Statistical methods for maximizing the utility of electronic health records data for the development, evaluation, and implementation of real-time prognostic models

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Overview

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- 2. Model Development (Topic 1)
- 3. Model Implementation (Topic 2)
- 4. Model Evaluation (Topic 3)
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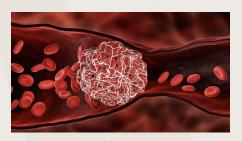


1. Motivation

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Clinical motivation: What is Venous thromboembolism?

- Venous thromboembolism (VTE) occurs when a blood clot forms in a vein.
- VTE is a common complication among hospital inpatients, contributing to longer hospital stays, increased morbidity, and higher mortality rates.
- Prompt medical attention is crucial, making real-time prediction of VTE imperative.



Statistical motivation: Challenge in electronic health records

- An Electronic Health Record (EHR) is a digital version of a patient's medical history, maintained by the healthcare provider over time. It may include all key administrative and clinical data relevant to the patient's care.
- EHR systems are valuable for developing and implementing prognostic models.
- A major challenge is synthesizing multiple predictors to develop accurate prediction models.
- Missing data, especially informative absences reflecting a patient's clinical status, potentially impact prognostic accuracy.
- Most imputation methods focus on accurate value estimation, leaving their effectiveness in real-time prediction modeling uncertain.



2. Model Development (Topic 1)



Data overview

- Data from VUMC's Research Derivative (RD) database.
- Derivation cohort: Inpatients from January 1, 2018, to December 31, 2020 (n=132,330).
- Validation cohort: Inpatients from January 1, 2021, to June 30, 2022 (n=62,546).
- Used ICD-10 codes to identify HA-VTE cases; manual review for accuracy.
- Screened 82 potential risk factors: demographic/clinical characteristics, diagnostic procedures, vital signs, and lab measurements.



Primary objectives

- Train and validate statistical and machine learning models for real-time clinical prediction of VTE.
 - Logistic regression
 - Random forest
 - XG boost
 - Neural network
 - Ensemble method
- Compare the performance between models.
 - C statistic, Integrated calibration index (ICI), Calibration slope, Brier score

Model description

- **Logistic regression:** A statistical model that captures the relationship between predictors and binary outcome using a logistic function.
- Random forest: This machine learning algorithm operates by taking the majority vote of multiple decision trees, improving prediction accuracy and robustness.
- XG boost: A machine learning algorithm that enhances prediction by sequentially updating decision trees to minimize the loss function and optimize the model.
- Neural network: This machine learning model identifies complex relationships in a dataset by mimicking the functions of the human brain through interconnected nodes.
- Ensemble method: An advanced algorithm that combines the predictions
 of the four models above by averaging their outputs to enhance overall
 prediction accuracy.



Model characteristics

Classifier	R package	Command	Requires dummy coding	Hyperparameters (argument)	Selected hyperparameter values	
Logistic regression	caret	glm	No	• None	None	
Random forest	randomforestSRC	rfsrc	No	 Number of tree (ntree) Number of split (mtry) Node size (nodesize) 	 ntree = 3000 mtry = 82 nodesize = 30 	
XG boost	caret	xgbTree	No	Learning rate (eta) Max depth of tree (max_depth) Min split loss (gamma) Percentage of features for tree (colsample_bytree) Min sum of instance weight (min child weight)	eta = 0.3 max_depth = 2 gamma = 0 colsample_bytree = 0.8 min_child_weight = 1	
Neural network	keras	keras_model_sequential	Yes	Number of layers (layer_dense) Number of neurons (units) Drop-out rates (layer_dropout) Activation function (activation) Loss function (loss), Optimizer (optimizer) Number of epochs (epochs), Batch size (batch size)	Ilayer_dense = 4 units = 84 -> 42 -> 20 -> 10 Ilayer_dropout = 0.3-0.4 activation = relu and sigmoid loss = binary_crossentropy optimizer = adam epochs = 10 batch_size = 50	

AUC is optimized via bootstrap in caret and cross validation in randomforestSRC and keras.

