



confeRence 2022

Introduction to PVBcorrect: An R Package for Partial Verification Bias Correction

27 November 2022

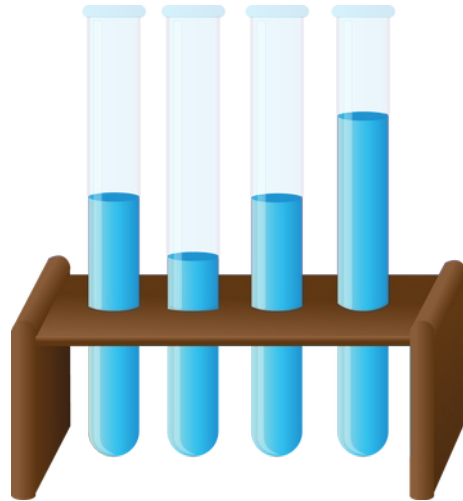
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- Background
- About PVBcorrect package
- Using PVBcorrect package
- Conclusion

Background

Background



Diagnostic Tests

Very important role in medical care¹

Objective assessment²

Disease vs No disease³



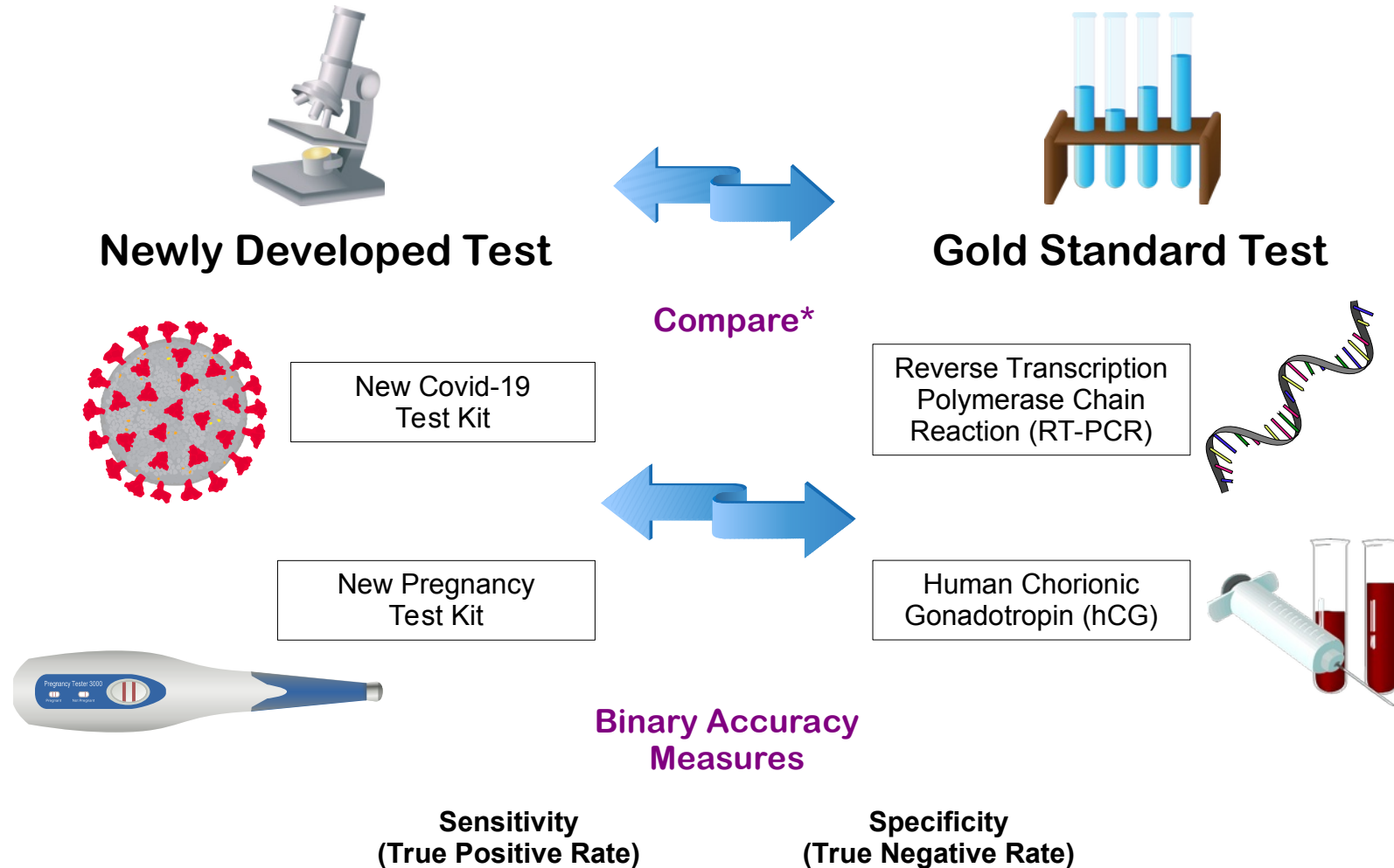
Newly Developed Tests

Evaluation⁴



Diagnostic Accuracy Studies

Diagnostic accuracy studies



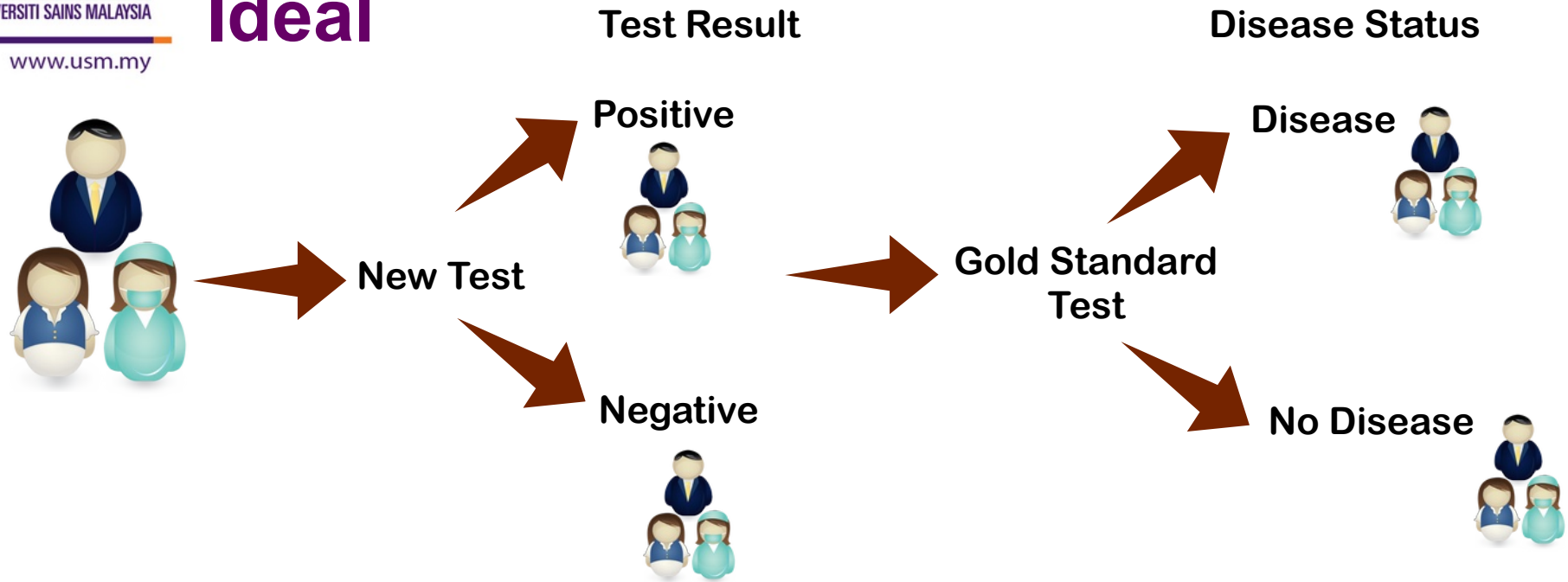
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*Hall et al. (2019), O’Sullivan et al. (2018).

