

Monthly Research Presentation

Partial Verification Bias Correction in Diagnostic Accuracy Studies

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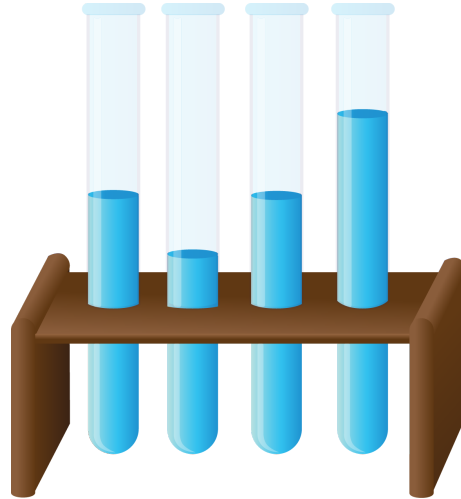
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Outlines

- Partial Verification Bias
- Bias Correction Methods
- PVBcorrect R Package
- Relevance to Medical Researchers
- Conclusion

Partial Verification Bias

Background



Diagnostic Tests

Extremely important role in medical care¹

Objective assessment²

Disease vs No disease³



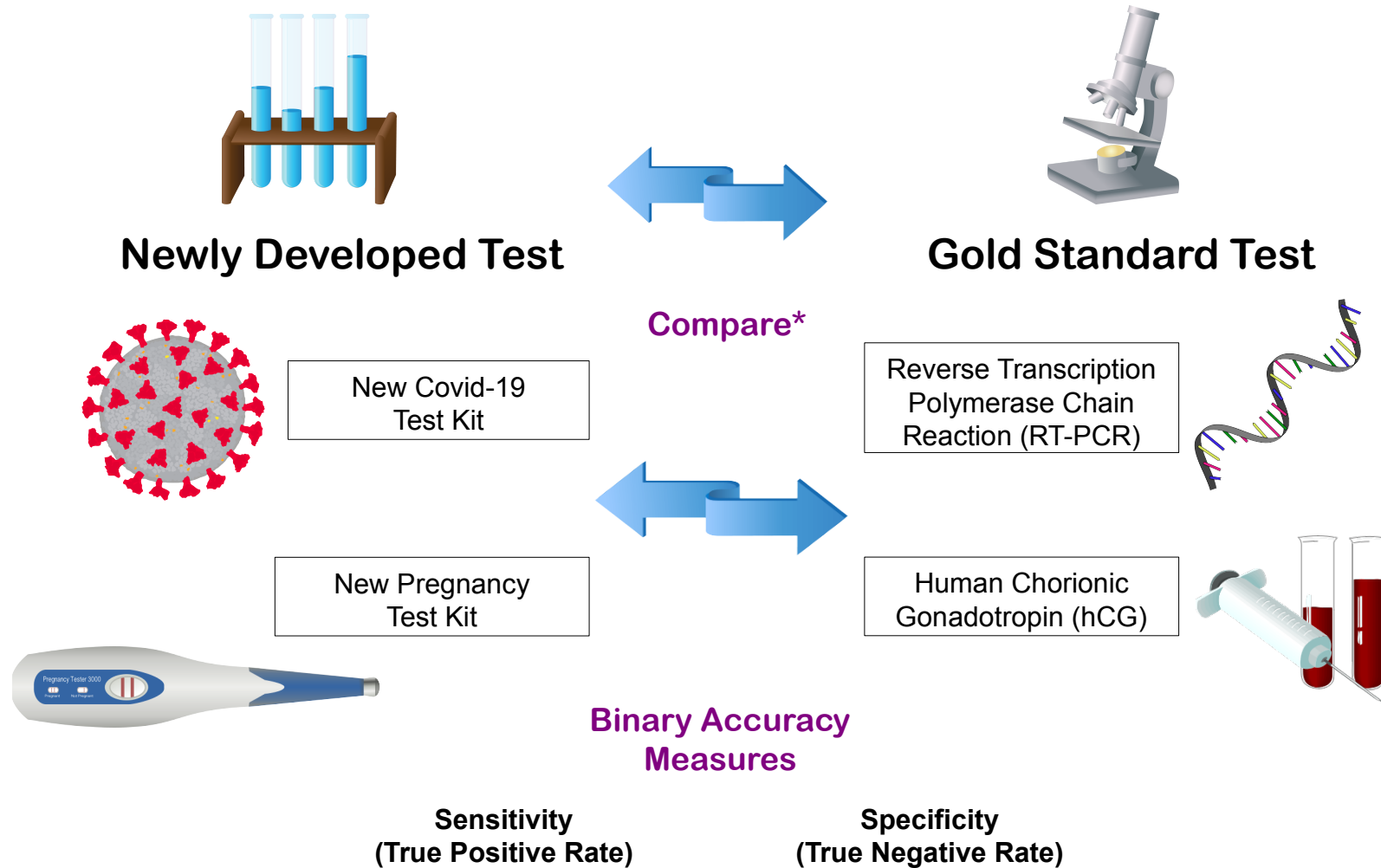
Newly Developed Tests

Evaluation⁴

