Série dos Seminários de Acompanhamento à Pesquisa

UEI Departamento De engenharia Industrial

Número 10 | 05 2021

Predicting the Acquisition of Resistant Pathogens in ICUs using Machine Learning Techniques

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Introduction

- Infections by antibiotic-resistant bacteria are one of the most significant current threats to global health;
- WHO: Gram-negative pathogens resistant to carbapenems are a critical priority;
- Increased attributable morbidity, mortality, hospitalization time, and economic costs;
- Mortality 1.78 times higher in patients with MDR-GNB infections;







Main

To develop models to predict CR-GNB acquisition in ICUs using machine-learning

techniques.

Specific Objectives

 To predict CR-GNB acquisition in ICUs, determining its risk factors, and assessing the impact on mortality rate using the logistic regression technique.

Paper:<u>https://www.journalofhospitalinfection.com/ar</u> ticle/S0195-6701(19)30182-3/fulltext

| | Journal of Hospital Infection 103 (2019) 121-127 | |
|---|--|--------------------------------------|
| | Available online at www.aciencedirect.com | |
| | Journal of Hospital Infection | Healthcare |
| SEVIER | journal homepage: www.elsevier.com/locate/jhin | - Sourcy |
| | | |
| redicting acqu | uisition of carbapenem-resistant | |
| ram-negative | pathogens in intensive care unit | ts |
| F. Dantas ^a . B. Dalm | as ^b , R.M. Andrade ^{c,d} , S. Hamacher ^a , F.A. Boz | za ^{e, f, *} |
| nes saint-etterne, Universite C pa D'Or Hospital, Rio de Janeir partment of General Medicine, Dr Institute for Research and E tional Institute of Infectious Di | ermoni Aldergne, Unes, Unes el la LUNG, Lentre CJ, sann-Ebenne, rri o, RJ, Brazil Federal University of the State of Rio de Janeiro, Rio de Janeiro, RJ, Bi duattan (1008), Rio de Janeiro, RJ, Brazil æase Evandro Chagas (IN), Oswaldo Cruz Foundation (FIOCRUZ), Rio de J | ance razil Janeiro, RJ, Brazil |
| RTICLEINFO | S U M M A R Y | |
| tcle history: | Background: Infections by multidrug-resistant Gram-negative (MD among the grantest contemporary health concerns, especially in it | PCN) bacteria are |

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Main

To develop models to predict CR-GNB acquisition in ICUs using machine-learning techniques.

Specific Objectives

2) Screening model to detects ICU patients who need to be tested;

3) Evaluating different machine learning and imbalanced learning method;

4) Developing a risk model that estimates ICU patients' probability of acquiring CR-GNB;

Differences between the models

| Model | Screening | Acquisition risk | |
|---|---|---------------------------------------|--|
| Туре | Discrimination/Classification | Prediction | |
| Study Population | All Screenings Tests | Screening Tests and Clinical Exams | |
| Unit of Analysis | Test | Patient | |
| Main Objective | To detect those who do NOT | To find the probability of each | |
| | need testing | patient to acquire the bacteria | |
| Sampling method | Different Balancing Strategies Matched Case-control | | |
| Hyperparameter Tuning Metric | AUC | Brier score | |
| Evaluation Metric | MCC and NPV | Brier score | |
| Interpretation | Error analysis (confusion matrix) | Calibration Belt | |
| Comparison of the techniques' performances | Yes | No | |
| Computational Time Analysis | Yes | No | |
| Analysis of the difference between hospitals | No | Yes | |
| Importance Factors | No | Yes | |
| Association Rules Mining | No | Yes | |

Study overview

• Hospitalized patients in 24 ICUs of five hospitals at a sizeable Brazilian network hospitals;

| - | Hospital | # ICUs | # ICU Beds | # Annual ICU admission |
|---|----------|--------|------------|------------------------|
| _ | Α | 1 | ~10 | ~600 |
| - | В | 2 | ~26 | ~1400 |
| - | С | 5 | ~52 | ~4500 |
| - | D | 9 | ~140 | ~5700 |
| | E | 7 | ~92 | ~5300 |

- The experiments were performed on an Intel® Core ™ i7 processor with 16GB of RAM and R 4.0.2 software;
- We used the CARET framework, imbalanced-learn packages, and others;
- We adapted all the functions of balancing strategies in CARET;

Database settings

- Epimed Monitor System®
 - Patient, ICU and hospital information;
 - Indexes (such as SAPS3 and Charlson);
 - Presence of comorbidities;
 - Use of the invasive devices during hospitalization;
 - Reasons for ICU admission;
- Business Intelligence (BI) System
 - Antibiotic data
- REAL system
 - Microbiology data \rightarrow Laboratory test results (positive/negative).

