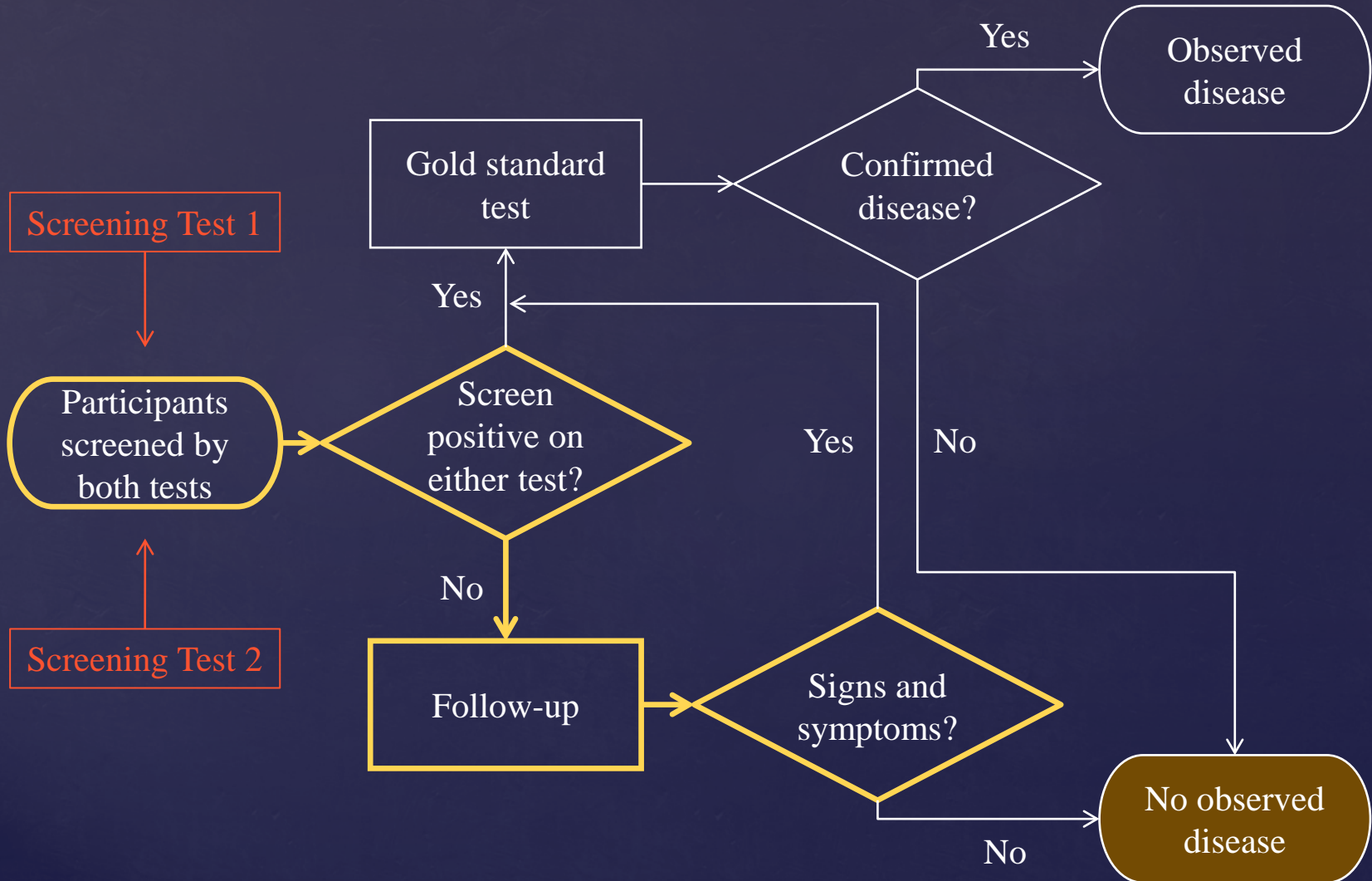
Paired Cancer Screening Trial



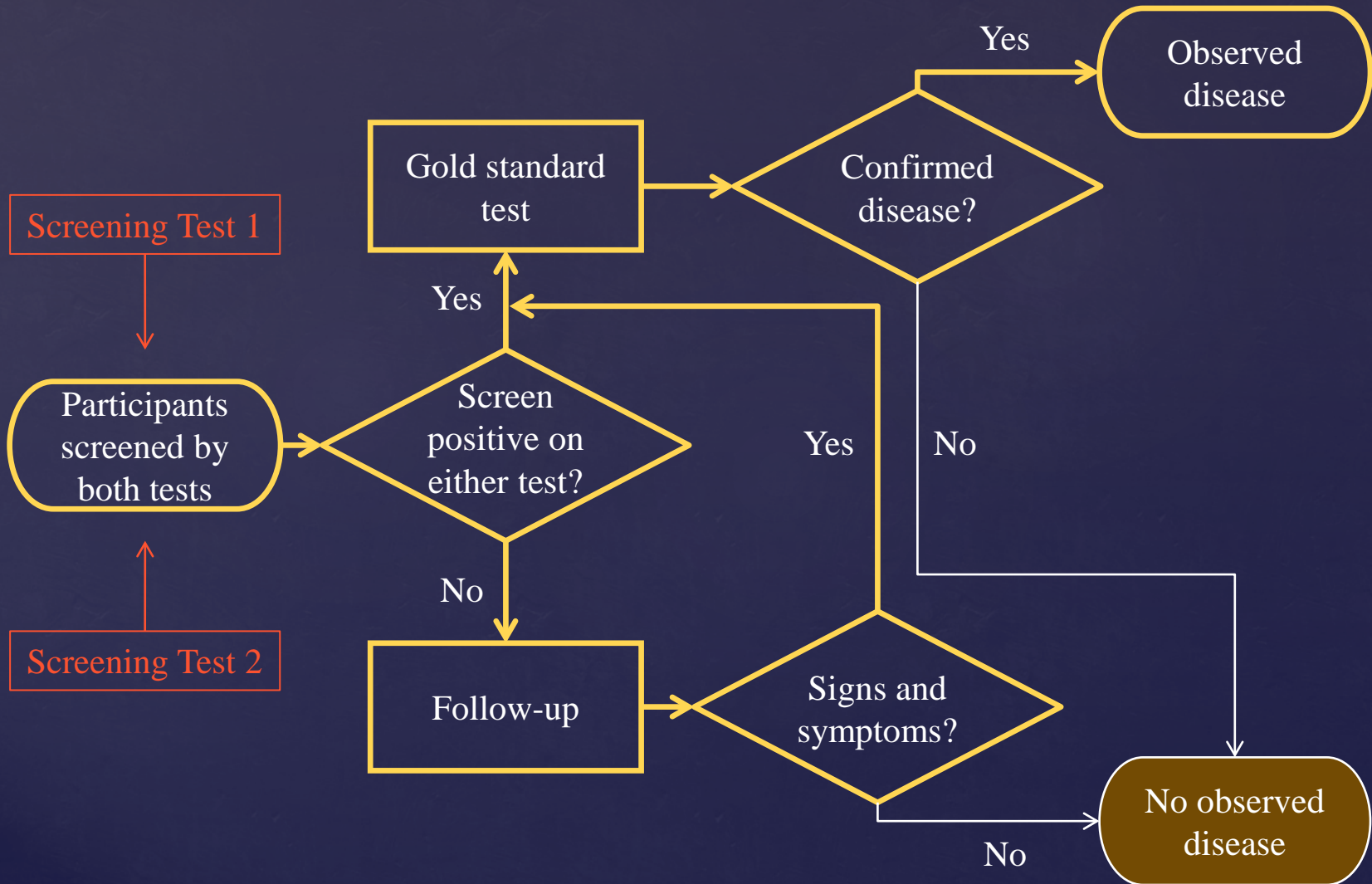
Paired Cancer Screening Trial



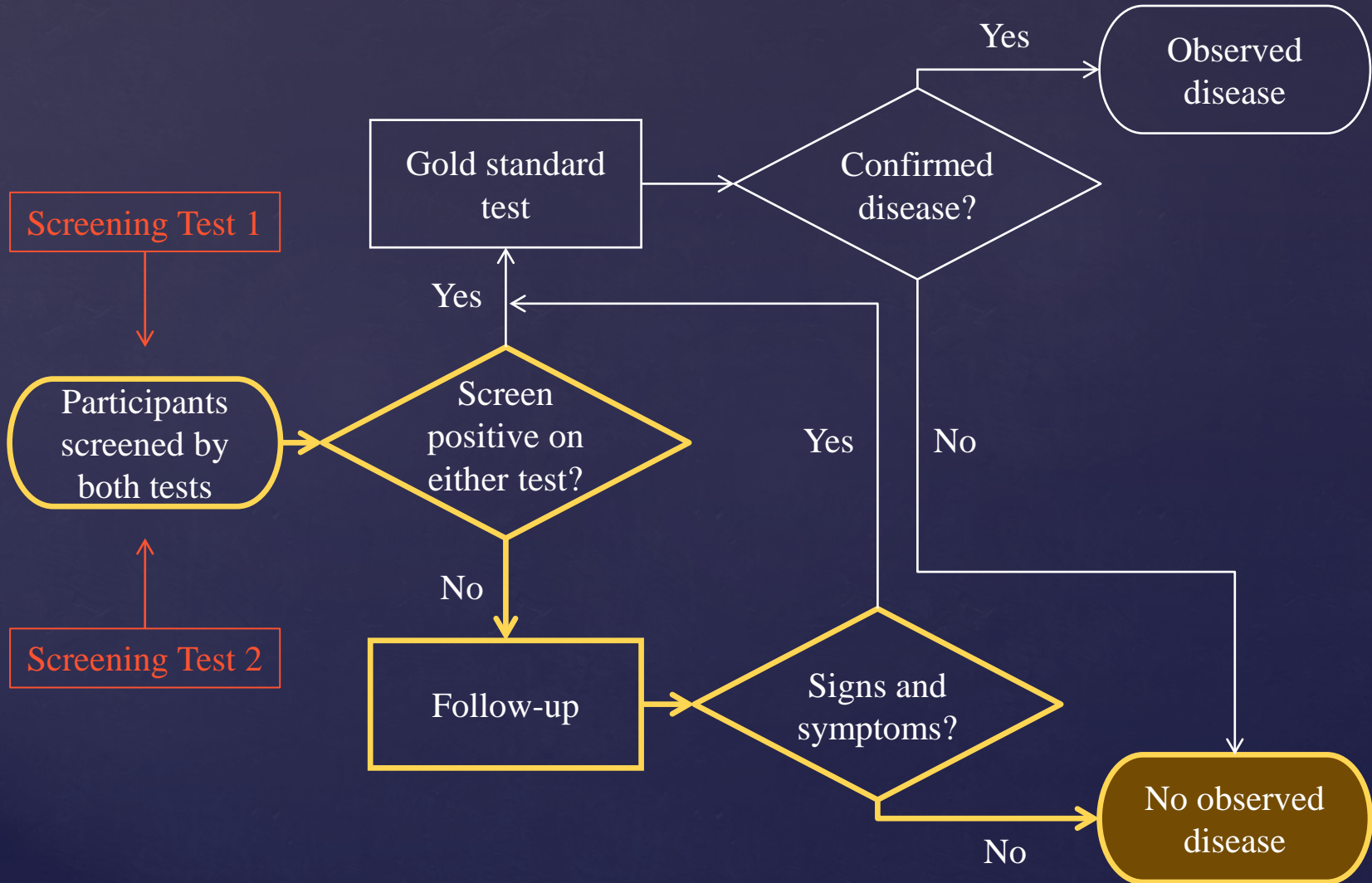
Paired Cancer Screening Trial



Paired Cancer Screening Trial

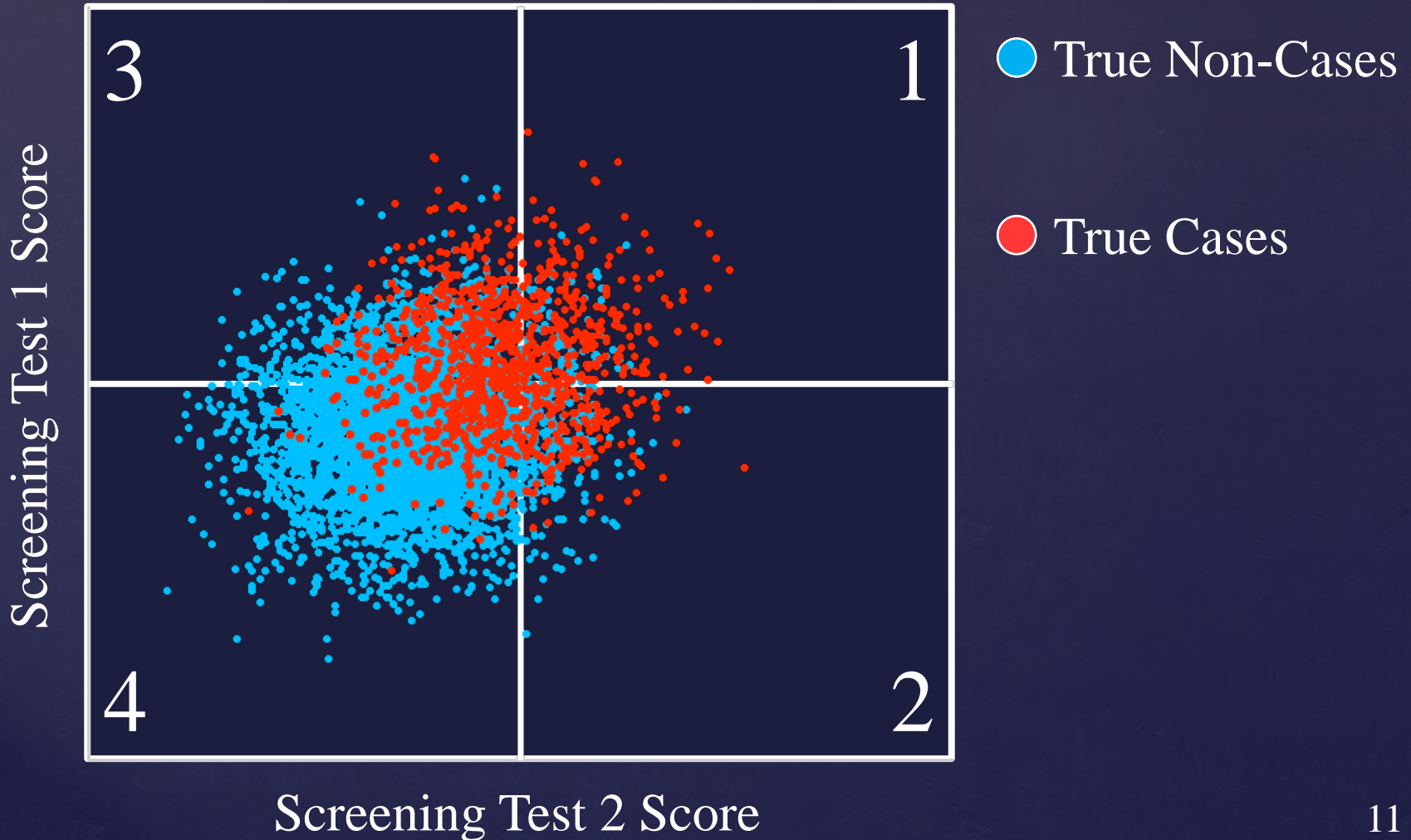


Paired Cancer Screening Trial



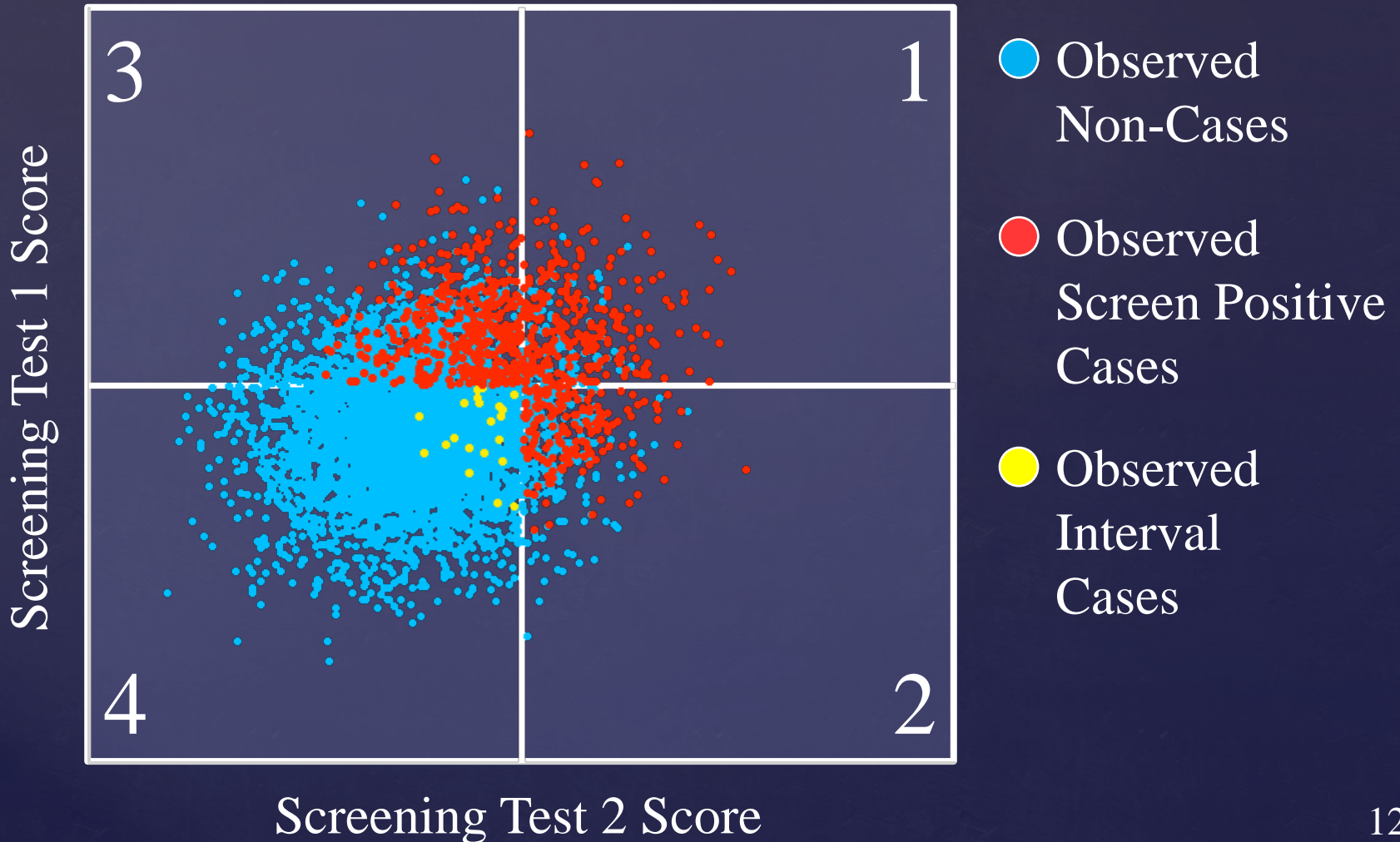
Hypothetical Cancer Screening Data

Omniscient Viewpoint



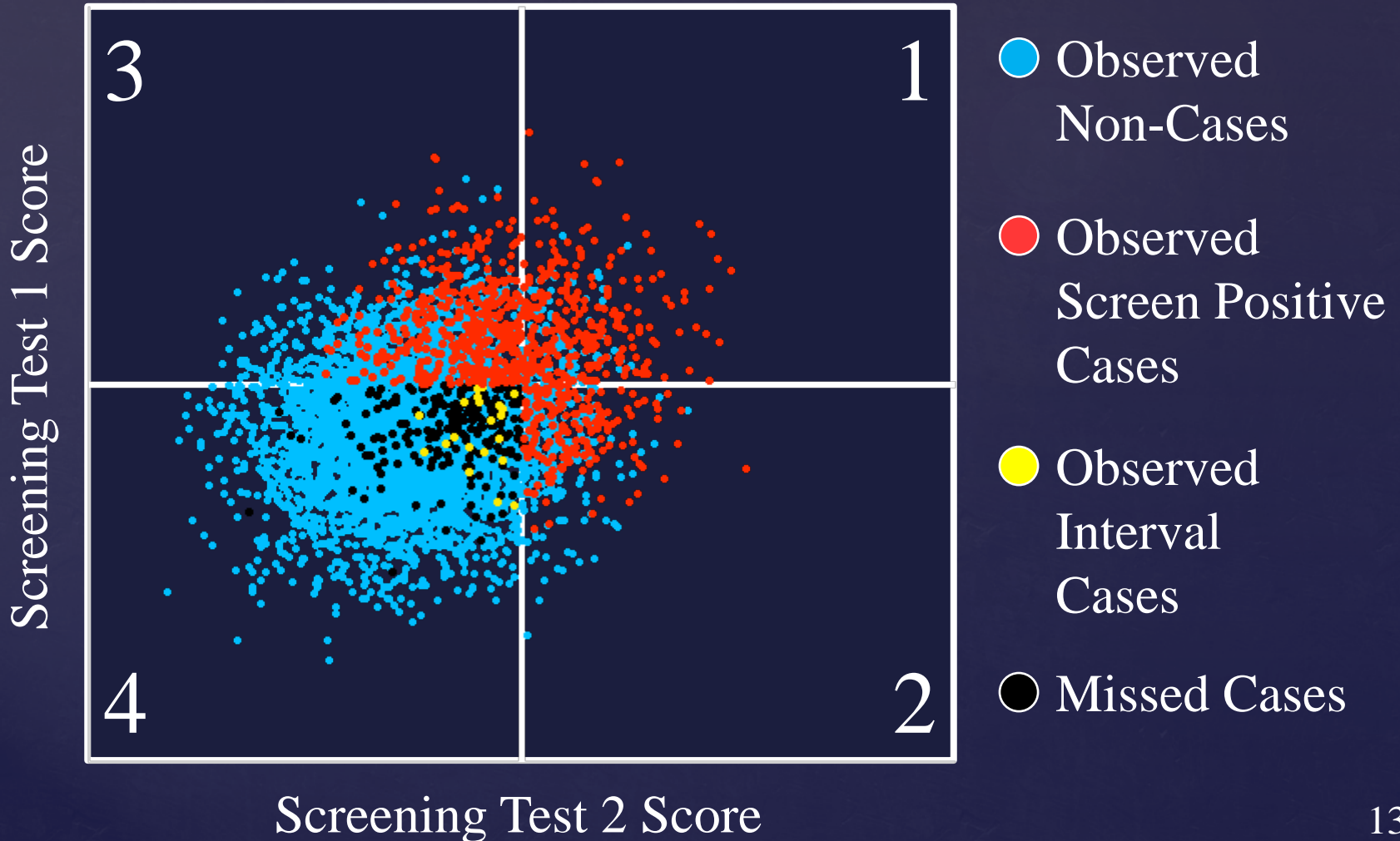
Hypothetical Cancer Screening Data

Study Investigator's Viewpoint



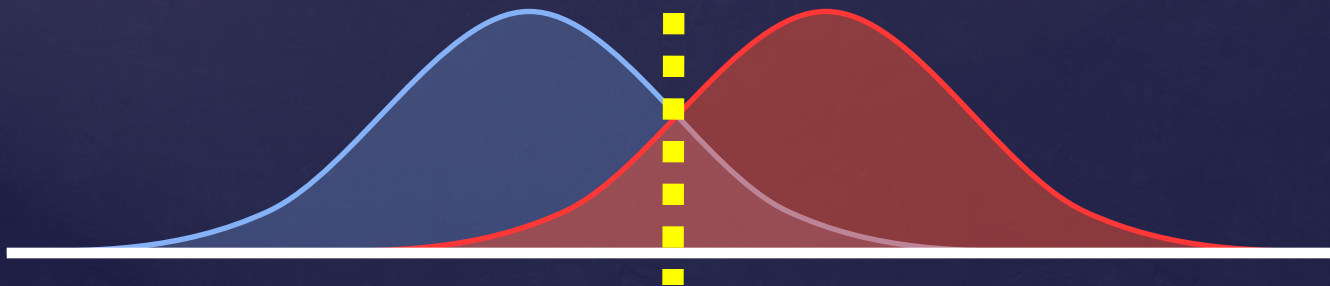