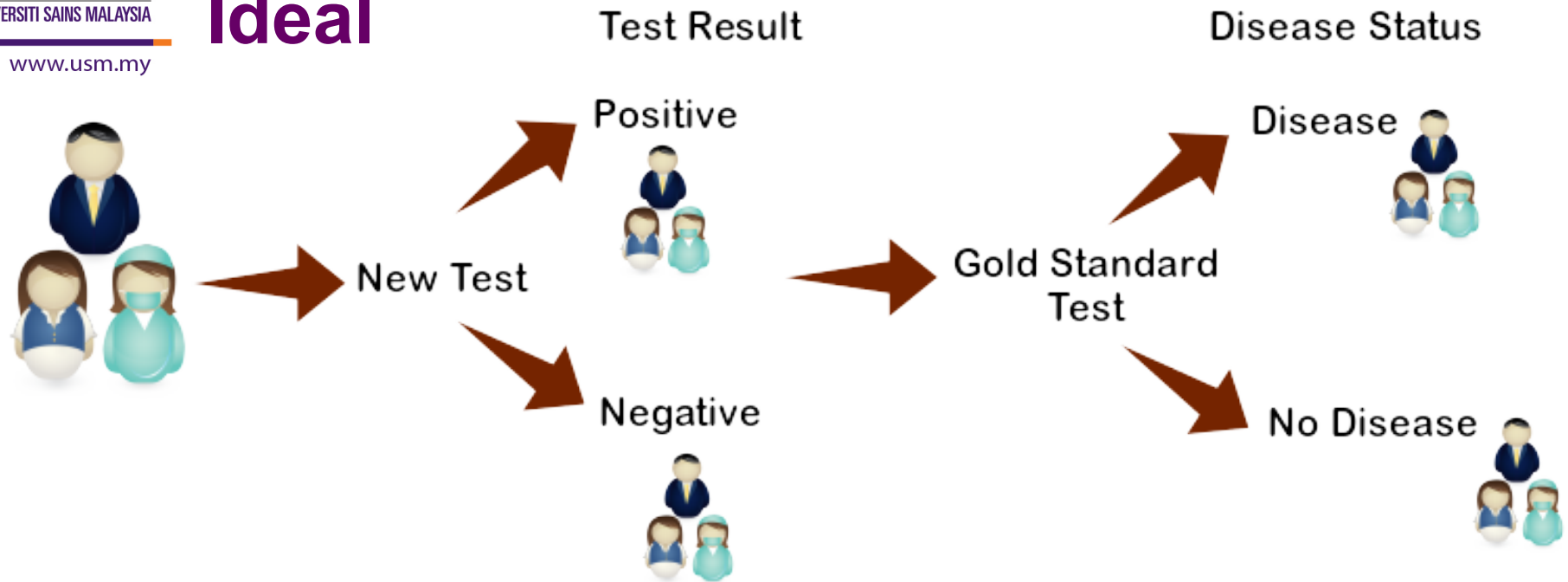


Diagnostic Accuracy Studies

Diagnostic accuracy studies

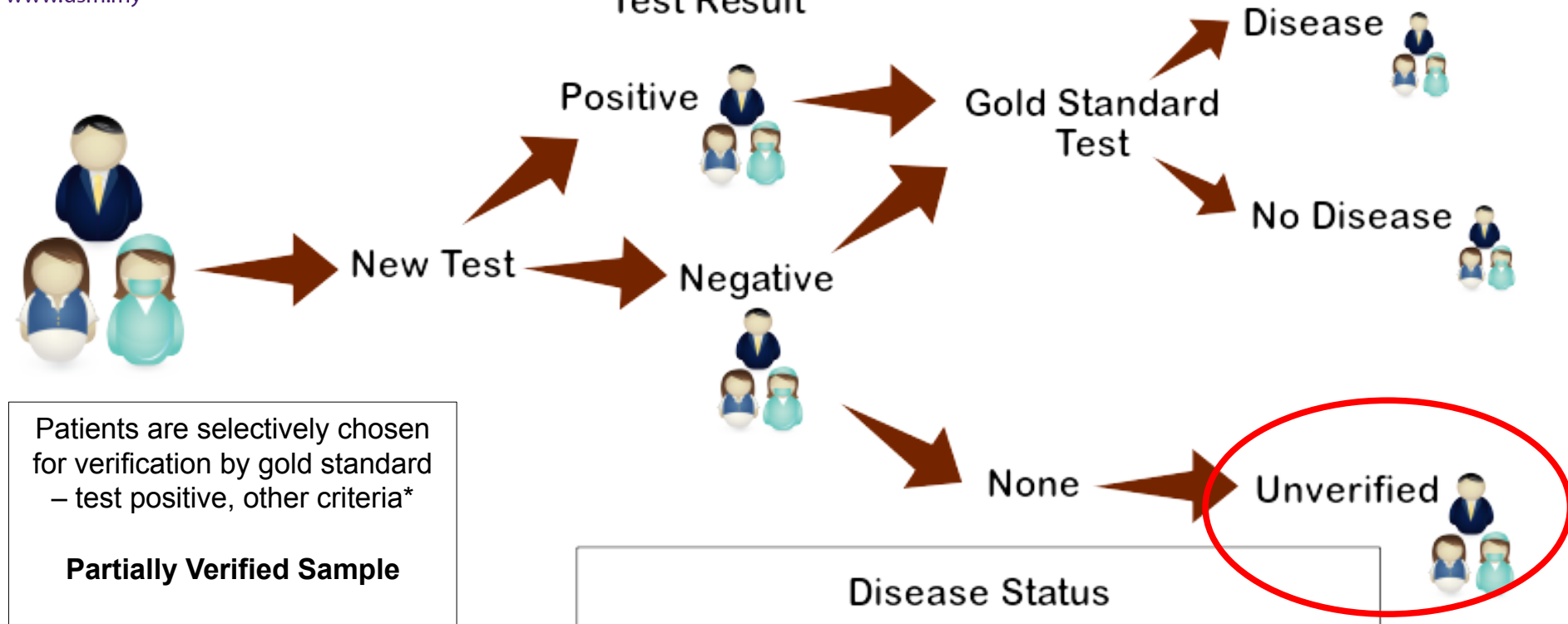


Ideal



Test Result	Disease Status	
	P	N
P	TP	FP
N	FN	TN
Sensitivity = $\frac{TP}{TP+FN}$		Specificity = $\frac{TN}{TN+FP}$

Partial Verification Bias (PVB)