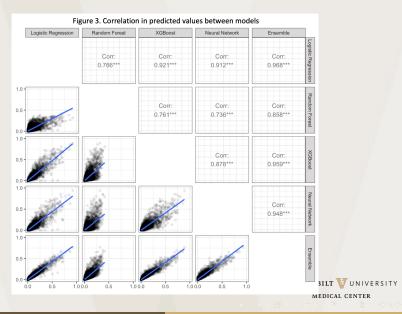
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Key results

Table 2. Performance comparison between models Derivation Validation cohort cohort Classifier C statistic C statistic ICI1 Calibration **Brier score** slope 0.896 0.894 0.0037 0.983 0.0122 Logistic regression Random forest 0.993 0.886 0.0030 0.951 0.0124 XG boost 0.908 0.893 0.0044 0.971 0.0122 Neural network 0.923 0.885 0.0048 0.969 0.0122 Ensemble method 0.975 0.901 0.0042 1.024 0.0121

¹ICI: Integrated Calibration Index

Key results



Key results

Table 3. Model accuracy measures at risk cut-off of 0.02

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Sensitivity	Specificity	PPV ¹	NPV ²	F1 ³						
0.690	0.904	0.092	0.995	0.162						
0.710	0.880	0.077	0.995	0.138						
0.670	0.914	0.098	0.995	0.171						
0.703	0.898	0.088	0.995	0.156						
0.719	0.900	0.092	0.996	0.163						
	0.690 0.710 0.670 0.703	Sensitivity Specificity 0.690 0.904 0.710 0.880 0.670 0.914 0.703 0.898	Sensitivity Specificity PPV¹ 0.690 0.904 0.092 0.710 0.880 0.077 0.670 0.914 0.098 0.703 0.898 0.088	Sensitivity Specificity PPV¹ NPV² 0.690 0.904 0.092 0.995 0.710 0.880 0.077 0.995 0.670 0.914 0.098 0.995 0.703 0.898 0.088 0.995						

¹PPV: Positive Predictive Value, ²NPV: Negative Predictive Value

F1 scores can range from 0 to 1, with 1 indicating a model with perfect classification.

³F1 score is the harmonic mean of PPV (precision) and sensitivity (recall).

Summary of the results

- All classifiers perform similarly well in terms of discrimination and calibration.
- More complex classifiers, known to handle interactions better, do not show improved performance compared to logistic regression.
- Random forest has a relatively low correlation with all other models, except for ensemble method. This is because its predicted probabilities cannot exceed 0.5.
- Model accuracy measures are more balanced when a cut-off of 0.02 (clinically meaningful cut-point) is used. Still, the sensitivity, PPV, and F1 are low due to a low prevalence of VTE.

3. Model Implementation (Topic 2)



Primary objectives

- Conduct a simulation study to evaluate imputation performance.
- Imputation involves replacing a missing value in the data with a reasonable estimate:
 - Fixed-value imputation: Median
 - Multiple imputation:
 - Data augmentation
 - Multivariate imputation by chained equations (MICE)
 - Conditional modeling
 - Joint modeling



Primary objectives

- Evaluate the performance of real-time VTE prediction models with imputed missing values.
 - Logistic regression
 - Random forest
 - XG boost
 - Neural network
 - Two-stage logistic regression model conditioning on a missing indicator M
 - Stage 1: Regression model conditioning on M = 1
 - Stage 2: Regression model conditioning on M = 0
 - This model does not require data imputation.



Simulation set-up

Define outcome Y and five predictors of interest, denoted as X_1, X_2, X_3, X_4, X_5 .

