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RetroClac: A Program for Harmonizing Reporting Paradigm in Diagnostic Accuracy Studies

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External link: <https://github.com/mdnx/RetroCalc>

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Protocol status: Working

We use this protocol and it's working

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Abstract

Accurate diagnostic test performance assessment is critical in clinical and research settings, where a wide range of metrics, including accuracy, sensitivity (recall), specificity, f1 score, positive predictive value (PPV), and negative predictive value (NPV), are used to evaluate test validity. However, heterogeneous reporting of these metrics across studies poses challenges for direct comparison and meta-analysis. RetroCalc, a novel program, addresses this gap by reconstructing essential confusion matrix values—true positives (TP), true negatives (TN), false positives (FP), and false negatives (FN)—from minimal input pairs. RetroCalc enables accurate estimation of all relevant diagnostic metrics, even when only limited data is provided. This feature supports meta-analyses by harmonizing data inputs, facilitating more consistent and comprehensive reviews of diagnostic test accuracy studies. RetroCalc provides a valuable tool for enhancing the interpretability and comparability of diagnostic test outcomes, supporting a more unified and accessible approach to diagnostic accuracy research.

Attachments



[RetroCalc.pdf](#)

258KB

Guidelines

Cochrane Handbook for Systematic Reviews of Diagnostic Test Accuracy.

Troubleshooting



Input Preparation

- 1 Prepare the total cohort sizes:
 - Positive cohort ($P = TP + FN$)
 - Negative cohort ($N = TN + FP$)
- 2 Identify one of the following pairs based on your data:
 1. Sensitivity (Sen) and Specificity (Spe)
 2. Sen and Positive Predictive Value (PPV)
 3. Sen and Negative Predictive Value (NPV)
 4. Sen and F1 Score (F1)
 5. Specificity (Spe) and Positive Predictive Value (PPV)
 6. Accuracy (Acc) and Sen
 7. Acc and Spe
 8. Spe and NPV
 9. Likelihood Ratios (LR+) and Spe

Install and Run RetroCalc

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```
# clone the retrocalc into your own enviroment
!git clone https://github.com/mdnx/RetroCalc.git

%cd RetroCalc
!pip install .

# input the required metrics, remember not to use percentage (e.g. 45% incorrect, .45 correct)

from retrocalc.retrocalc import get_user_inputs

get_user_inputs()
```

RetroCalc Mechanism

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 1. Launch the program from the directory:
 2. Provide the input metrics and cohort sizes when prompted.



4. Metric Validation & Reconstruction

1. RetroCalc verifies the input metrics, avoiding division by zero or invalid values.
2. It computes TP, TN, FP, and FN using predefined pairwise formulas.
3. If the input pair is insufficient, the program suggests additional metrics for input.

5. Output and Visualization

1. RetroCalc displays the reconstructed confusion matrix and recalculated diagnostic metrics.
2. Visualize the matrix using the built-in plotting function for easier interpretation.

References

5 References

- RetroCalc GitHub Repository: <https://github.com/mdnx/RetroCalc>
- Cochrane Handbook for Systematic Reviews of Diagnostic Test Accuracy.

Protocol references

Deeks JJ, Bossuyt PM. Chapter 3: Methods and presentation of systematic reviews of test accuracy. In: Deeks JJ, Bossuyt PM, Leeflang MM, Takwoingi Y editor(s). Cochrane Handbook for Systematic Reviews of Diagnostic Test Accuracy Version 2.0 (updated July 2023).