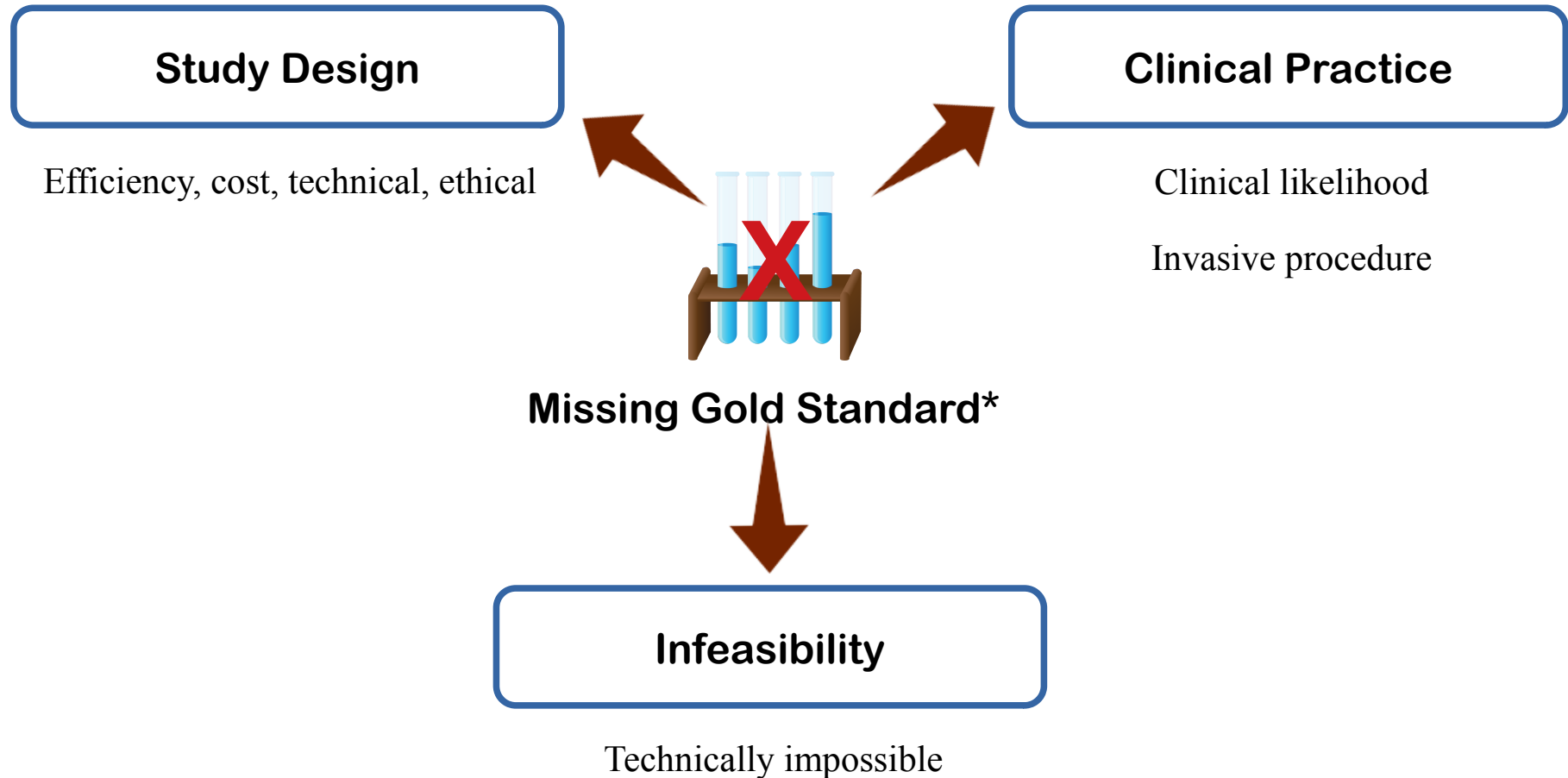


Test Result	Disease Status		
	P	N	Unverified