Hypothetical Cancer Screening Data

Omniscient Viewpoint



Analysis of Cancer Screening Studies

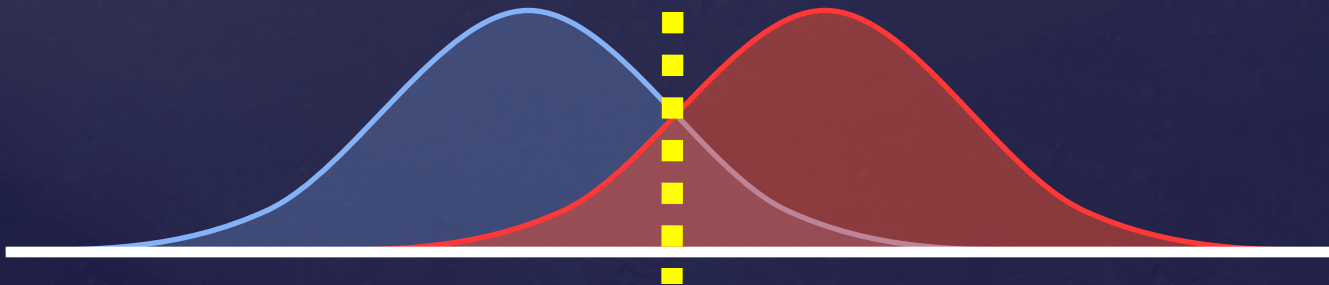
		Disease Status	
		+	-
Screening Test	+	a	b
	-	c	d



Analysis of Cancer Screening Studies

		Disease Status	
		+	-
Screening Test	+	a	b
	-	c	d

$$\text{Sensitivity} = \frac{a}{a+c}$$

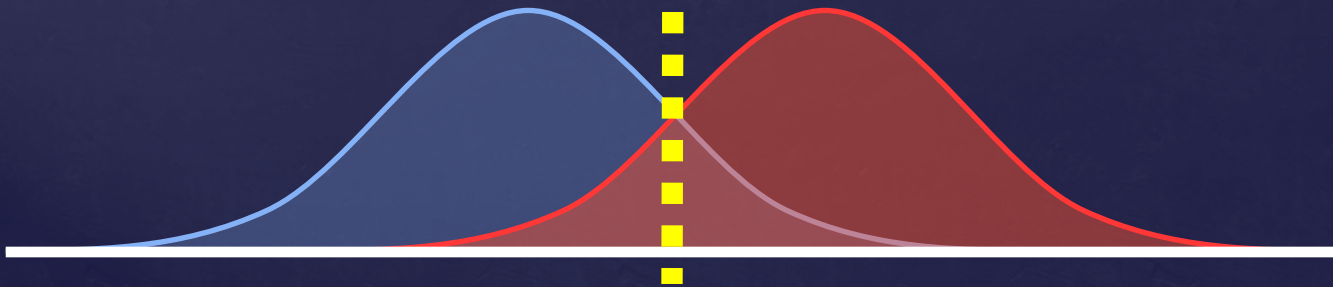


Analysis of Cancer Screening Studies

		Disease Status	
		+	-
Screening Test	+	a	b
	-	c	d

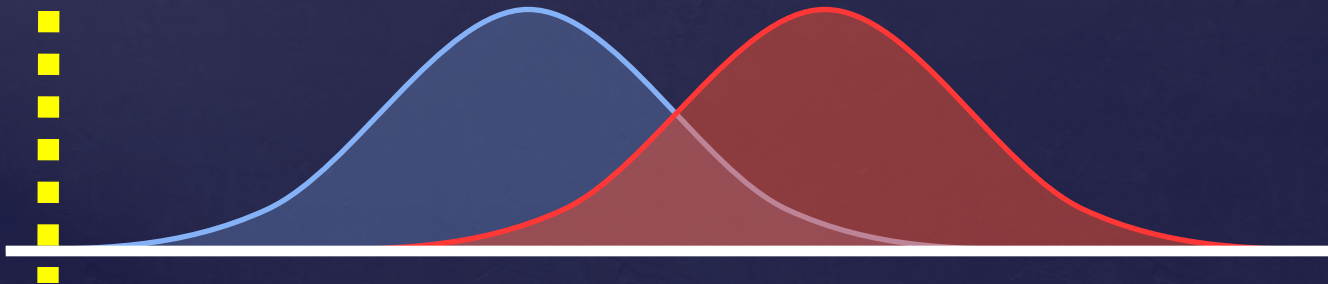
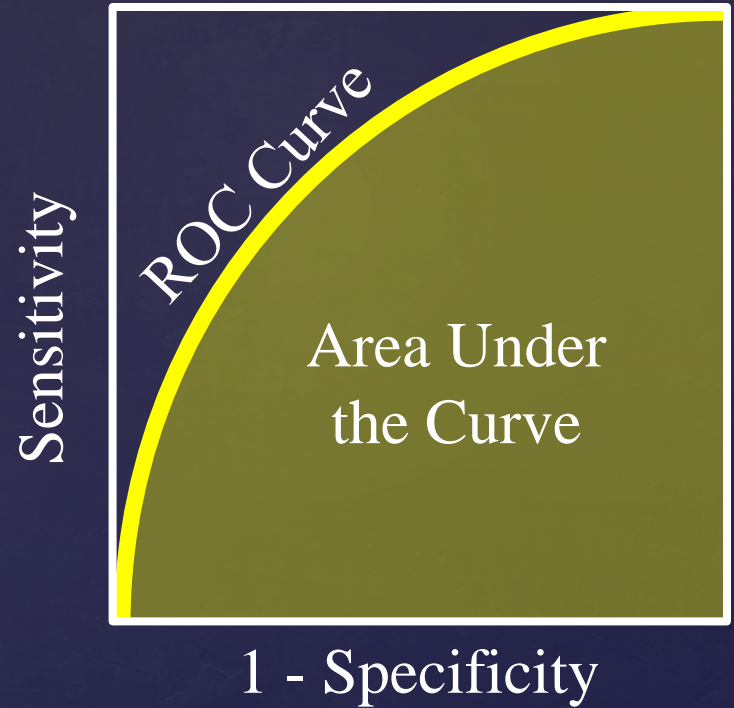
$$\text{Sensitivity} = \frac{a}{a+c}$$