We included 44 new input attributes.

Conducting a Machine learning analysis

• Our analysis adds to the current studies in four respects: machine learning techniques, balancing strategies, feature selection, and performance evaluation;



• Machine Learning Techniques and Balancing strategies

| Method | Algorithm |
|---|-------------|
| LINEAR CLASSIFICATION MODELS | |
| Logistic Regression | glm |
| Logistic Regression with regularization | glmnet |
| Linear Discriminant Analysis (LDA) | lda |
| Nearest Shrunken Centroids (NSC) | pam |
| Support Vector Machine (SVM) - Linear | svmLinear |
| NONLINEAR CLASSIFICATION MODELS | |
| Neural Network | nnet |
| Support Vector Machine (SVM) - Radial | svmRadial |
| k-Nearest Neighbors (kNN) | kNN |
| Naive Bayes | naive_bayes |
| CLASSIFICATION TREES | |
| Decision Tree C45 | J48 |
| Decision Tree CART | rpart |
| Decision Tree C50 | C5.0 |
| Random Forest (RF) | rf |
| Gradient Boosting Machines (GBM) | gbm |
| Bagging | treebag |
| AdaBoost | AdaBoost.M1 |

Balancing approaches

SAMPLING

Random downsampling (or undersampling)

Random upsampling (or oversampling)

SMOTE

DATA CLEANING TECHNIQUES

Tomek Links

Neighbourhood Cleaning Rule (NCL)

One-sided selection (OSS)

SMOTE + Tomek

SMOTE + NCL

SMOTE + OSS

ENSEMBLE-BASED METHODS

SMOTEBoost

RUSBoost

SMOTEBagging

UnderBagging

Differences between the models

| Model | Screening | |
|------------------------------------|--------------------------------|--|
| Туре | Discrimination/Classification | |
| Study Population | All Screenings Tests | |
| Unit of Analysis | Test | |
| Main Objective | To detect those who do NOT | |
| | need testing | |
| Sampling method | Different Balancing Strategies | |
| Hyperparameter Tuning Metric | AUC | |
| Evaluation Metric | MCC and NPV | |
| Interpretation | Error analysis (confusion | |
| | matrix) | |
| Comparison of the techniques' | Vec | |
| performances | 1 63 | |
| Computational Time Analysis | Yes | |
| Analysis of the difference between | No | |
| hospitals | | |
| Importance Factors | No | |
| Association Rules Mining | No | |
| | | |

Prediction Screening Tests and Clinical Patient Fo find the probability of each patient to acquire the bacteria Matched Case-control Study No No

Setting and Study population

Inclusion criteria:

- Tests realized between 48h and 60days after patient admission and made in adult ICUs.
- Testing in patients with admission date after May 8th, 2017 until August 31st, 2019;
- Patients aged ≥18 years old;

- Total: 3,911 tests

- Positive = 394
- Negative = 3,517



Descriptive analysis

- Patients more likely to be colonized:
 - High length of stay in hospital or ICU;
 - Higher severity indices;
 - Antibiotics use;
 - Invasive devices between 24 hours to 30 days before the test;
 - Prolonged use time of mechanical ventilation and catheters;
 - Higher duration that a procedure is used between one test and another and the number of times they were changed;
 - Admitted from sepsis/infection or neurological disease and by the operation room or other ICU from the hospital;

Setting and Study population

| Hospital | #Screening Tests | #Positive Tests | #Negative Tests | % Positive Tests |
|----------|---------------------|--------------------|--------------------|------------------------|
| Α | 310 | 60 | 250 | 19.4% |
| В | 806 | 57 | 749 | 7.1% |
| С | 1081 | 81 | 1000 | 7.5% |
| D | 1714 | 196 | 1518 | 11.4% |
| All | 3911 | 394 | 3517 | 11.3% |