P	TP	FP	?TP/FP
N	FN	TN	?FN/TN

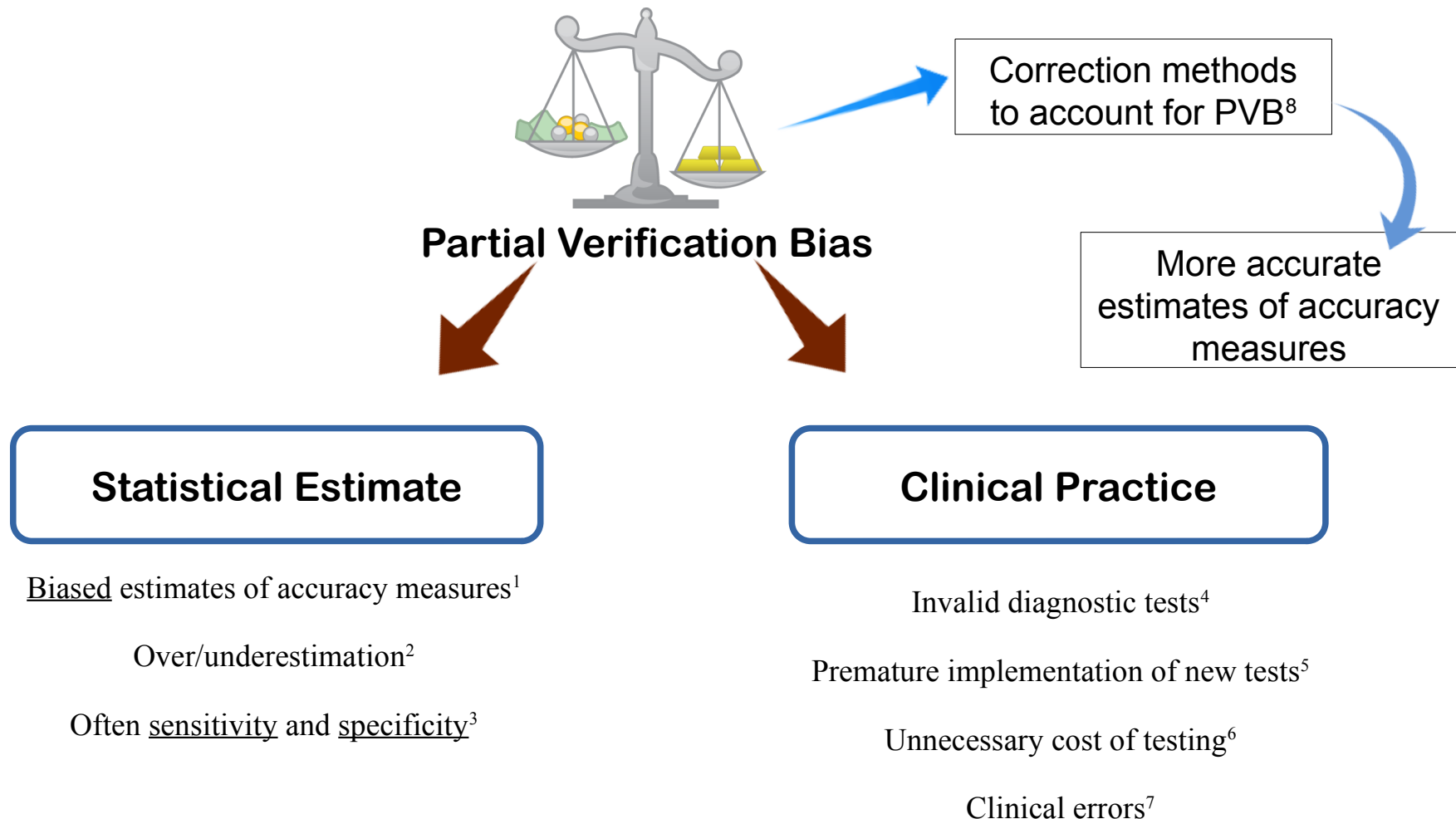
Sensitivity?