Attributions: ¹https://commons.wikimedia.org/wiki/File:Coronavirus_SVG_Vector_Image.svg ²<https://commons.wikimedia.org/wiki/File:Pregnancy-test.svg>

³https://commons.wikimedia.org/wiki/File:201904_RNA.svg

Ideal



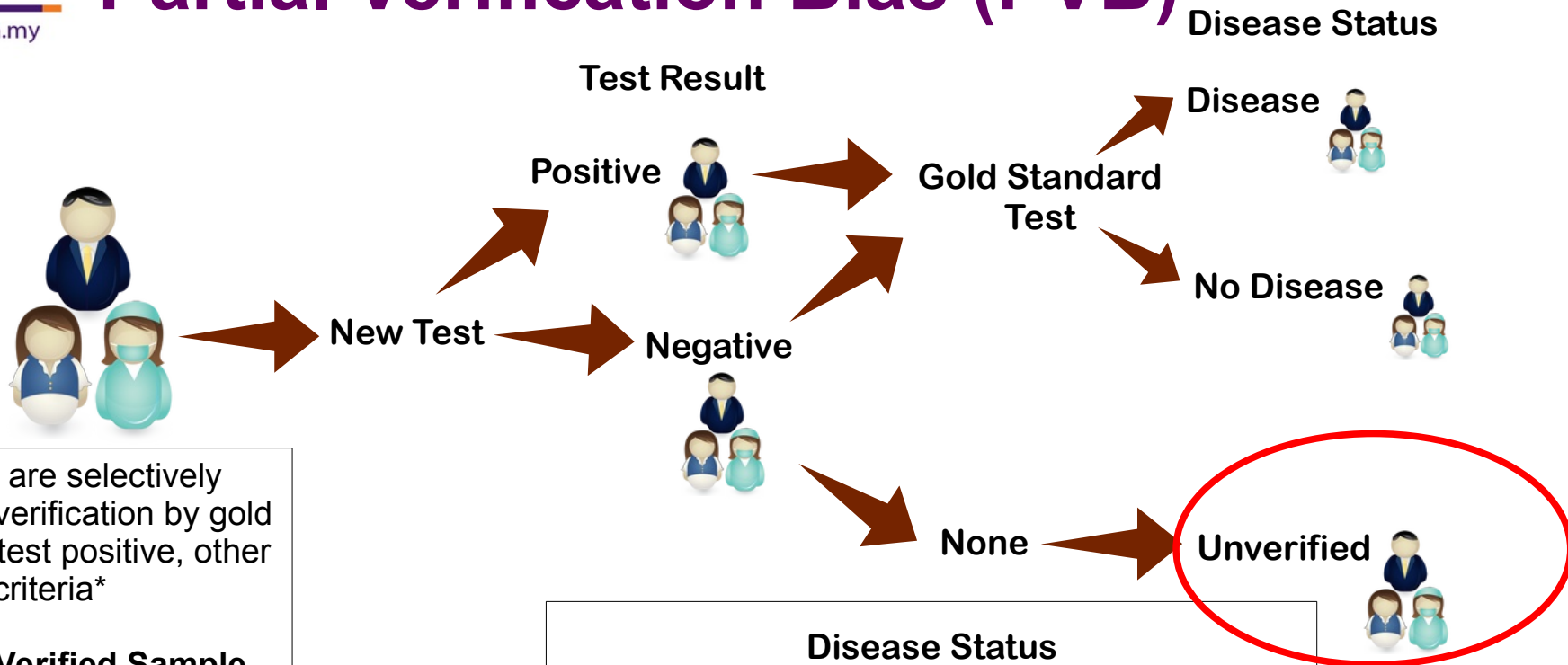
		Disease Status	
		P	N
Test Result	P	TP	FP
	N	FN	TN

Sensitivity=
 $TP / (TP + FN)$

Specificity=
 $TN / (TN + FP)$

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Partial Verification Bias (PVB)



Patients are selectively chosen for verification by gold standard – test positive, other criteria*

Partially Verified Sample

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

Sensitivity?

Specificity?