Data Preprocessing

Feature Selection

| | Mean AUC values | | | | Number of | |
|--|-----------------|------------|-------|-------|-----------|-----------|
| | C45 | SVM Radial | KNN | LR | AK | variables |
| Recursive Feature Elimination with random forest (RF-RFE) | 0.632 | 0.690 | 0.642 | 0.713 | 1.25 | 35 |
| Selection by Filter (SBF) | 0.624 | 0.674 | 0.625 | 0.709 | 2.75 | 42 |
| Class Decomposition with filter (D.SBF) | 0.568 | 0.658 | 0.641 | 0.702 | 3.75 | 76 |
| Class Decomposition with random forest (D.RF) | 0.607 | 0.687 | 0.658 | 0.708 | 2.25 | 24 |
| Friedman test (p-value) | | 0. | 007 | | | |

- We proposed an approach combining feature selection and cluster techniques: D.RF.
 - The second best when comparing AUC;
 - The best one to discriminate the positive classes comparing the Sensitivity;

Building model – Training



✓ Evaluating different machine learning and imbalanced learning method

NPV

Friedman chi-squared \rightarrow p-value < 0.001



✓ Evaluating different machine learning and imbalanced learning method

MCC



✓ Computational Time

Timing Final Model

- The sampling strategies have the lowest medians, followed by data cleaning strategies;
- Tree-based strategies take longer to build the final model;
- The linear models are more efficient, followed by decision trees;
- The SVM Radial and Adaboost are the slowest;

| Strategies | Strategies Timing Everything (min) Tim Median | |
|--------------|--|------|
| Downsampling | 24.1 | 0.1 |
| Upsampling | 40.9 | 0.5 |
| OSS | 43.3 | 0.9 |
| SMOTE | 67.0 | 0.4 |
| Tomek | 74.4 | 0.5 |
| UnderBagging | 75.3 | 2.4 |
| NCL | 121.4 | 1.3 |
| SMOTE_Tomek | 125.1 | 0.7 |
| SMOTE_OSS | 135.2 | 0.9 |
| SMOTE NCL | 157.6 | 1.3 |
| RUSBoost | 1001.9 | 6.5 |
| SMOTEBoost | 1402.7 | 9.3 |
| SMOTEBagging | 1474.3 | 10.3 |

| Methods | Timing Everything (min) Median | Timing Final Model (min) Median |
|-------------------|-----------------------------------|------------------------------------|
| LDA | 6.1 | 0.5 |
| LR | 6.4 | 0.5 |
| NSC | 6.5 | 0.5 |
| CART | 7.6 | 0.6 |
| BAGGING | 9.8 | 0.9 |
| C50 | 39.2 | 1.6 |
| KNN | 44.8 | 0.5 |
| LR_regularization | 51.1 | 0.6 |
| SVM_LINEAR | 180.8 | 4.2 |
| RF | 292.4 | 2.0 |
| NN | 297.0 | 1.0 |
| NB | 319.8 | 0.8 |
| C45 | 413.5 | 0.6 |
| ADABOOST | 451.8 | 11.8 |
| GBM | 815.5 | 2.4 |
| SVM_RADIAL | 1562.2 | 3.6 |

✓ Model Analysis

- We analyze each false-negative case found in the confusion matrices of the best-classifiers using the

781 (78 positives and 703 negatives) data test;

CONSERVATIVE MODEL (BY NPV) - NB, RF, and LR regularized

| RF (downsampling) | | | | |
|-------------------|------|------|------|------|
| Sens | Spec | PPV | NPV | AUC |
| 0.92 | 0.39 | 0.14 | 0.98 | 0.75 |
| Reference | | | | |
| | | Pos | Neg | |
| Dradiatad | Pos | 72 | 429 | |
| Predicted | Neg | 6 | 274 | |

✓ Reduce 280 tests (36%), but 6 patients non-isolated;

MODERATE MODEL (BY MCC) - NN and SVM Radial

| NN (SMOTE+Tomek) | | | | | | |
|------------------|-----------|------|------|------|--|--|
| Sens | Spec | PPV | NPV | MCC | | |
| 0.76 | 0.64 | 0.19 | 0.96 | 0.24 | | |
| | Reference | | | | | |
| | | Pos | Neg | | | |
| Prodicted | Pos | 59 | 251 | - | | |
| | Neg | 19 | 452 | - | | |