Specificity?

Partial Verification Bias (PVB)

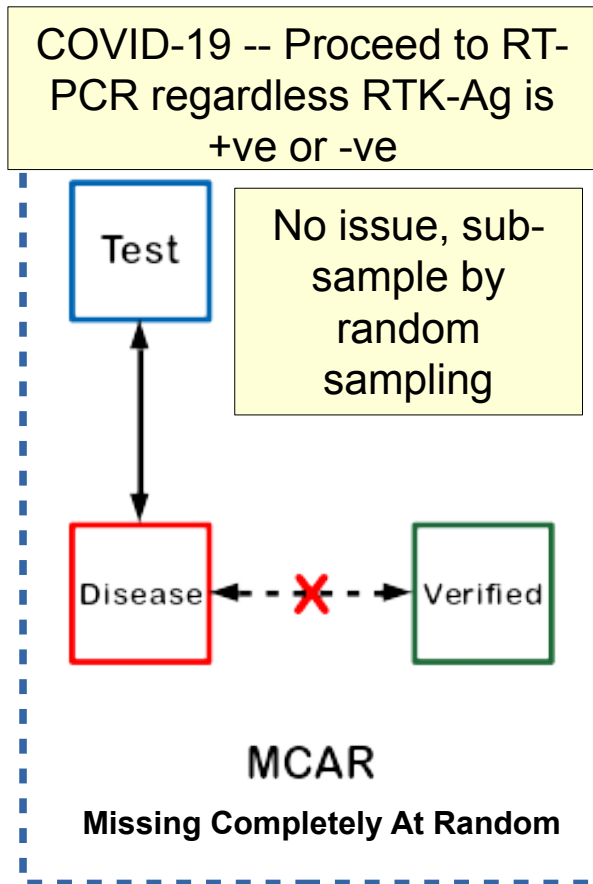


Impacts of Partial Verification Bias



Bias Correction Methods

Missing Data Mechanisms in PVB



No selection bias

*Adapted from Figure 2 in Schafer & Graham (2002)