$$\text{Specificity} = \frac{d}{b+d}$$

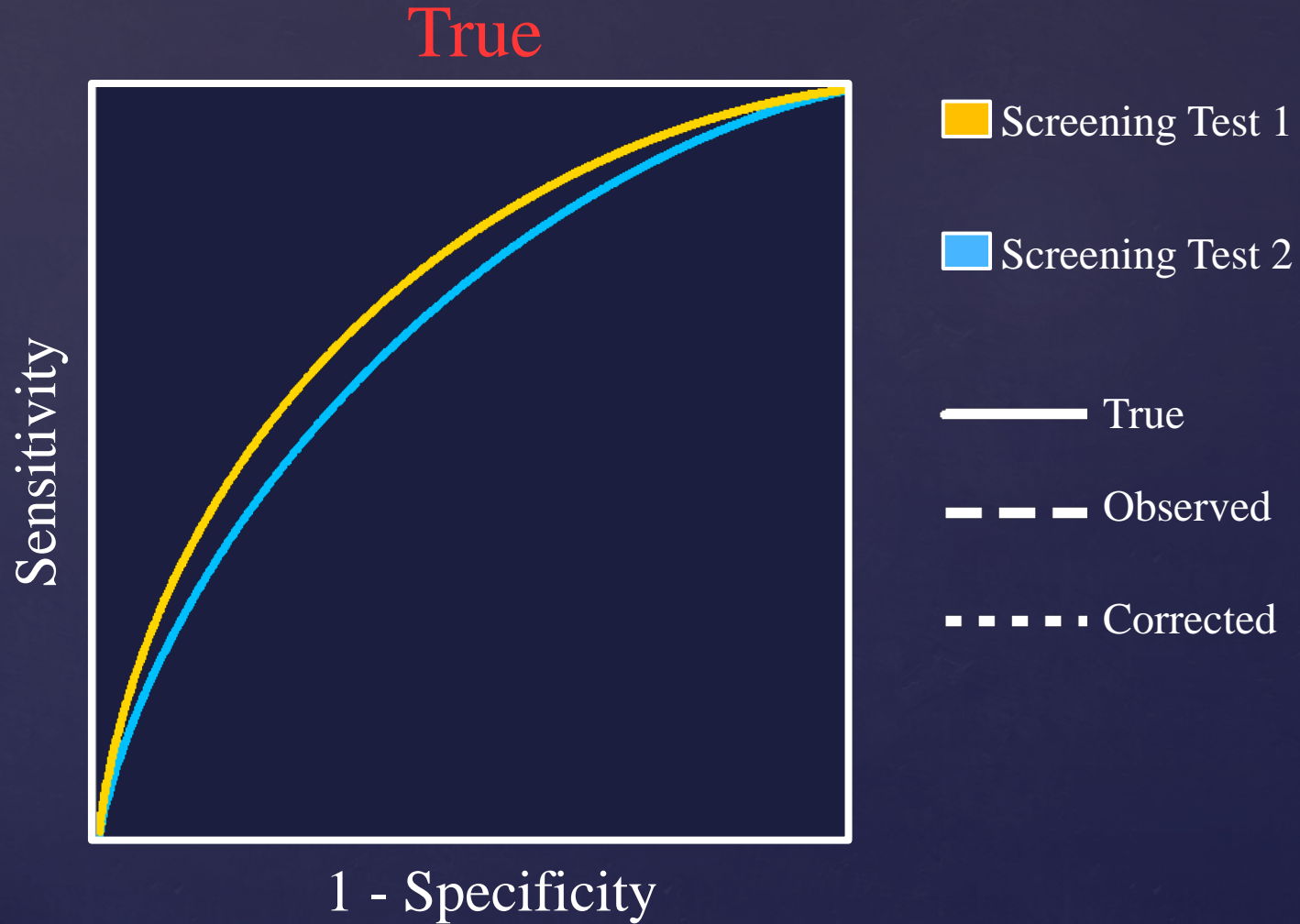


Analysis of Cancer Screening Studies

		Disease Status	
		+	-
Screening Test	+	a	b
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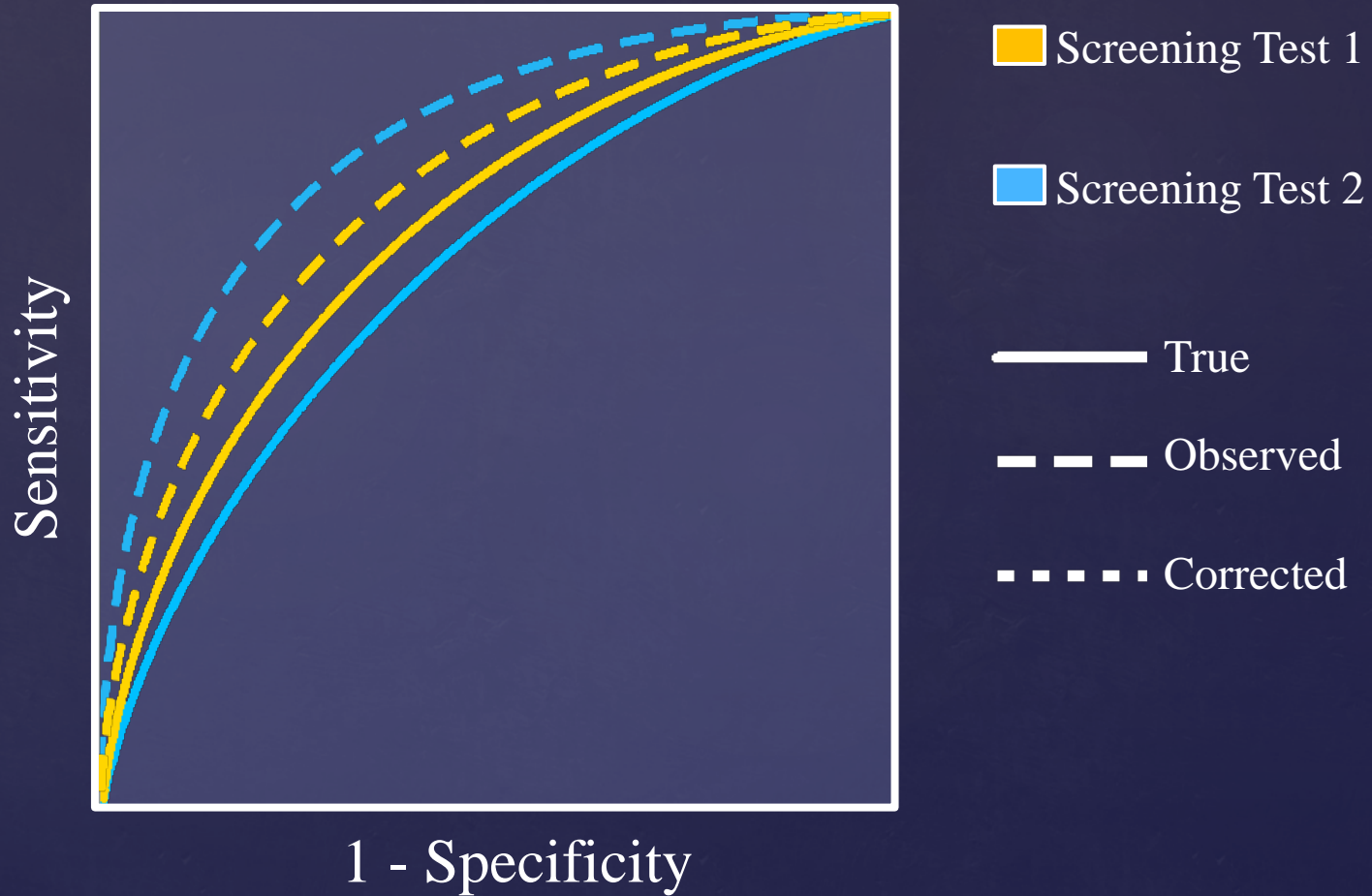


Comparing Two Continuous Screening Tests



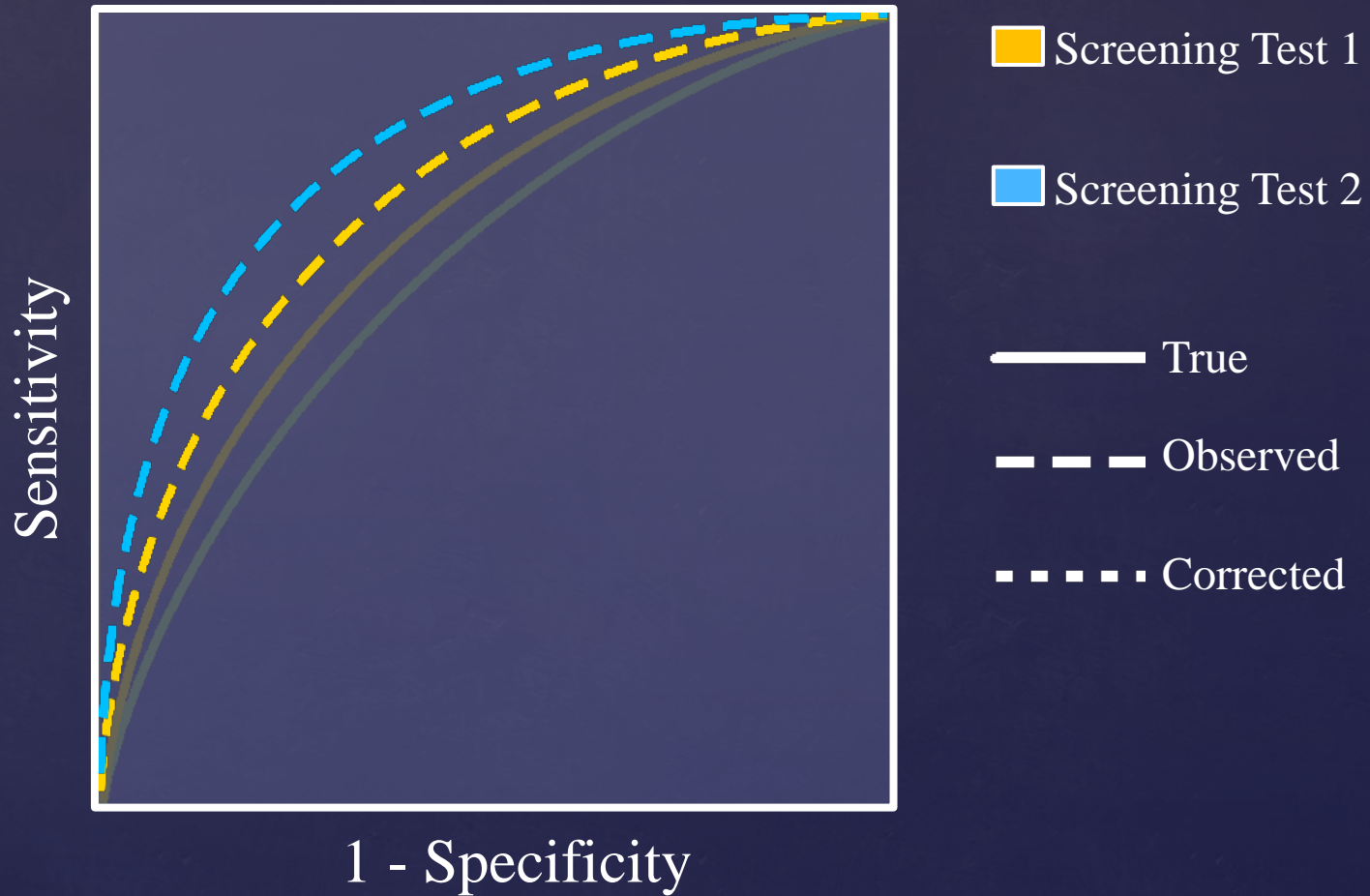
Comparing Two Continuous Screening Tests

Observed



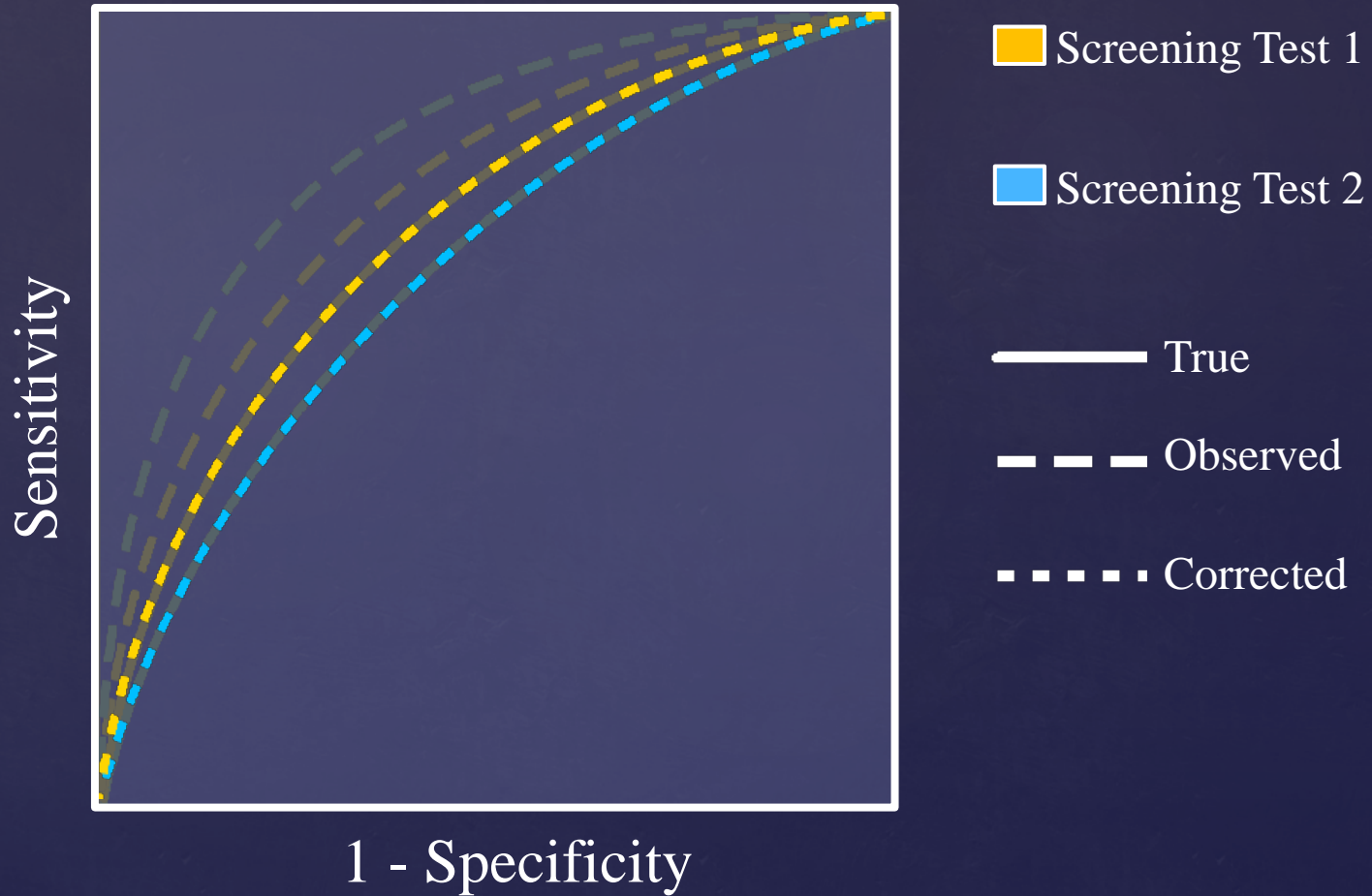
Comparing Two Continuous Screening Tests

Observed



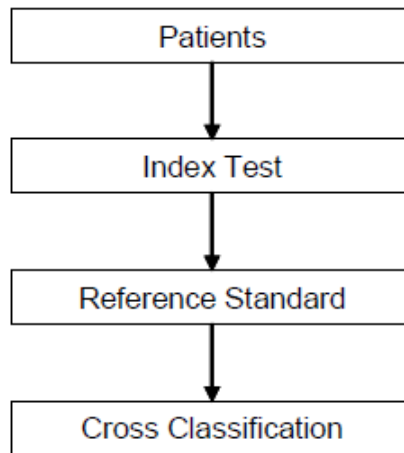
Comparing Two Continuous Screening Tests

Corrected

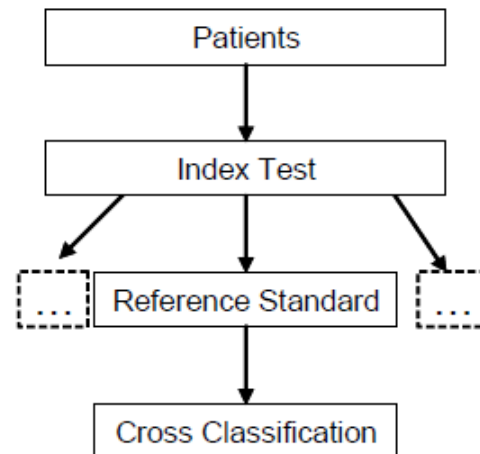


Types of Bias

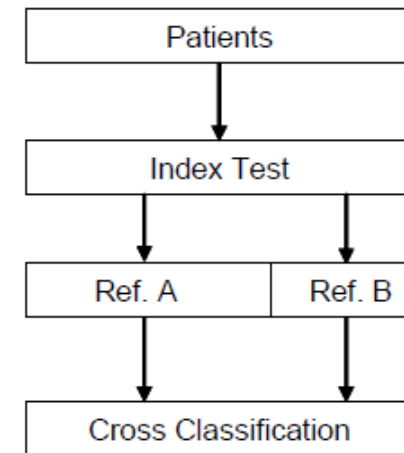
A: Classic Design



B: Partial Verification



C: Differential Verification

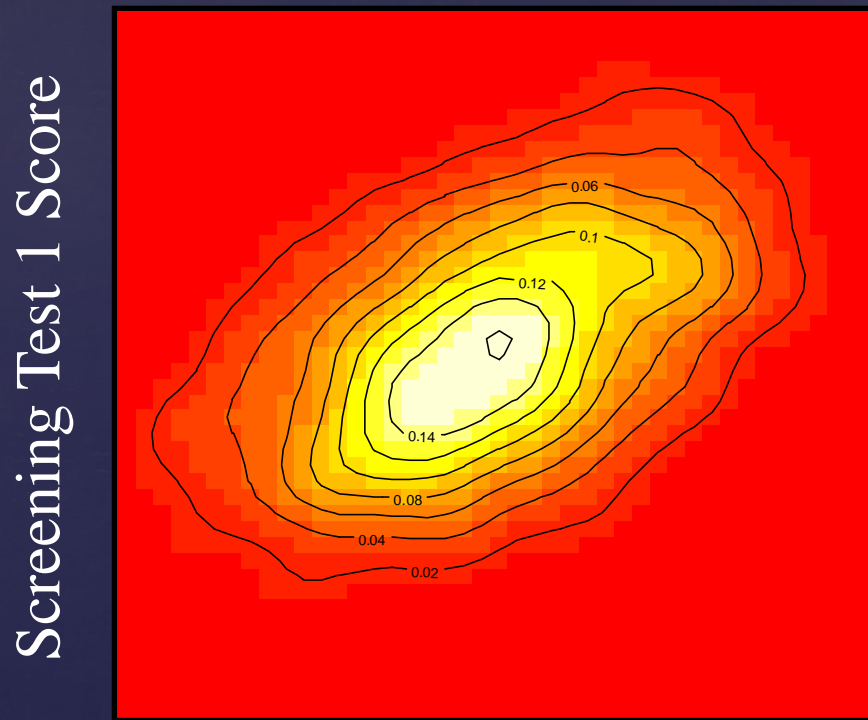


Bias Correction Algorithm

1. Find the maximum likelihood estimates of the parameters of the bivariate Gaussian distribution of test scores for the cases.
2. Use the maximum likelihood estimates and the sampling fractions in each partition to calculate weighted estimates.