✓ Reduce 471 tests (64%), but 19 patients non-isolated;

Differences between the models

| Model | | Acquisition risk |
|------------------------------------|--------------------------------------|---------------------------------|
| Туре | Discrimination/Classification | Prediction |
| Study Dopulation | All Sereeninge Teete | Screening Tests and Clinical |
| Study Population | All Screenings rests | Exams |
| Unit of Analysis | Test | Patient |
| Main Objective | To detect those who do NOT | To find the probability of each |
| Main Objective | need testing | patient to acquire the bacteria |
| Sampling mothod | Different Balancing | Matched Case control Study |
| Sampling method | Strategies | Matched Case-control Study |
| Hyperparameter Tuning Metric | AUC | Brier score |
| Evaluation Metric | MCC and NPV | Brier score |
| Interpretation | Error analysis (confusion | Calibration Balt |
| Interpretation | matrix) | Calibration Beit |
| Comparison of the techniques' | Vac | No |
| performances | 165 | INO |
| Computational Time Analysis | Yes | No |
| Analysis of the difference between | Ne | Voc |
| hospitals | NO | 165 |
| Importance Factors | No | Yes |
| Association Rules Mining | No | Yes |



✓ Setting and study population

| Hospital | # Tests | # Negative Tests | # Positive Tests | # Patients |
|----------|---------|---------------------|------------------|------------|
| A | 404 | 341 | 63 | 214 |
| В | 1,039 | 971 | 68 | 469 |
| С | 1,540 | 1,452 | 88 | 611 |
| D | 3,849 | 3,616 | 233 | 1,658 |
| E | 1,157 | 1,082 | 75 | 652 |
| All | 7,989 | 7,462 | 527 | 3,604 |

- Unit of analysis \rightarrow Patient;
- A matched case-control design by the hospital and admission date (3:1);

✓ Database Preparation



✓ Model building and evaluation



✓ Model Building and Evaluation

- General Model

- NSC is the best model to estimate CR-GNB acquisition risk;
- NSC, GBM, CART, LR, LR regularized, and LDA are calibrated models, suitable for prediction;
- NB, Bagging, and RF overestimate the colonization for medium and high-risk patients and underestimates low-risk patients;
- The NB is out almost the whole diagonal line and presented the worst Brier score.

| | Brier score | | | Confidence level (80%) | | |
|----------------|-------------|-------|---------|------------------------|-------------------|--|
| Methods | | MCC | p-value | Under the bisector | Over the bisector | |
| NSC | 0.152 | 0.327 | V | V | v | |
| GBM | 0.159 | 0.312 | V | V | V | |
| CART | 0.167 | 0.379 | V | V | V | |
| LR | 0.163 | 0.338 | V | V | V | |
| LR regularized | 0.155 | 0.318 | V | V | X | |
| LDA | 0.159 | 0.327 | V | V | V | |
| SVM RADIAL | 0.171 | 0.109 | X | V | X | |
| C45 | 0.165 | 0.383 | X | X | X | |
| NN | 0.160 | 0.335 | X | V | X | |
| ADABOOST | 0.172 | 0.295 | X | X | V | |
| C50 | 0.160 | 0.399 | X | V | X | |
| kNN | 0.173 | 0.296 | X | V | X | |
| RF | 0.176 | 0.326 | X | X | X | |
| BAGGING | 0.183 | 0.308 | X | X | X | |
| SVM LINEAR | 0.177 | 0.345 | Χ | X | X | |
| NB | 0.196 | 0.339 | X | X | X | |

- ✓ Model Building and Evaluation
 - General Model



NEAREST_SHRUNKEN_CENTROIDS

- No evidence of the lack of calibration emerges from the calibration belt;
- The model calibration on the development is
 acceptable (p-value = 0.440).