COVID-19 -- Proceed to RT-PCR when RTK-Ag is +ve

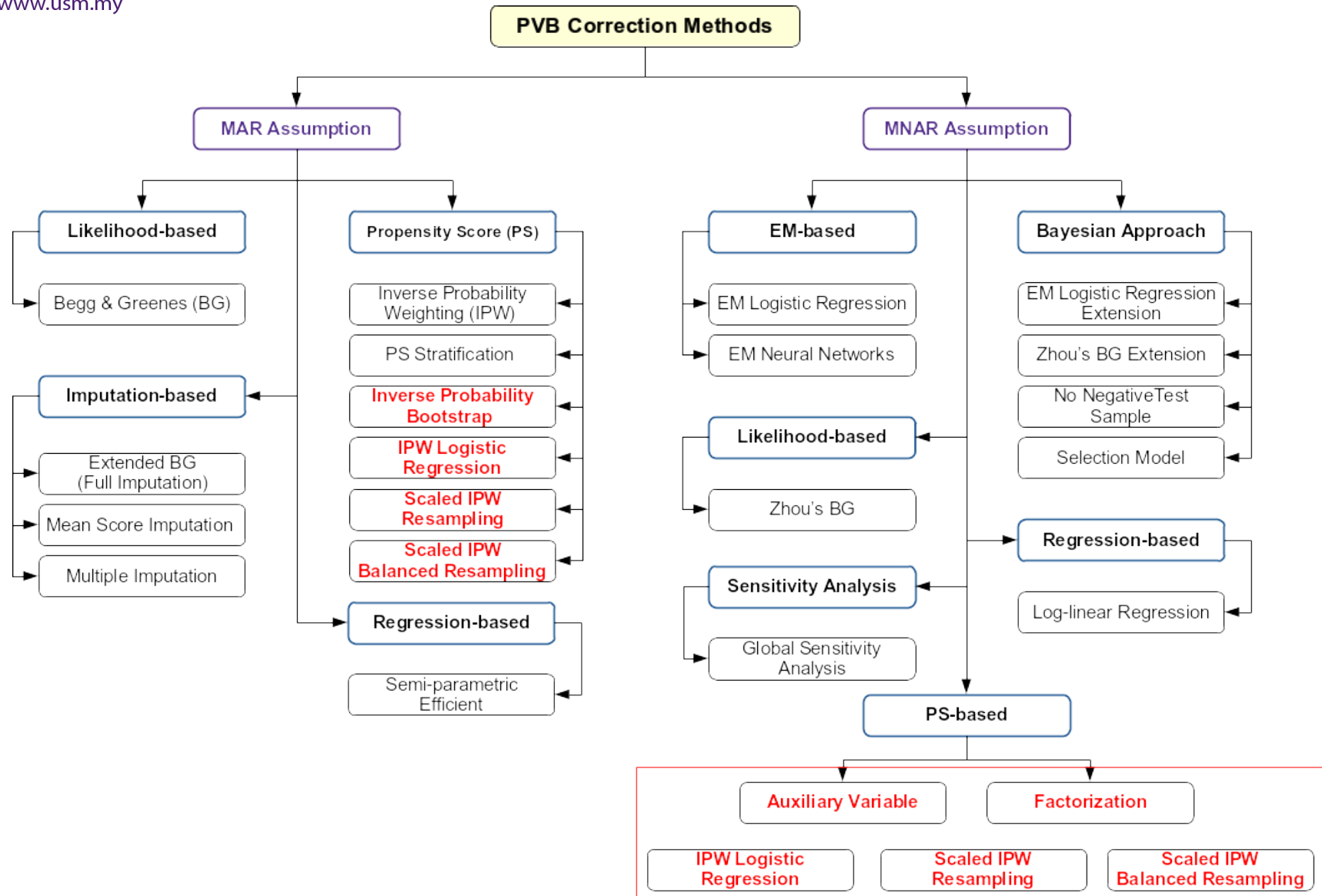
COVID-19 -- Proceed to RT-PCR when RTK-Ag is +ve AND patient looks sick

Cause of selection bias known & recorded

Cause of selection bias not known

With Partial Verification Bias

Existing PVB Correction Methods



PVBcorrect R Package

About

- Available in GitHub



github.com/wnarifin/PVBcorrect

- Tutorial published in



doi.org/10.1002/sim.9311

Features

- Version 0.1.1
- Available methods:
 - Begg and Greenes' methods
 - Multiple imputation method by logistic regression
 - EM-based logistic regression method
- Point estimates & CIs
- Other methods still in coding stage

Installation

In R command:

```
install.packages("devtools")  
devtools::install_github("wnarifin/PVBcorrect")
```


Some example outputs

No correction

```
cca_out = acc_cca(data = cad_pvb, test = "T", disease = "D", ci = TRUE)
cca_out$acc_results
```

```
## Estimates of accuracy measures
## Uncorrected for PVB: Complete Case Analysis
##
##           Est           SE      LowCI      UppCI
## Sn  0.9750000  0.01103970  0.9533626  0.9966374
## Sp  0.1439114  0.02132173  0.1021216  0.1857013
## PPV  0.4566745  0.02410569  0.4094282  0.5039207
## NPV  0.8863636  0.04784519  0.7925888  0.9801385
```