Bias Correction Algorithm

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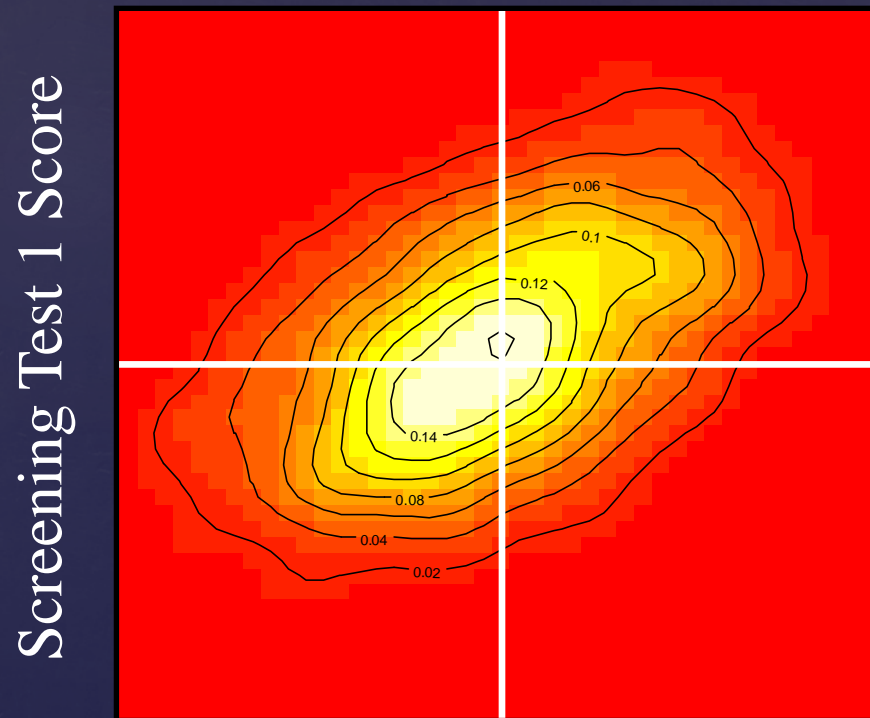


(Nath, 1971)

Screening Test 2 Score

Bias Correction Algorithm

1. Find the maximum likelihood estimates of the parameters of the bivariate Gaussian distribution of test scores for the cases.

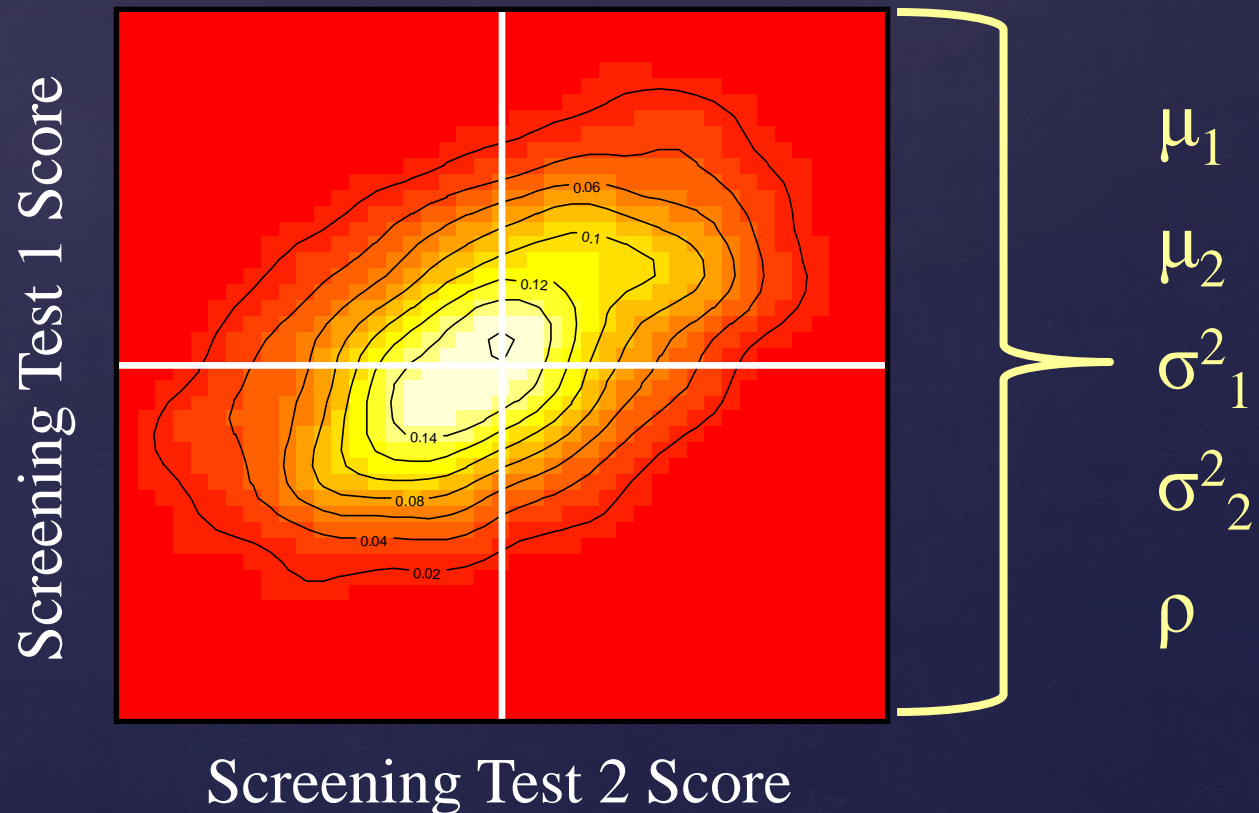


(Nath, 1971)

Screening Test 2 Score

Bias Correction Algorithm

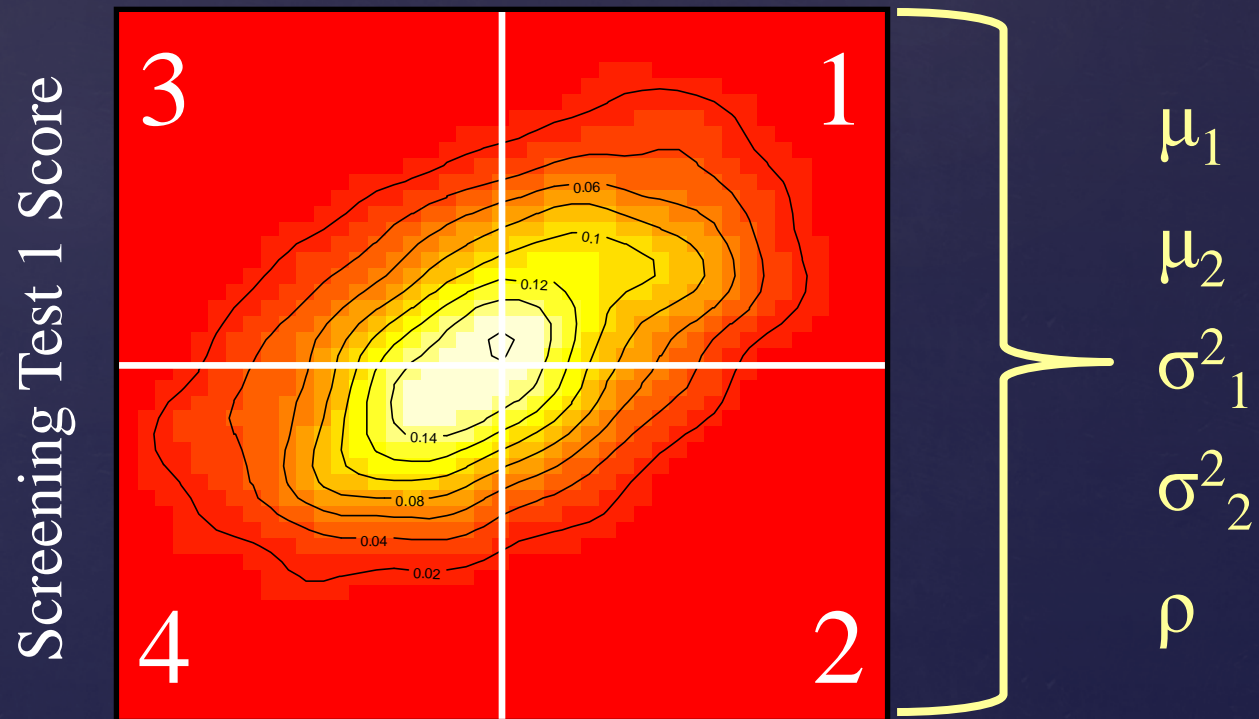
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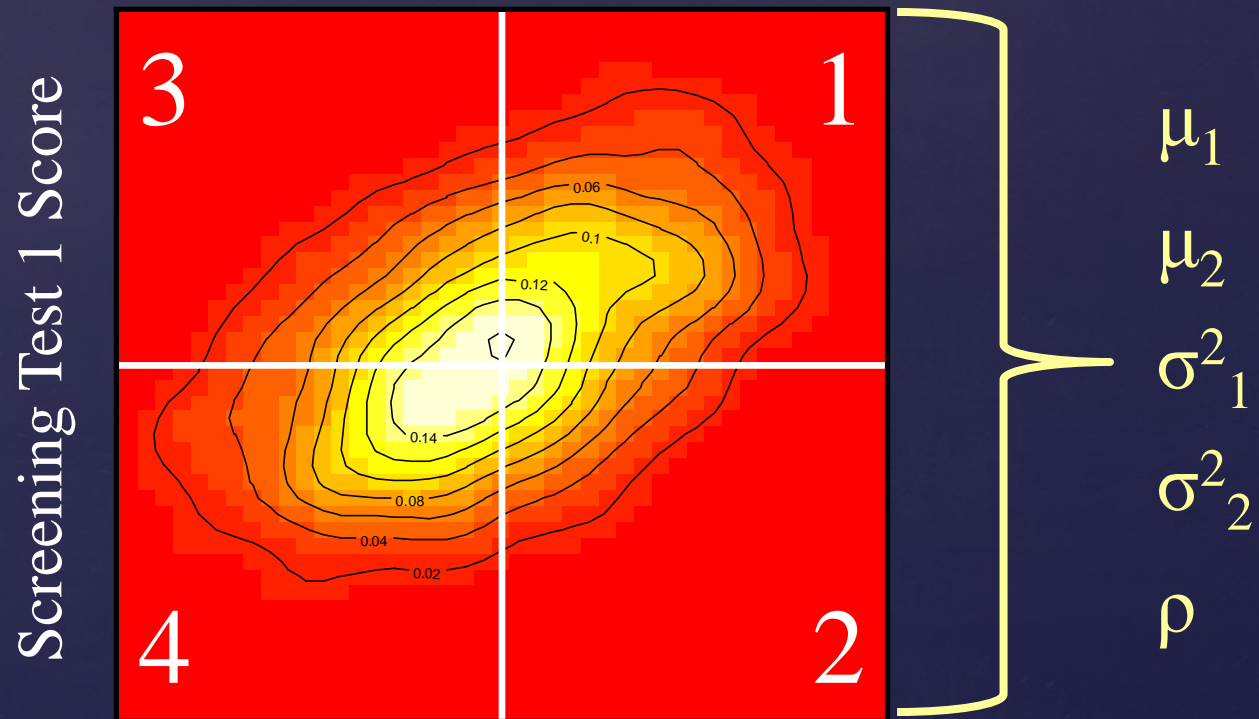


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Screening Test 2 Score

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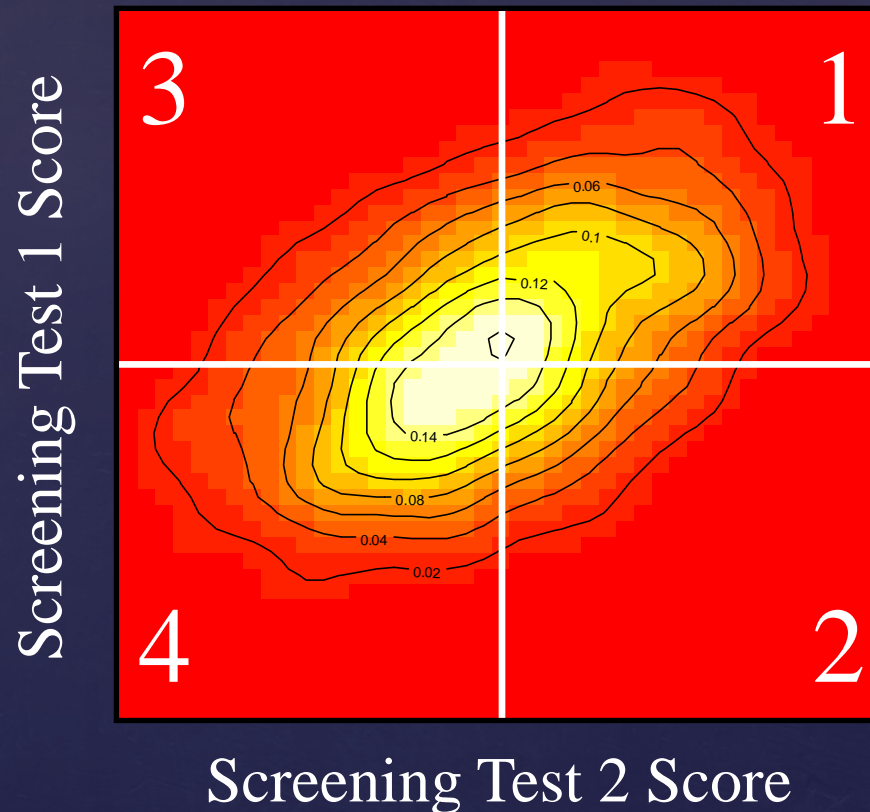
Screening Test 2 Score