- Y: Fully observed binary outcome (VTE status)
- X₁: Fully observed binary non-lab variable (history of VTE)
- X_2 : Fully observed five-level categorical non-lab variable (type of admission)
- X₃: Fully observed continuous lab variable (BUN)
- X₄: Partially observed continuous lab variable (CRP)
- X_5 : Fully observed continuous (auxiliary) lab variable associated with missingness in X_4 (CL)
- M: Missingness in X_4 , where M = 1 if X_4 is absent



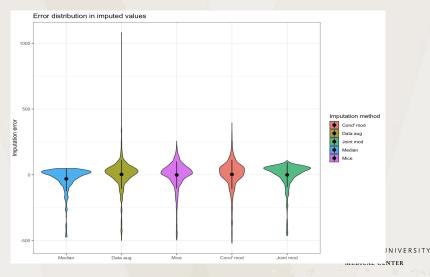
Simulation set-up

- Step 1: Generate data for a total population of 10,000 patients and compute true risks of each individual. Generate missing indicator (M) for all patients from a logistic distribution under MAR.
- Step 2: Develop five models on the true population with predictors X_1, X_2, X_3, X_4 and outcome Y.
- Step 3: Draw a sample of 100 patients (n_{sample}) from the total population. Draw 99 patients from non-missing group (M=0) and draw 1 patient from missing group (M=1).
- Step 4: Define a patient from the missing group as a new patient D^i . Impute the missing value of a new patient, D^i , using various imputation methods.
- Step 5: Predict risk of VTE for the patient i with the five models developed in Step 2.
- Step 6: Evaluate the performance of the imputation methods.
- Step 7: Compare the true risks obtained from the true population (Step 1) with the risk predictions based on imputation (Step 5). Compare performance and system time between different imputation methods and between five models. Repeat the step 3-7 1000 times.



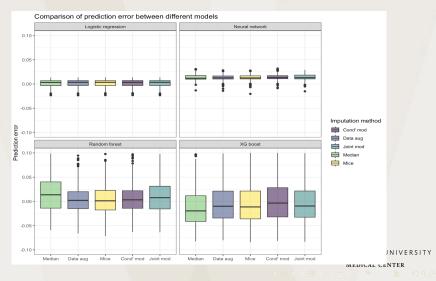
Simulation results

Event rate (0.1), Absence rate (0.1), MAR Imputation error = Imputed value - True value



Simulation results

Event rate (0.1), Absence rate (0.1), MAR Prediction error = Predicted risk - True risk



Simulation results

Event rate (0.1), Absence rate (0.1), MAR The units for time spent is seconds.

	Mean time spent for each imputation method									
Median	Data augmentation	MICE	Cond' modeling	Joint modeling						
0	1.382	1.588	1.393	1.393						



Summary of the results

- Median imputation: Errors from median imputation tend to center around 0 and exhibit much less variability compared to other imputation methods. However, median imputation does not work well with tree-based methods.
- Performance of imputation methods: While all imputation methods perform similarly, data augmentation shows the highest variability with some outliers. Despite this, its performance in prediction models is quite good.
- Logistic regression: Logistic regression has the lowest bias and variability, since it is the model used to generate the outcome in the simulation.
- Neural network: This model has low variability but tends to overestimate the probability of VTE.
- Random forest and XG boost: These tree-based models show higher variability in predictions overall. In particular, XG boost has the highest variability because it is more likely to overfit compared to robust bagging methods. Still, prediction errors in any imputation method other than median imputation are centered around 0.

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4. Model Evaluation (Topic 3)



Motivation

- Prediction models for VTE ignore potentially important information regarding the event.
 - Type of VTE: Pulmonary embolism, Deep vein thrombosis, Obstetric thromboembolism, etc.
 - Severity of VTE
- Additional information could facilitate more accurate prediction.
 - A model could have limited accuracy for all VTE.
 - The model might accurately predict a certain type of VTE.
 - The model might accurately predict more severe VTE.



Sketch of statistical approach

- Develop ROC-based summaries for predication accuracy
- Considering event timing
- Considering event severities on a continuous scale (marks)
- Extension of time-dependent prediction accuracy summaries introduced by Heagerty and colleagues

5. Future Plans

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Current progress and future plans

- Topic 1: Analysis is complete, and a manuscript is in preparation.
- Topic 2: Conduct a comprehensive simulation study considering a range of simulation settings.
- Topic 3: Development of an estimator and evaluate its properties.

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Thank You!

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