*O'Sullivan et al. (2018). Abbreviations: TP, true positive; TN, true negative; FP, false positive; FN, false negative.

Background

- Reasons:
 - Study design: Efficiency, technical, ethical.
 - Clinical practice: Clinical likelihood.
 - Infeasibility: Invasive procedures, postmortem diagnosis.

Background

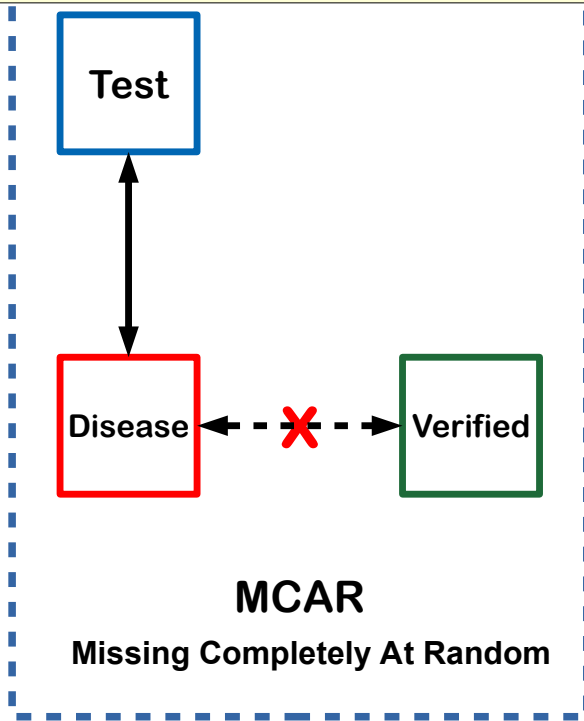
- PVB → Inaccurate estimates of accuracy measures^{1,2}
- Impact on the clinical practice
 - Invalid diagnostic tests³
 - Clinical errors²
- Cannot eliminate verification bias in medical data
→ Need methods to correct PVB⁴