Bias Correction Algorithm

2. Use the maximum likelihood estimates and the sampling fractions in each partition to calculate weighted estimates.

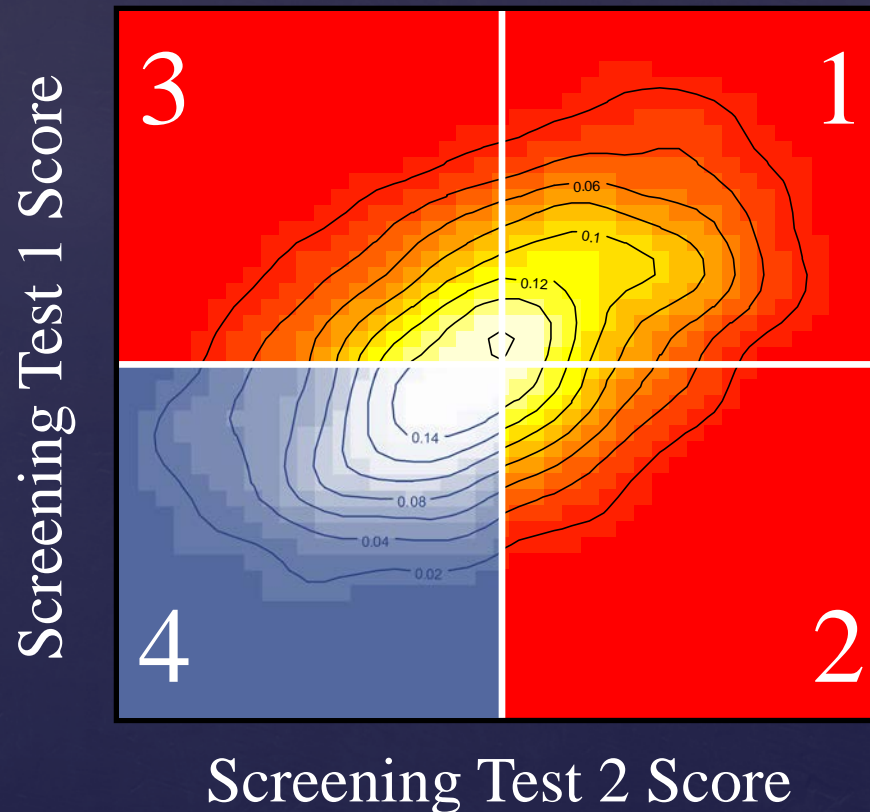
Bias Correction Algorithm

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Bias Correction Algorithm

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Weighted Estimates

$$\hat{\lambda} = 1 - \Phi \left(\frac{a_1 - \hat{\mu}_{11, N}}{\hat{\sigma}_{11, N}}, \frac{a_2 - \hat{\mu}_{21, N}}{\hat{\sigma}_{21, N}}, \hat{\rho}_{1, N} \right)$$

Weighted Estimates

$$\hat{\lambda} = 1 - \Phi \left(\frac{a_1 - \hat{\mu}_{11, N}}{\hat{\sigma}_{11, N}}, \frac{a_2 - \hat{\mu}_{21, N}}{\hat{\sigma}_{21, N}}, \hat{\rho}_{1, N} \right)$$

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Weighted Estimates

$$\hat{\mu}_{11, W} = \hat{\lambda} \bar{X}_{11, A} + (1 - \hat{\lambda}) \bar{X}_{11, B}$$

$$\hat{\mu}_{21, W} = \hat{\lambda} \bar{X}_{21, A} + (1 - \hat{\lambda}) \bar{X}_{21, B}$$

$$\hat{\sigma}_{11, W}^2 = \mathcal{G}_1 + \mathcal{H}_1 - \hat{\mu}_{11, W}^2$$

$$\hat{\sigma}_{21, W}^2 = \mathcal{G}_2 + \mathcal{H}_2 - \hat{\mu}_{21, W}^2$$

$$\hat{\rho}_{1, W} = \hat{\sigma}_{11, W}^2 \hat{\sigma}_{21, W}^2 (\mathcal{P} + \mathcal{Q} - \hat{\mu}_{11, W} \hat{\mu}_{21, W})$$

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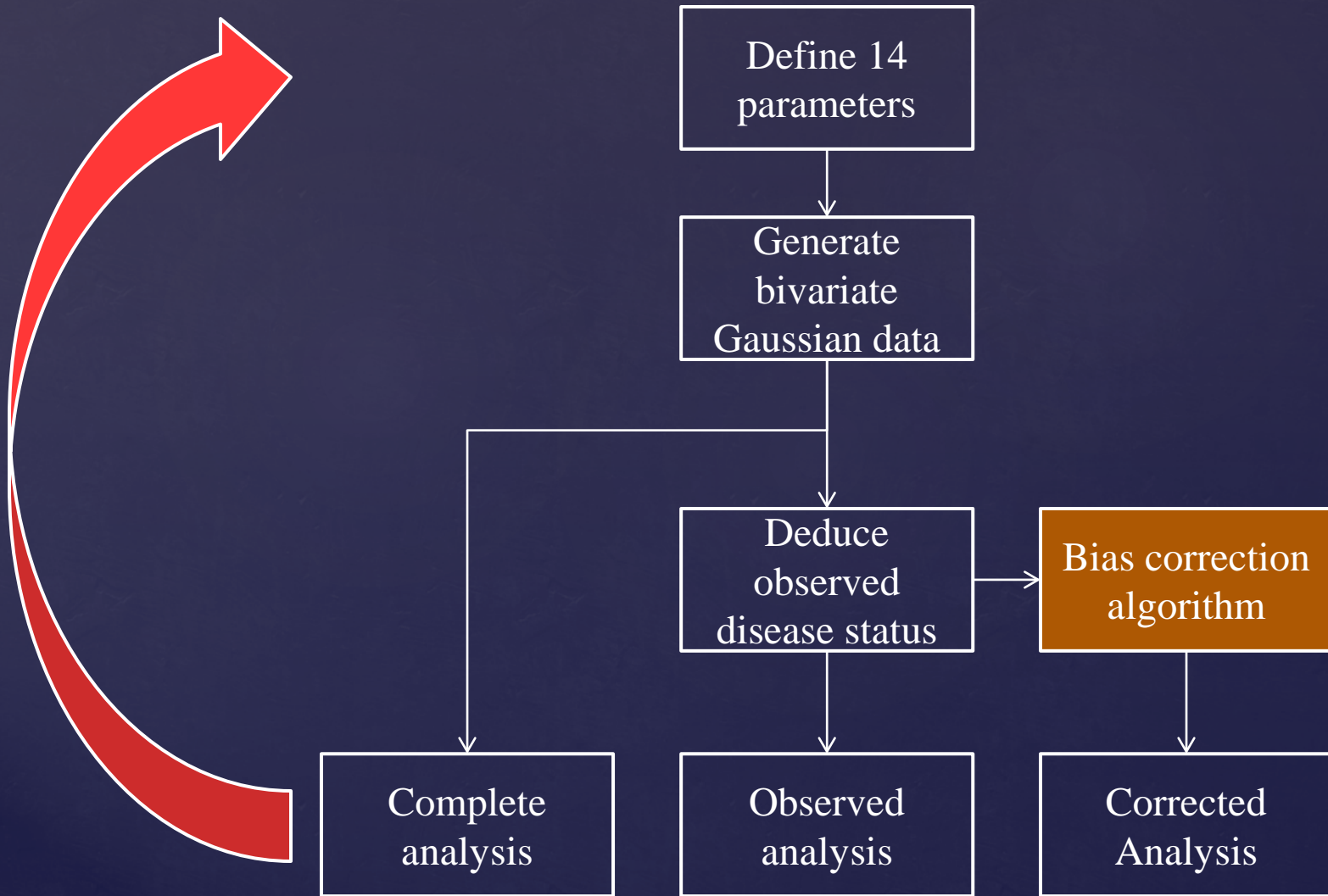
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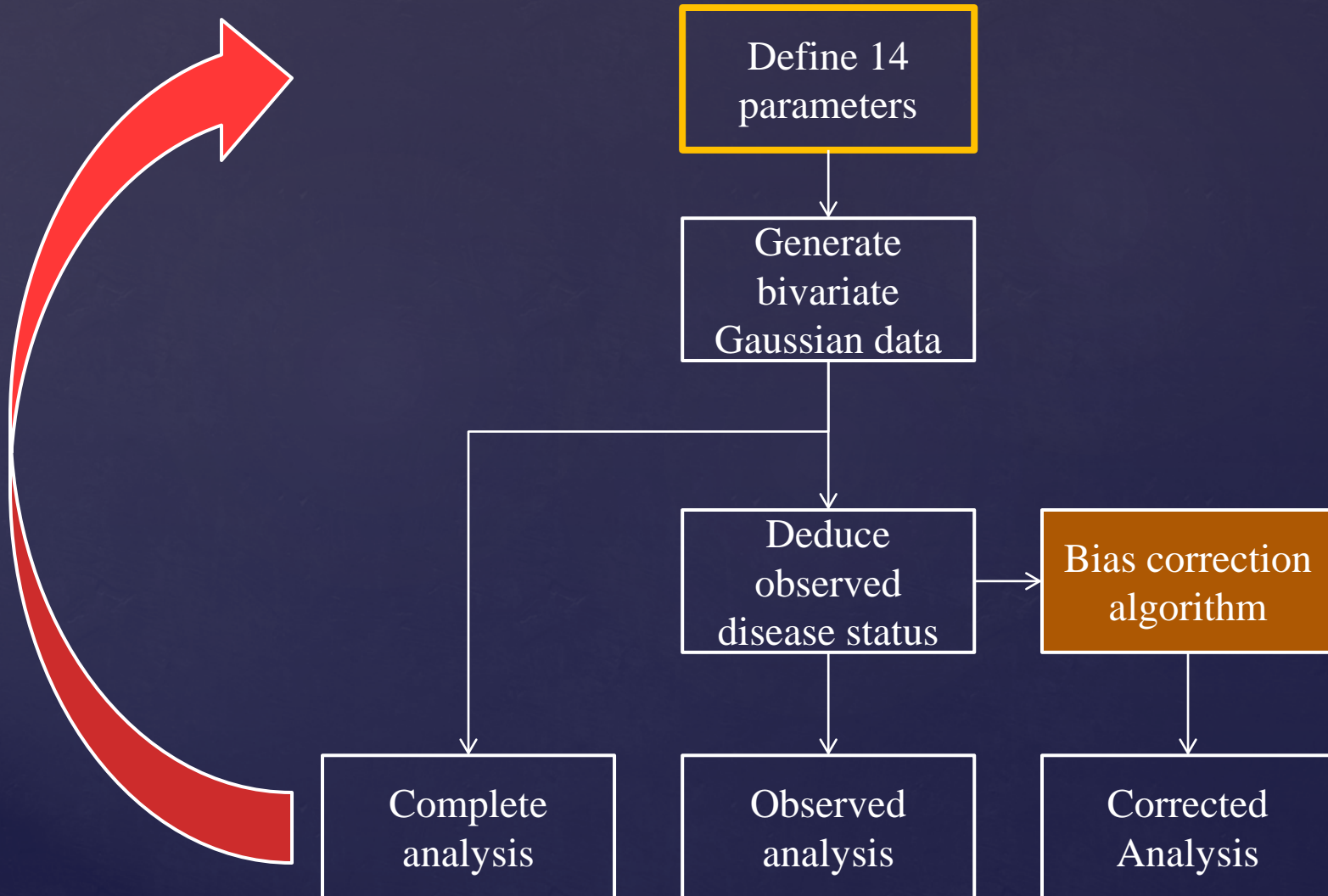
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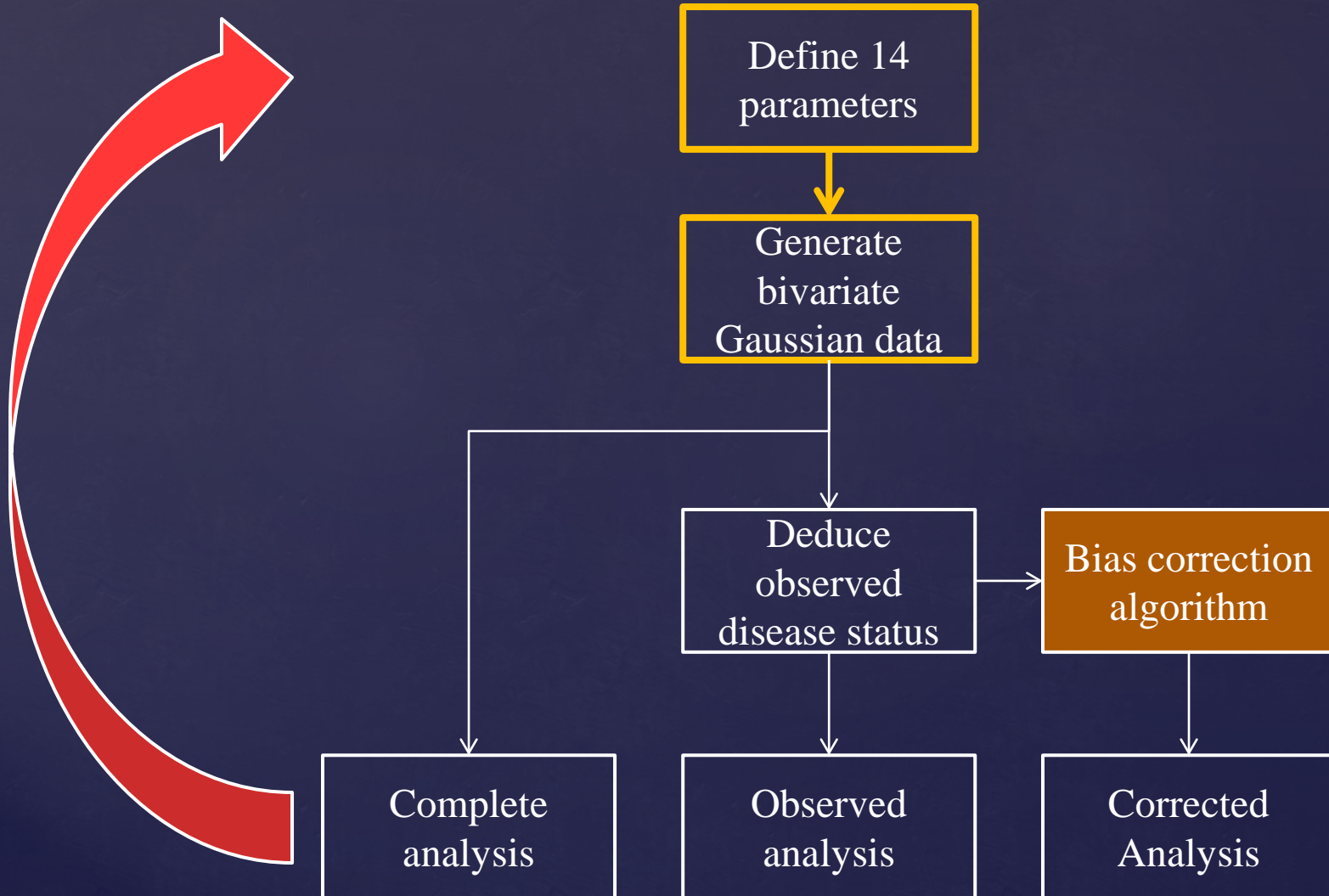
Design for Simulation Studies