Larger Sn
Lower Sp

Corrected by Extended B&G

```
ebg_out = acc_ebg(data = cad_pvb, test = "T", disease = "D", ci = TRUE,
ebg_out$acc_results
```

```
## Estimates of accuracy measures
## Corrected for PVB: Extended Begg and Greenes' Method
##
##           Est           SE      LowCI      UppCI
## Sn  0.8188629  0.06438696  0.6860317  0.9360441
## Sp  0.5918754  0.01759285  0.5587741  0.6285913
## PPV  0.4566745  0.02421467  0.4063396  0.5059415
## NPV  0.8863636  0.04925696  0.8005051  0.9949495
```

Corrected Sn &
Sp

Corrected by Multiple Imputation

```
mi_out = acc_mi(data = cad_pvb, test = "T", disease = "D", ci = TRUE, seednum = 12345)
mi_out$acc_results
```

```
## Estimates of accuracy measures
## Corrected for PVB: Multiple Imputation Method
##
##           Est           SE      LowCI      UppCI
## Sn  0.8031807  0.06451495  0.6753082  0.9310532
## Sp  0.5874592  0.02199350  0.5440683  0.6308500
## PPV  0.4578988  0.02400632  0.4105655  0.5052321
## NPV  0.8699605  0.05498024  0.7609444  0.9789765
```

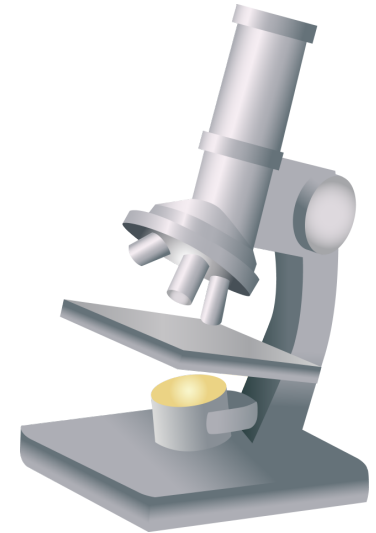
Corrected Sn &
Sp

https://wnarifin.github.io/workshop/conf_r2022/PVBinR_Demo.html

Relevance to Medical Researchers

Scenario 1

- Already-conducted diagnostic accuracy study – might be biased – PVB
- Gold standard test – only done to selected patients
- Identify reasons (variables) why patients are not selected
- Reanalyze / Reevaluate – Perform PVB correction for the diagnostic test performance results and compare w/out correction

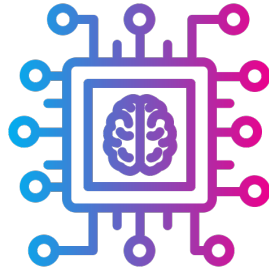


Scenario 2



- Limitation with gold standard test – invasive, costly, infeasible
- With planning – in view of selection bias – document reasons why patients are not selected
- Prioritize patients with higher clinical possibility of having disease (i.e. positive gold standard test)
- Ethical – avoid unnecessary procedure on healthy patients
- Save cost

Scenario 3



- Assume Machine Learning = Diagnostic Test
- Machine learning – the issue with disease labeling / confirmation
- Selective labeling – save clinicians' time
- May consider PVB correction as well – select only cases with high probability of disease for labeling

Conclusion

Take-home messages

- Although gold standard test is invasive/expensive/not feasible, with proper research planning in view of selection bias, diagnostic test performance can be accurately evaluated
- Bias correction is easy when potential sources of selection bias are identified and documented, even if we cannot avoid the bias
- PVBcorrect R Package helps researchers to perform the correction – GUI is planned

Publications

Arifin, W. N., & Yusof, U. K. (2022). Correcting for partial verification bias in diagnostic accuracy studies: A tutorial using R. *Statistics in Medicine*, 41(9), 1709–1727. (Published. Indexed by WoS, Impact Factor: 2.497, Q1 in Statistics & Probability)

Arifin, W. N., & Yusof, U. K. (2022). Correcting for partial verification bias in diagnostic accuracy measures using inverse probability bootstrap sampling. *Diagnostics*, 12(11), 2839. (Published. Indexed by WoS, Impact Factor: 3.992, Q2 Medicine, General & Internal)

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Thank You