- ✓ Model Building and Evaluation
 - Model by hospital

<u>Objective</u>: To understand if the built general model can be used for all hospitals;

- There is not a significant difference between general and the five individual models by Brier scores;
 - T-test \rightarrow P-value = 1

We can use the general model for all hospitals without losing performance;

External Validation

• Our final risk model for the acquisition of CR-GNB is the NSC;

| Hospital | # Patient | #Positive | #Negative | % Positive Tests | Brier Score | MCC |
|--|-----------|------------------|-----------|--------------------------|-------------|-------|
| F | 267 | 39 | 228 | 14.61% | 0.128 | 0.261 |
| G | 357 | 34 | 323 | 9.52% | 0.079 | 0.261 |
| All hospitals - General Model (testing set) | 413 | 105 | 308 | Case-control study (3:1) | 0.152 | 0.327 |

- The model does not classify well the non-acquisition of CR-GNB (MCC = 0.261) but can predict the probability of acquiring (Brier score = 0.128 and 0.079);
- The model is well-calibrated and acceptable to be introduced at Hospital G;
- Hospital F model overestimates the colonization of patients;



✓ Important of variables

• We identified the attribute importance by Information Gain;



Attribute Importance - All hospitals

- Duration and use of invasive devices, especially mechanical ventilation;
- Antibiotic groups;
- Admission Source and Admission Reason;
- Criticality indices such as Saps3;
- Length of stay before test.

✓ Association Rules

- We extracted a list of 157 association rules with predictive value "positive";
- Example:

| # | Rules | Support | Confidence | Lift |
|---|--|---------|------------|-------|
| 1 | {MVDURTOTAL=[4,57],VesDURTOTAL=[6,58],J01D=TRUE,Antibiotic=TRUE,VESICAL=YES} => {RESULT=pos} | 0.100 | 0.575 | 2.257 |

If a patient is hospitalized with these conditions, this patient has a 57.5% probability of acquisition;

• All the conditions selected include some information about invasive dispositive use;

✓ Main Findings – Screening Model

- SMOTEBagging and UnderBagging approaches obtained better results than the data cleaning;
- The more straightforward linear techniques is not significantly different from the more complex classifiers;
- Screening models:
 - Conservative: Random forest \rightarrow the unnecessary test is avoided **39%** and **8%** of false-negatives.
 - Moderate: Neural Network \rightarrow the unnecessary test is avoided 64% and 24% of false-negatives.

✓ Main Findings – Risk Model

- NSC is the best model to estimate acquisition risk;
- Naïve Bayes technique has better discrimination power but the worst Brier score value;
- We can use the general model for all hospitals without losing performance;
- The variables related to the duration of the use of invasive devices, especially mechanical ventilation, are the most important;

Contributions

Literature

- A literature review on prediction in the healthcare context, focusing on multi-resistant bacteria acquisition;
- Evaluation of the different machine learning techniques and balancing strategies;

Methodological

- A framework about "how to conduct a machine learning analysis";
- Combination of feature selection and cluster techniques;
- An approach to screening modeling considering weekly tests and variables that consider actions that happened between one test and another;

Contributions

Applied

- Rules of strongly associated features that indicate that a patient is at risk of acquired CR-GNB;
- Two screening models: one more conservative and the other moderate;
- A risk model for the acquisition of CR-GNB;

✓ Limitations

- These results cannot be directly extrapolated to other healthcare institutions;
- Heterogeneous Gram-negative bacteria were analyzed collectively;
- We do not know precisely how the patient acquired the bacteria;
- Patients with the same conditions may have different types and timing of observations;
- Some records may be lost due to data imputation human errors.

✓ Future Researches

- To perform an external validation using the best screening model in new periods and hospitals;
- To develop time-series models considering variable changes over time;
- To compare the relationship of antibiotic use between the periods before and during the pandemic;
- To analyze the influence of acquisition for the patient's outcome within 30 days after a positive test using a survival model.

✓ Final Consideration

- Identifying patients who don't need a weekly culture test decreases hospital costs and laboratory waiting times;
- The models for predicting resistance can offer utility where rapid diagnostics are unavailable or resource impractical;
- Infection control policies can be established to control the spread of these bacteria;
- The framework on how to conduct a machine learning analysis and the code developed can be reusable and easily adaptable;

Publications

Articles in Scientific Journals

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Articles in R1 - Submitted

App-based symptom tracking to optimize SARS-CoV-2 testing strategy using machine learning \rightarrow PLoS One Socio-demographic factors associated with COVID-19 in-hospital mortality in Brazil \rightarrow Public Health

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Awards and Titles

2017 Best report Academic Category, BPI Challenge, Business Process Intelligence Workshop.

Expanded Summary published in proceedings of conferences

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THANK YOU!!

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