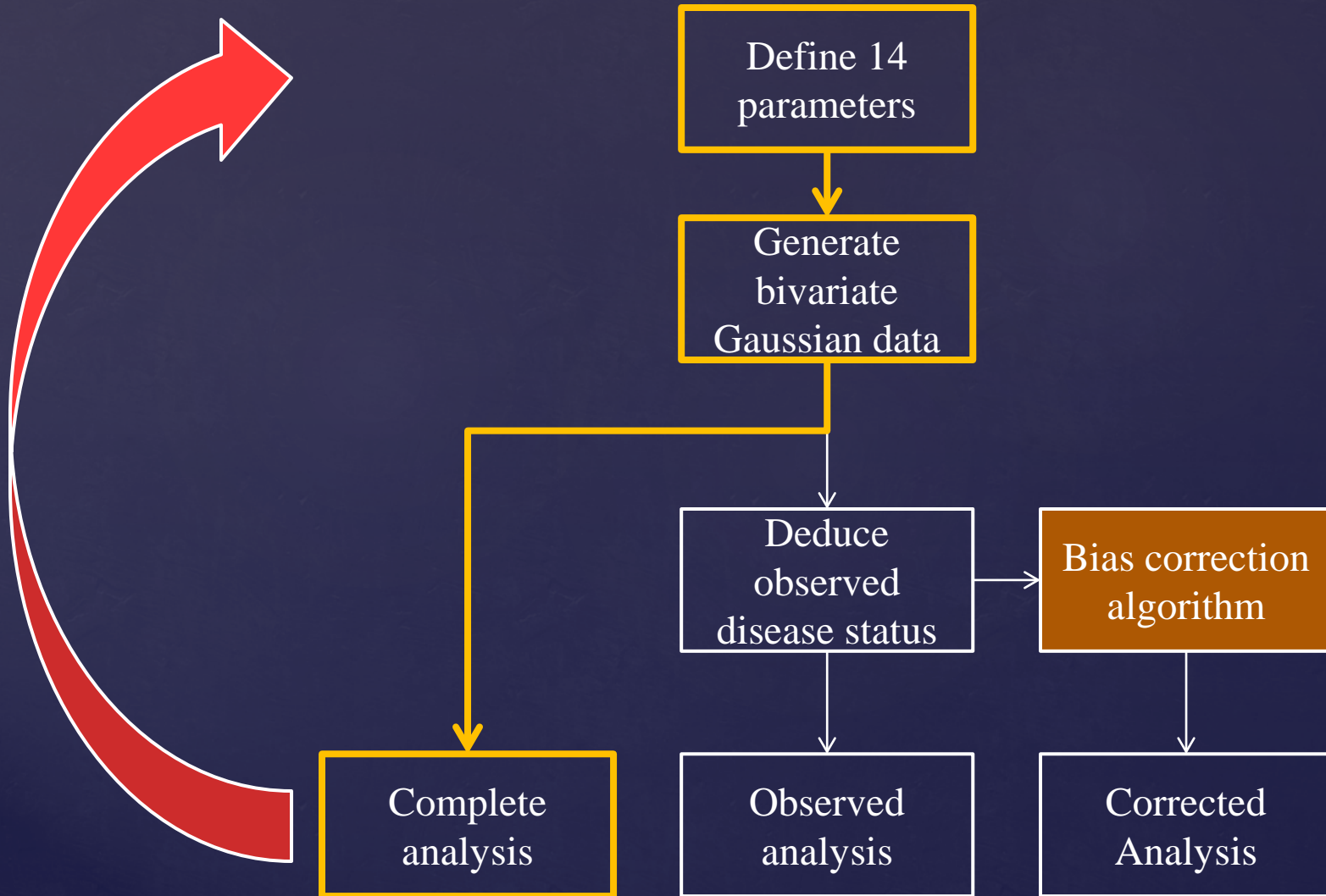
Design for Simulation Studies



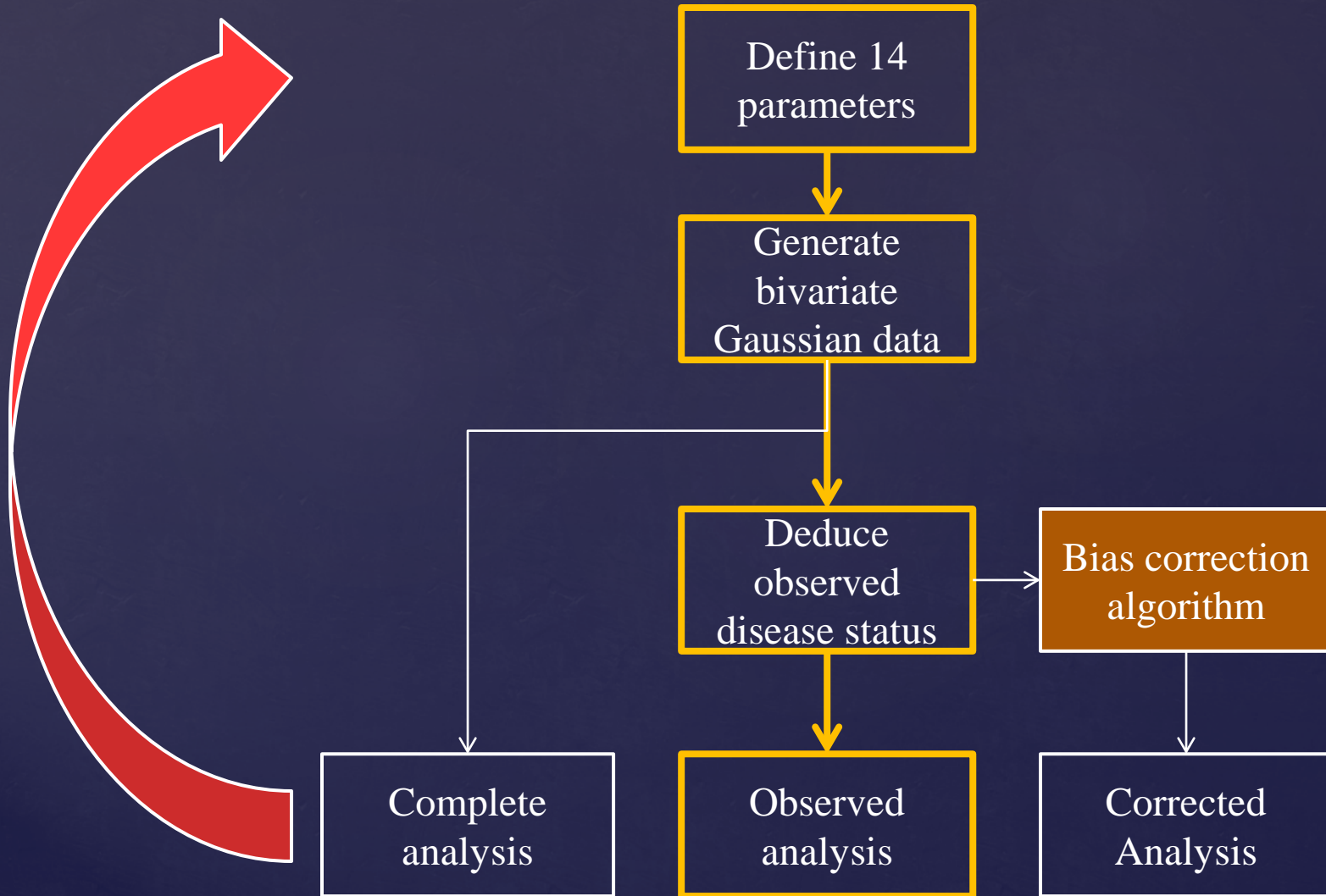
Design for Simulation Studies



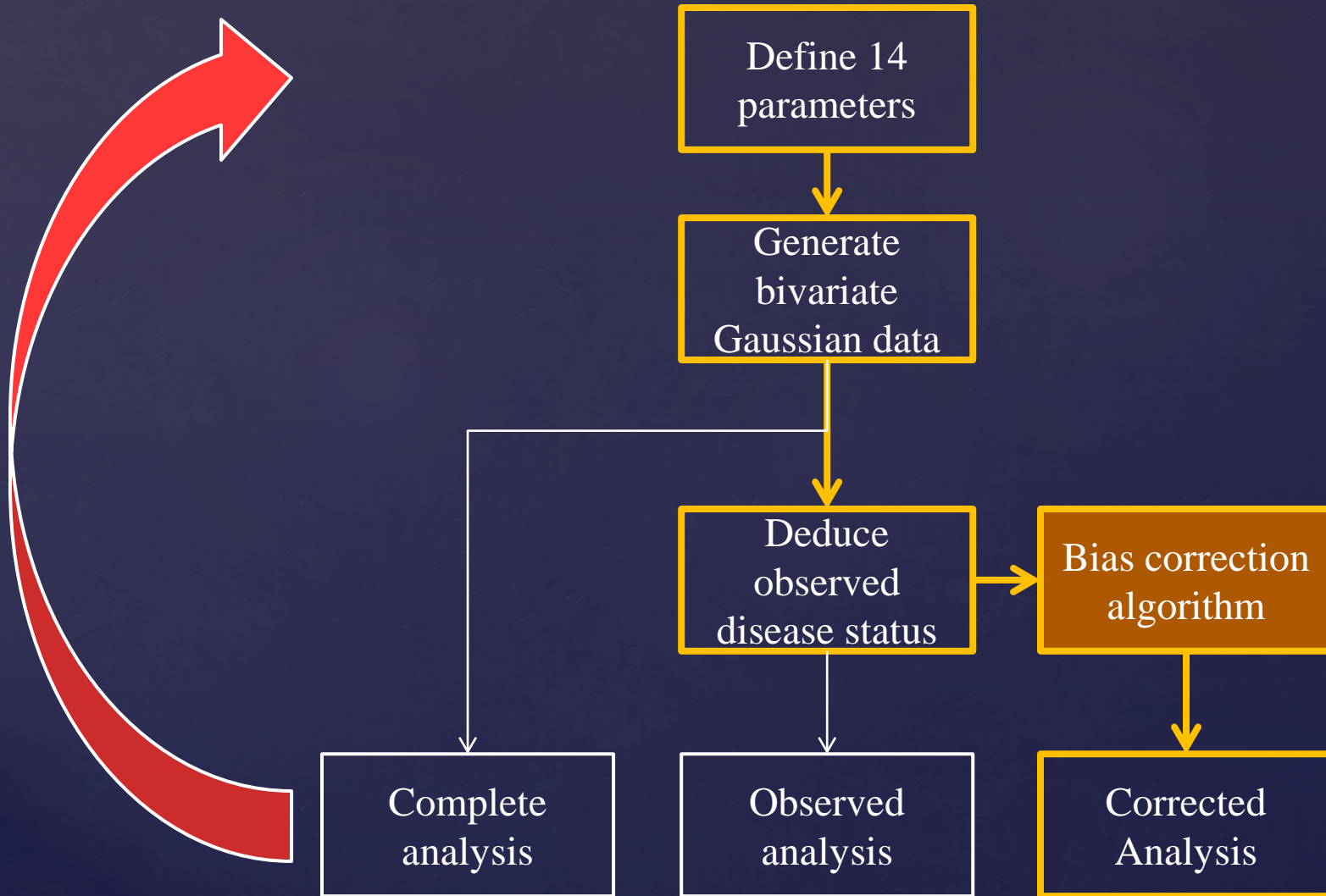
Design for Simulation Studies



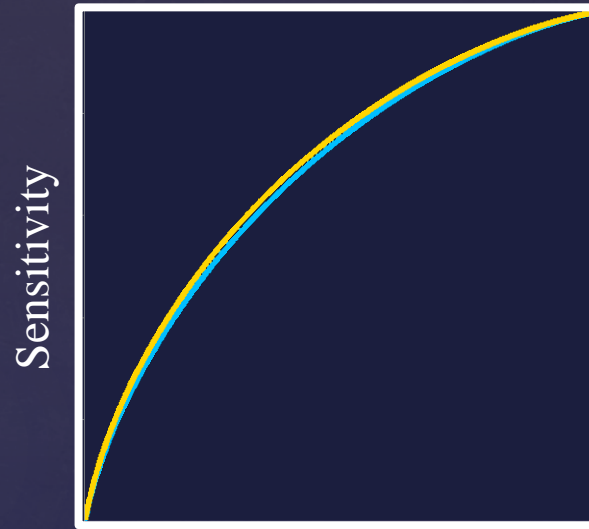
Design for Simulation Studies



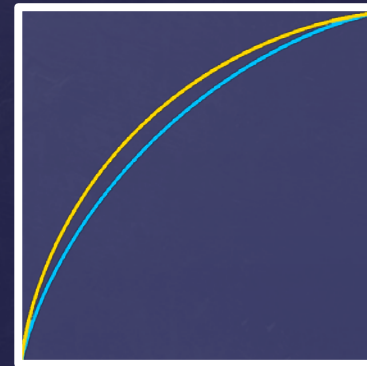
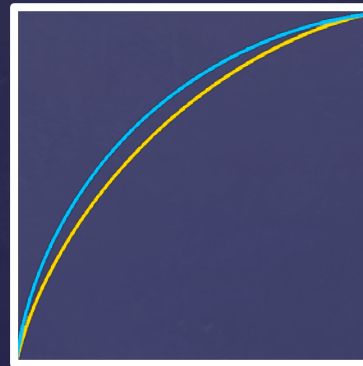
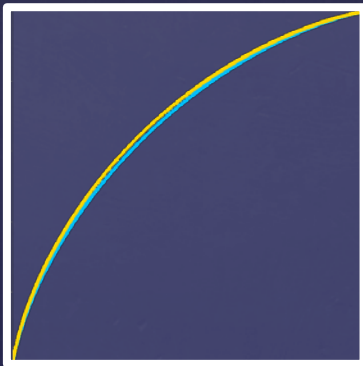
Design for Simulation Studies