Missing Data Mechanisms in PVB

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

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



MCAR

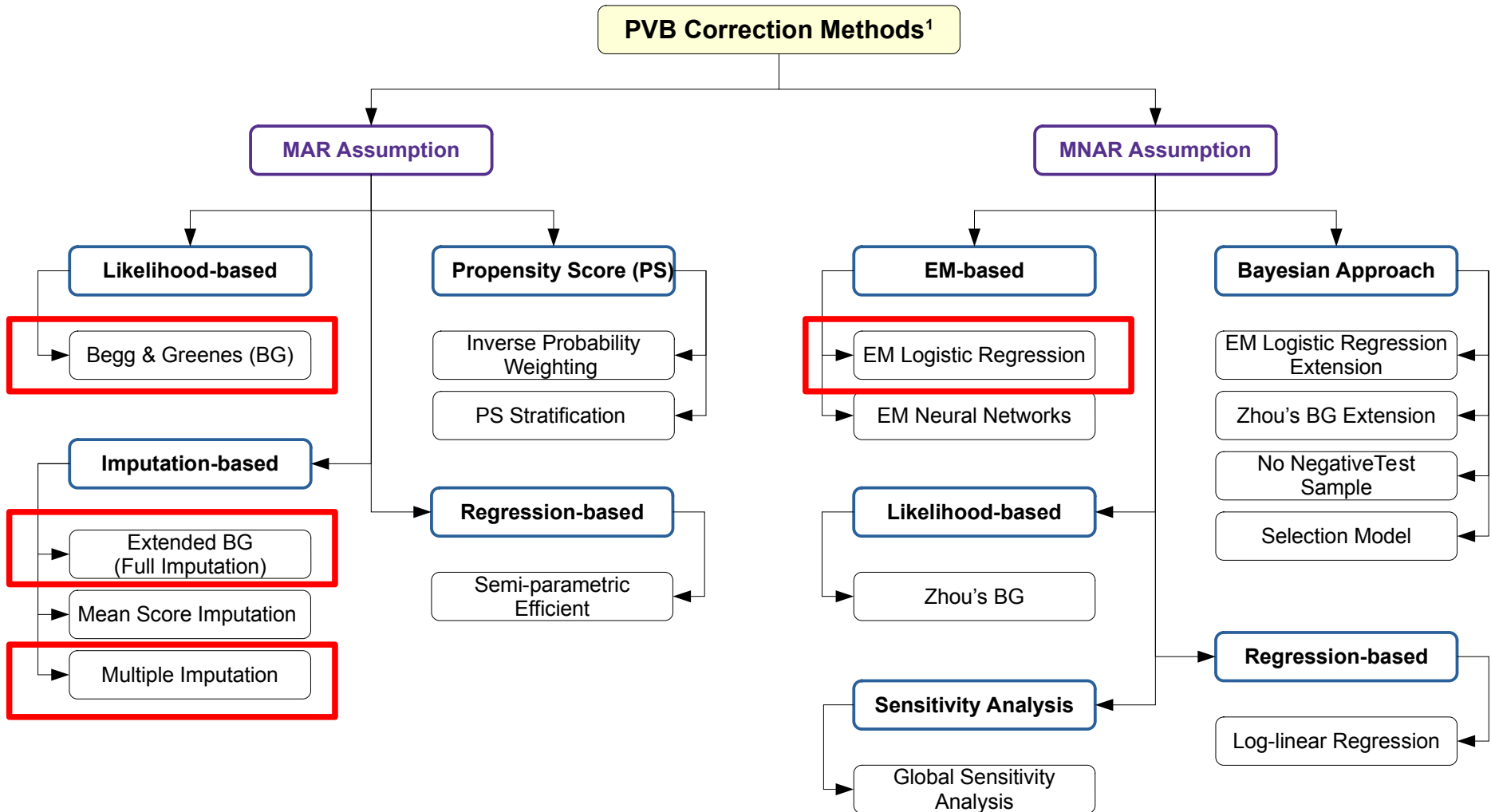
Missing Completely At Random

No selection bias

With Partial Verification Bias

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

Existing PVB Correction Methods



¹Arifin & Yusof, 2022. EM, expectation maximization; MAR, missing at random; MNAR, missing not at random; PVB, partial verification bias.

About PVBcorrect

About



- Available in GitHub

github.com/wnarifin/PVBcorrect

- Tutorial published in

Statistics
in Medicine



doi.org/10.1002/sim.9311

Features

- Version 0.1.1
- Available methods:
 - Begg and Greenes' method
 - Begg and Greenes' methods 1 and 2
 - Multiple imputation method by logistic regression
 - EM-based logistic regression method
- Point estimates & CIs

Installation

- Pre-requisites

```
install.packages("boot", "mice")
```

- Installation

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

Using PVBcorrect

Data Preparation

- Categorical variables, code as 1/0:
 - Outcome, Test variables: Binary (Yes = 1, No = 0)
 - Categorical variables: Binary/Dummy (Yes = 1, No = 0)
- Numerical variables: No issue

Load Data Set

- Load PVBcorrect

```
library(PVBcorrect)
```

- Built-in data set

```
?cad_pvb
```

SPECT Thallium test data set

- Single-photon-emission computed-tomography (SPECT) thallium is a non-invasive diagnostic test used to diagnose coronary artery disease (CAD).
- $n = 2688$ patients, only 471 verified, 2217 unverified (82.5%).
- Data set, five variables:
 1. SPECT thallium test, T: Binary, 1 = Positive, 0 = Negative
 2. CAD, D: Binary, 1 = Yes, 0 = No
 3. Gender, X1: Binary, 1 = Male, 0 = Female
 4. Stress mode, X2: Binary, 1 = Dipyridamole (Medication for stress test when the patient is unable to exercise), 0 = Exercise
 5. Age, X3: Binary, 1 = 60 years and above, 0 = Below 60 years

PVBcorrect Demo

wnarifin.github.io/workshop.html

Conclusion

- To-do list for future updates:
 - Existing methods listed in LR
 - New methods developed by the presenter
 - Methods for numerical diagnostic test
 - GUI interface via R shiny

References

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Thank You