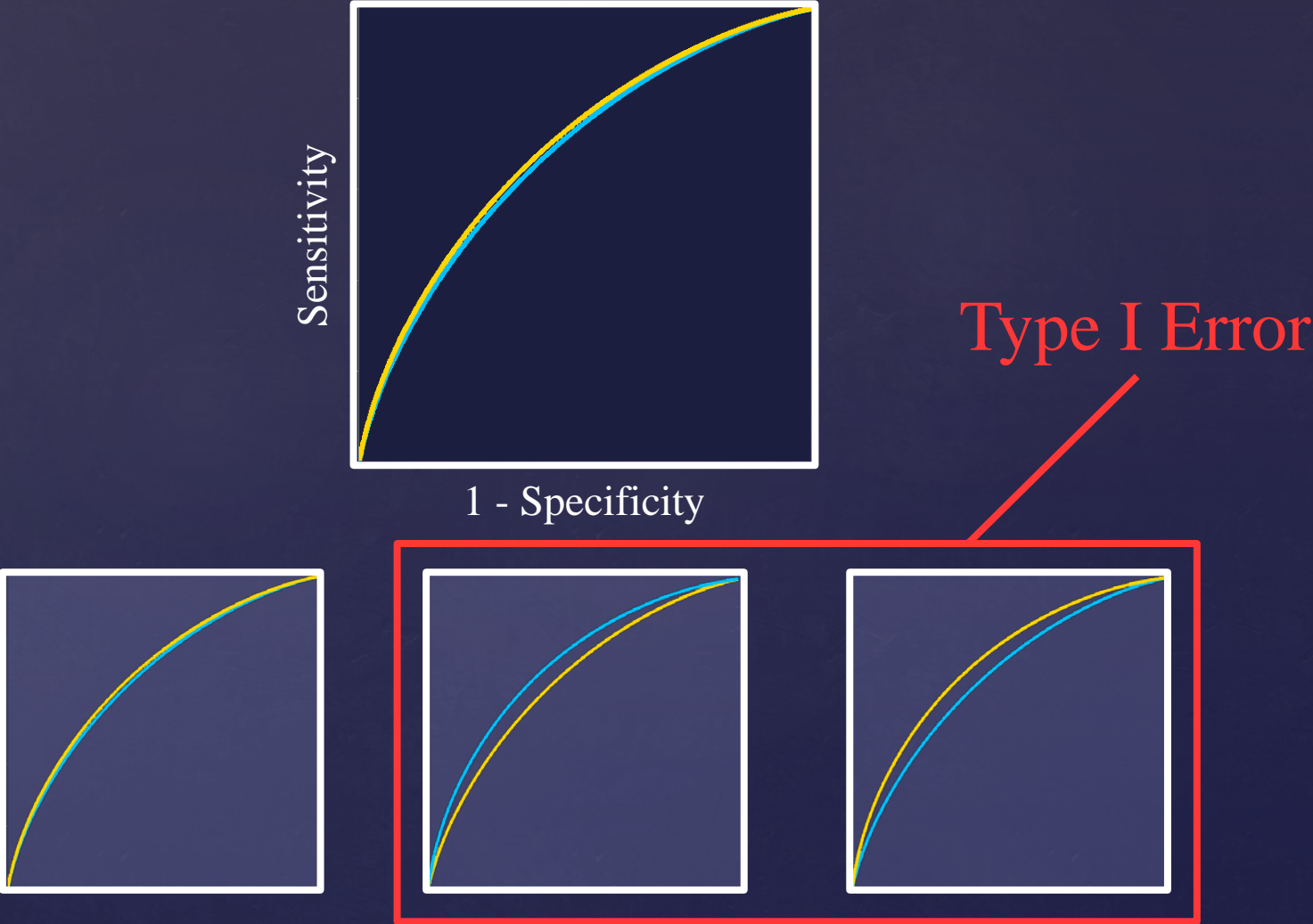
Evaluation of the Method



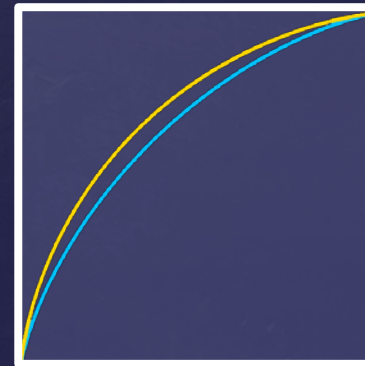
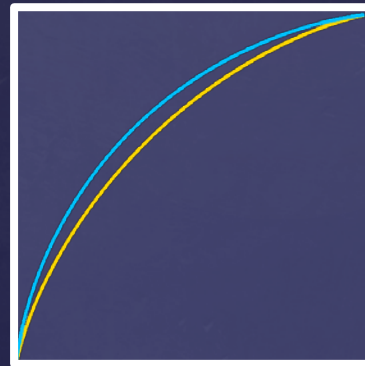
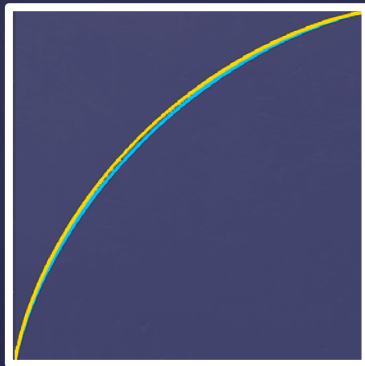
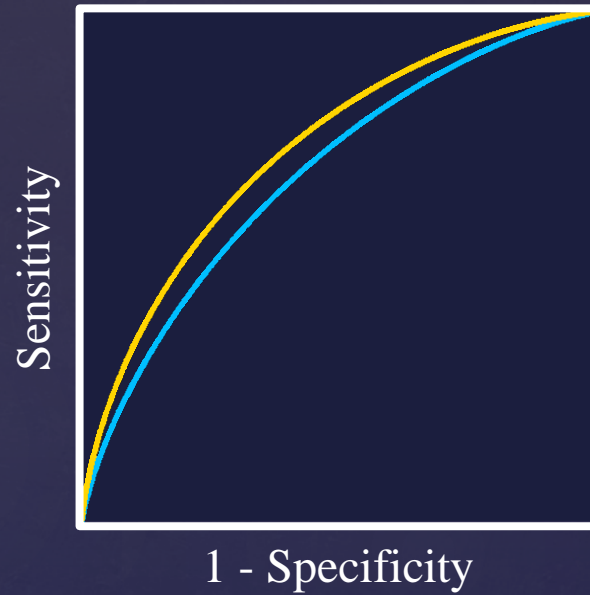
1 - Specificity



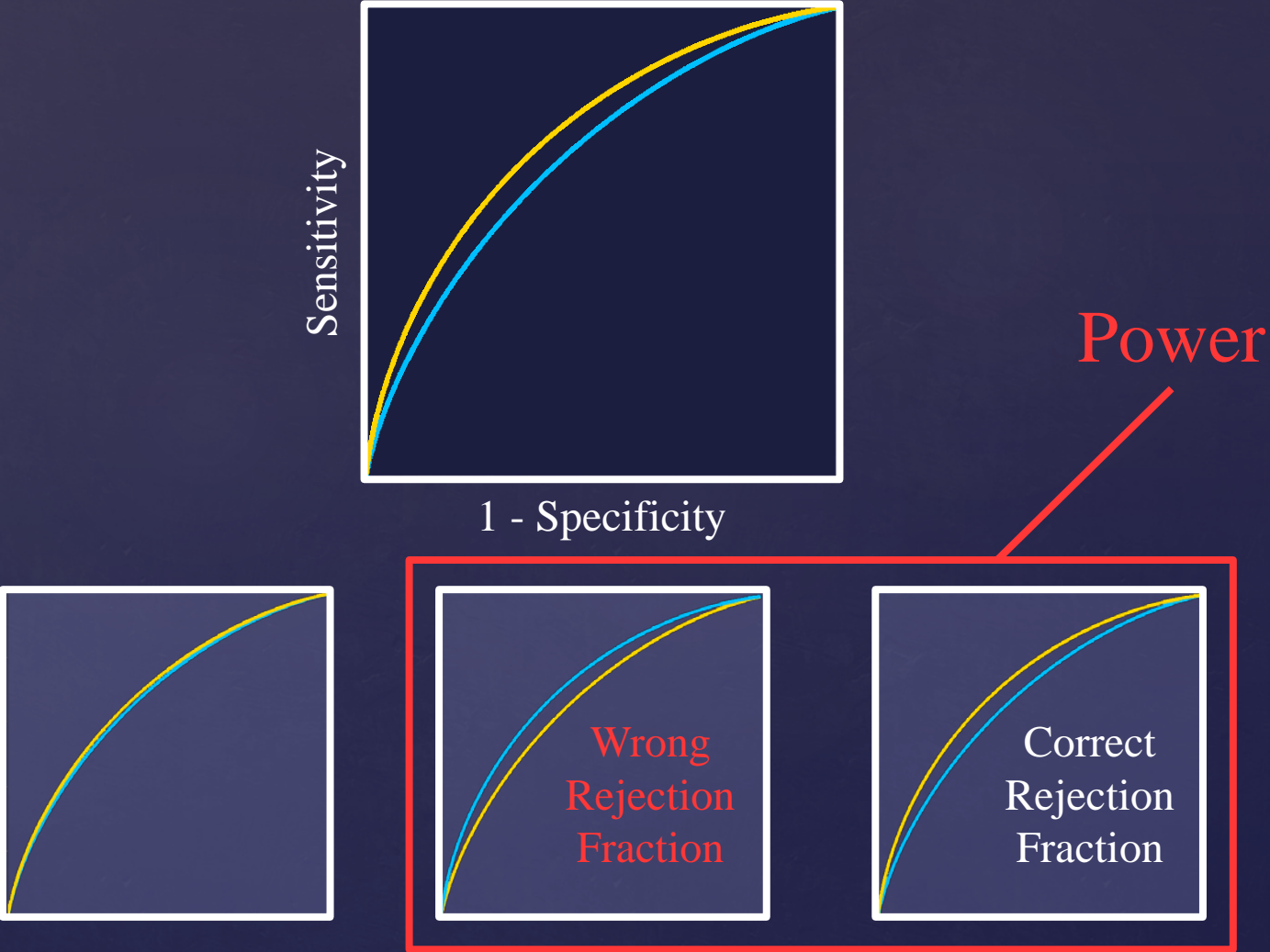
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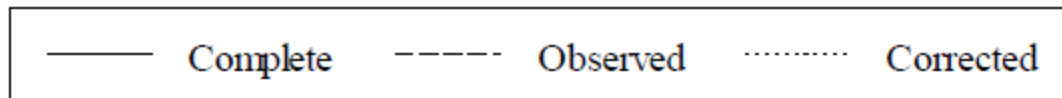
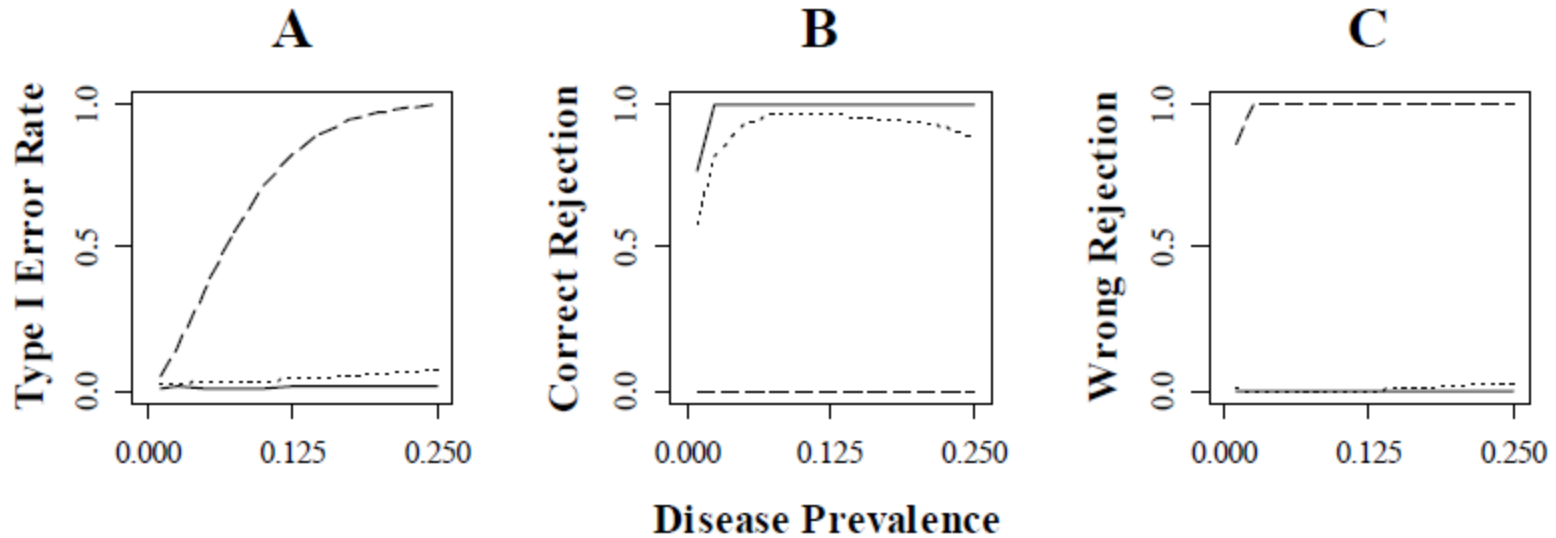
Evaluation of the Method



Evaluation of the Method



Disease Prevalence Simulation Study



Percent Identified Simulation Study

Bias?	Percent Identified (Test 1 / Test 2)	Complete Type I Error	Observed Type I Error	Corrected Type I Error
Yes	15/50	0.01	0.89	0.36
	15/80	0.02	0.95	0.25
	50/80	0.01	0.23	0.12
No	15/15	0.01	0.02	0.23
	50/50	0.01	0.02	0.12
	80/80	0.02	0.02	0.18

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No	15/15	0.01	0.02	0.23
	50/50	0.01	0.02	0.12
	80/80	0.02	0.02	0.18

Recommendation

- Study investigators should conduct a **simulation of their study** using both the standard analysis and the bias correction method.
- Study investigators should choose the analysis plan that has a **nominal Type I error rate** and the **highest power** for the correct decision.

Oral Cancer Screening Demonstration

VISIBLE LIGHT



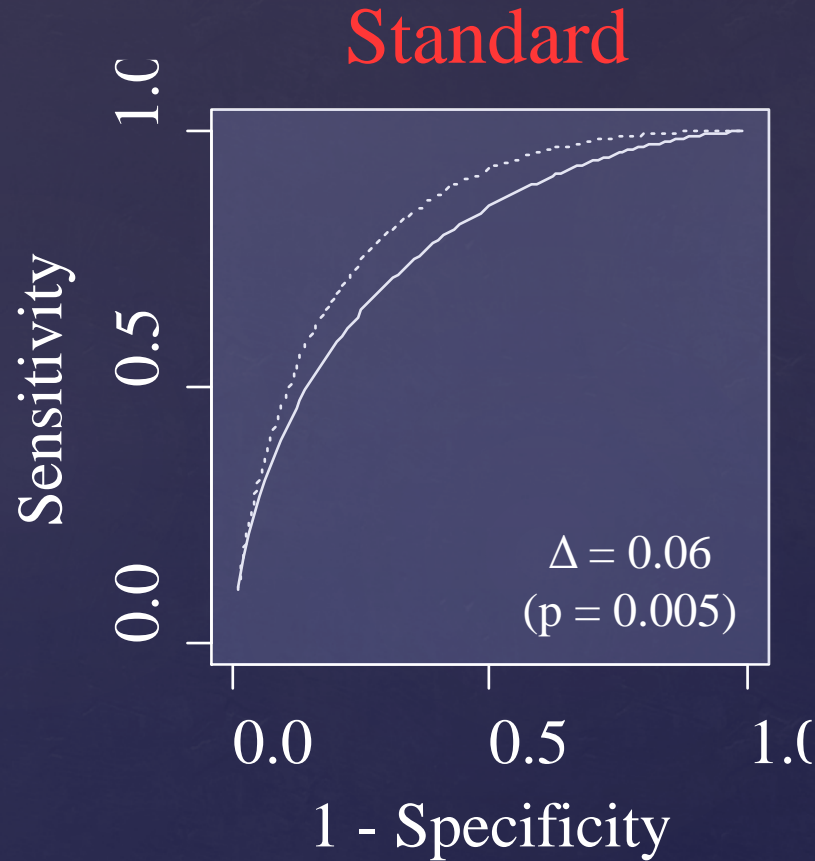
No visible lesion

AUTOFLUORESCENCE



Dark region confirmed to be
carcinoma in situ

Oral Cancer Screening Analysis

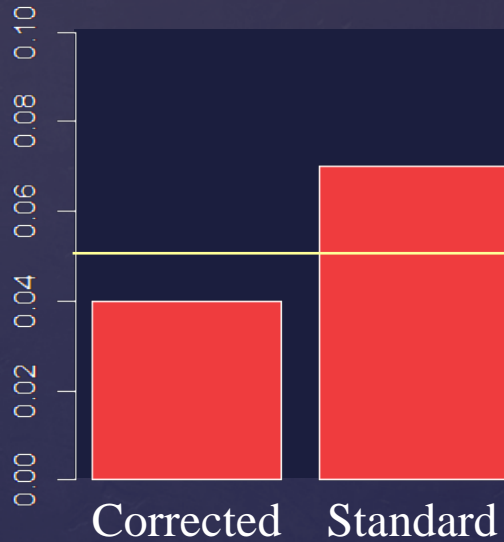


..... Autofluorescence

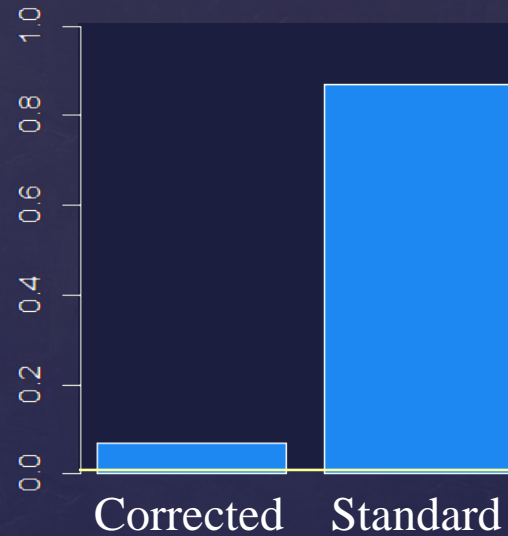
— Visible Light

Decision Errors Simulation

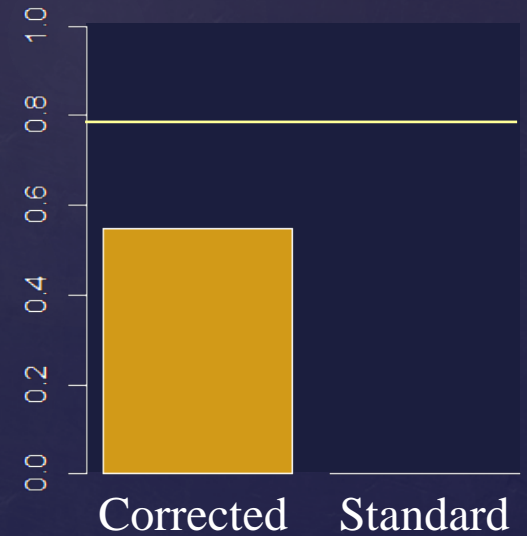
Type I Error



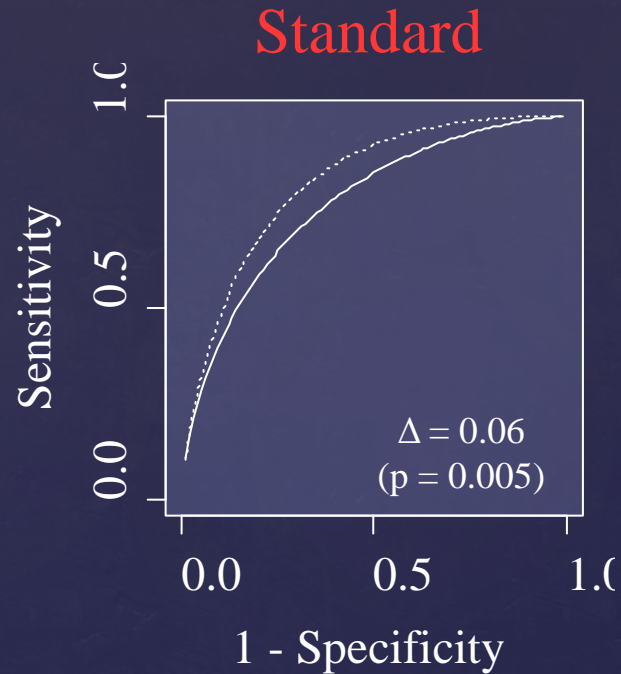
Wrong Rejection



Correct Rejection



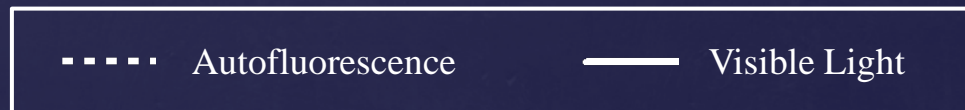
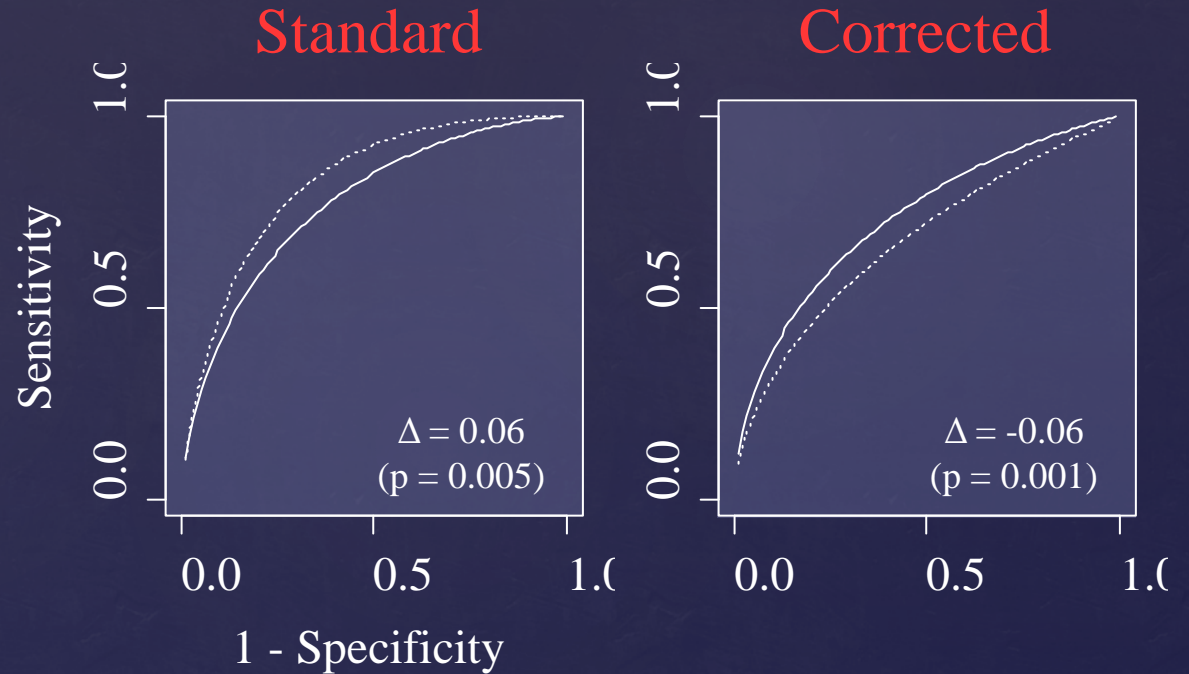
Oral Cancer Screening Analysis



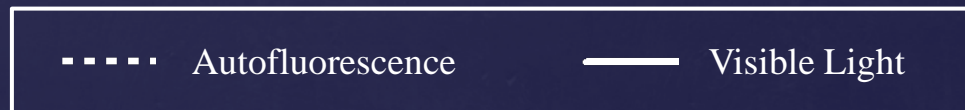
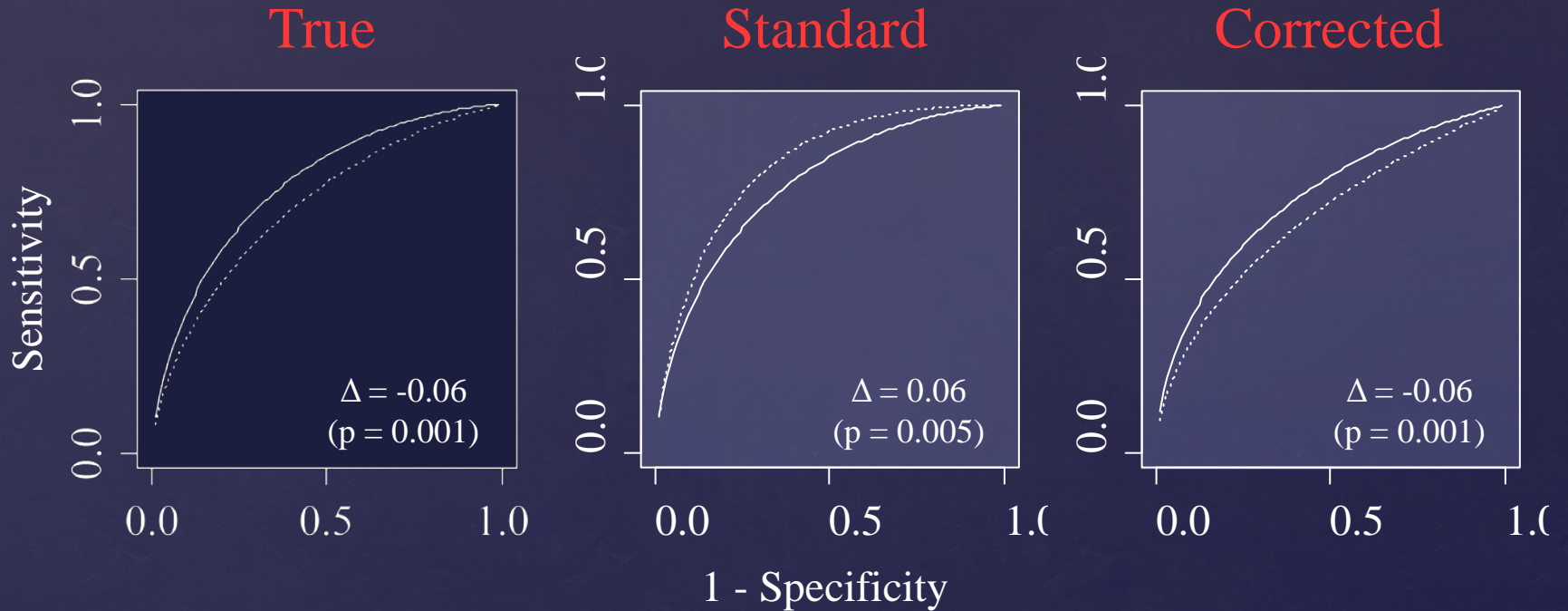
- - - - - Autofluorescence

— Visible Light

Oral Cancer Screening Analysis



Oral Cancer Screening Analysis



References

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Literature Review

- Re-weighting, imputation, and Bayesian approaches have been proposed to reduce the effect of partial verification bias
- Maximum likelihood methods and latent class models have been proposed to estimate diagnostic accuracy in the presence of imperfect reference standard bias
- A method using general estimating equations can correct for missing disease status, but does not account for misclassification of disease status.
- We have not found any methods that reduce the effect of paired screening trial bias.

Nath Algorithm

$$\hat{\mu}_1 = \bar{x} - \left(\hat{P} + \hat{\rho} \hat{Q} \right) \hat{\sigma}_1$$

$$\hat{\mu}_2 = \bar{y} - \left(\hat{Q} + \hat{\rho} \hat{P} \right) \hat{\sigma}_2$$

$$\hat{\sigma}_1 = s_1 \left[\left(1 + \hat{\xi} \hat{P} \right) + \hat{\rho}^2 \left(\hat{A} + \hat{\eta} \hat{Q} \right) - \left(\hat{P} + \hat{\rho} \hat{Q} \right)^2 \right]^{-\frac{1}{2}}$$

$$\hat{\sigma}_2 = s_2 \left[\left(1 + \hat{\eta} \hat{Q} \right) + \hat{\rho}^2 \left(\hat{A} + \hat{\xi} \hat{P} \right) - \left(\hat{Q} + \hat{\rho} \hat{P} \right)^2 \right]^{-\frac{1}{2}}$$

$$\hat{\rho} = r s_1 s_2 (\hat{\sigma}_1 \hat{\sigma}_2)^{-1} \left[\left(1 + \hat{\xi} \hat{P} + \hat{\eta} \hat{Q} + \hat{A} \right) - \left(\hat{P} + \hat{\rho} \hat{Q} \right) \left(\hat{Q} + \hat{\rho} \hat{P} \right) / \hat{\rho} \right]^{-1}$$

Nath Algorithm

$$\hat{\mu}_1 = \bar{x} - \left(\hat{P} + \hat{\rho} \hat{Q} \right) \hat{\sigma}_1$$

$$\hat{\mu}_2 = \bar{y} - \left(\hat{Q} + \hat{\rho} \hat{P} \right) \hat{\sigma}_2$$

$$1 \quad \hat{\sigma}_1 = s_1 \left[\left(1 + \hat{\xi} \hat{P} \right) + \hat{\rho}^2 \left(\hat{A} + \hat{\eta} \hat{Q} \right) - \left(\hat{P} + \hat{\rho} \hat{Q} \right)^2 \right]^{-\frac{1}{2}}$$

$$1 \quad \hat{\sigma}_2 = s_2 \left[\left(1 + \hat{\eta} \hat{Q} \right) + \hat{\rho}^2 \left(\hat{A} + \hat{\xi} \hat{P} \right) - \left(\hat{Q} + \hat{\rho} \hat{P} \right)^2 \right]^{-\frac{1}{2}}$$

$$\hat{\rho} = r s_1 s_2 (\hat{\sigma}_1 \hat{\sigma}_2)^{-1} \left[\left(1 + \hat{\xi} \hat{P} + \hat{\eta} \hat{Q} + \hat{A} \right) - \left(\hat{P} + \hat{\rho} \hat{Q} \right) \left(\hat{Q} + \hat{\rho} \hat{P} \right) / \hat{\rho} \right]^{-1}$$

Nath Algorithm

$$2 \quad \hat{\mu}_1 = \bar{x} - \left(\hat{P} + \hat{\rho} \hat{Q} \right) \hat{\sigma}_1$$

$$2 \quad \hat{\mu}_2 = \bar{y} - \left(\hat{Q} + \hat{\rho} \hat{P} \right) \hat{\sigma}_2$$

$$\hat{\sigma}_1 = s_1 \left[\left(1 + \hat{\xi} \hat{P} \right) + \hat{\rho}^2 \left(\hat{A} + \hat{\eta} \hat{Q} \right) - \left(\hat{P} + \hat{\rho} \hat{Q} \right)^2 \right]^{-\frac{1}{2}}$$

$$\hat{\sigma}_2 = s_2 \left[\left(1 + \hat{\eta} \hat{Q} \right) + \hat{\rho}^2 \left(\hat{A} + \hat{\xi} \hat{P} \right) - \left(\hat{Q} + \hat{\rho} \hat{P} \right)^2 \right]^{-\frac{1}{2}}$$

$$\hat{\rho} = r s_1 s_2 (\hat{\sigma}_1 \hat{\sigma}_2)^{-1} \left[\left(1 + \hat{\xi} \hat{P} + \hat{\eta} \hat{Q} + \hat{A} \right) - \left(\hat{P} + \hat{\rho} \hat{Q} \right) \left(\hat{Q} + \hat{\rho} \hat{P} \right) / \hat{\rho} \right]^{-1}$$

Nath Algorithm

$$\hat{\mu}_1 = \bar{x} - \left(\hat{P} + \hat{\rho} \hat{Q} \right) \hat{\sigma}_1$$

$$\hat{\mu}_2 = \bar{y} - \left(\hat{Q} + \hat{\rho} \hat{P} \right) \hat{\sigma}_2$$

$$\hat{\sigma}_1 = s_1 \left[\left(1 + \hat{\xi} \hat{P} \right) + \hat{\rho}^2 \left(\hat{A} + \hat{\eta} \hat{Q} \right) - \left(\hat{P} + \hat{\rho} \hat{Q} \right)^2 \right]^{-\frac{1}{2}}$$

$$\hat{\sigma}_2 = s_2 \left[\left(1 + \hat{\eta} \hat{Q} \right) + \hat{\rho}^2 \left(\hat{A} + \hat{\xi} \hat{P} \right) - \left(\hat{Q} + \hat{\rho} \hat{P} \right)^2 \right]^{-\frac{1}{2}}$$

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$$\hat{\rho} = r s_1 s_2 (\hat{\sigma}_1 \hat{\sigma}_2)^{-1} \left[\left(1 + \hat{\xi} \hat{P} + \hat{\eta} \hat{Q} + \hat{A} \right) - \left(\hat{P} + \hat{\rho} \hat{Q} \right) \left(\hat{Q} + \hat{\rho} \hat{P} \right) / \hat{\rho} \right]^{-1}$$

We derived expressions for the weighted parameter estimates using the conditional covariance formula (Proposition 5.2, p. 348, Ross, 2009) and the definition of the weighted mean (Equation 3.2.1, p. 77, Kish, 1995). We define the estimates as follows:

$$\hat{\mu}_{11,W} = \hat{\lambda}\bar{X}_{11,A} + (1 - \hat{\lambda})\bar{X}_{11,B}, \quad (3)$$

$$\hat{\mu}_{21,W} = \hat{\lambda}\bar{X}_{21,A} + (1 - \hat{\lambda})\bar{X}_{21,B}, \quad (4)$$

$$\hat{\sigma}_{11,W}^2 = \mathcal{G}_1 + \mathcal{H}_1 - \hat{\mu}_{11,W}^2, \quad (5)$$

$$\hat{\sigma}_{21,W}^2 = \mathcal{G}_2 + \mathcal{H}_2 - \hat{\mu}_{21,W}^2, \quad (6)$$

and

$$\hat{\rho}_{1,W} = \hat{\sigma}_{11,W}^2 \hat{\sigma}_{21,W}^2 (\mathcal{P} + \mathcal{Q} - \hat{\mu}_{11,W} \hat{\mu}_{21,W}) \quad (7)$$

where

$$\mathcal{G}_j = \hat{\lambda}(\bar{X}_{j1,A}^2 + S_{j1,A}^2),$$

$$\mathcal{H}_j = (1 - \hat{\lambda})(\bar{X}_{j1,B}^2 + S_{j1,B}^2),$$

$$\mathcal{P} = \hat{\lambda}\bar{X}_{11,A}\bar{X}_{21,A} + \hat{\lambda}S_{11,A}S_{21,A}r_{1,A},$$

and

$$\mathcal{Q} = (1 - \hat{\lambda})\bar{X}_{11,B}\bar{X}_{21,B} + (1 - \hat{\lambda})S_{11,B}S_{21,B}r_{1,B}.$$