

Random Forest vs Logistic Regression for Binary Classification

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Abstract. Selecting a learning algorithm to implement for a particular application on the basis of performance still remains an ad-hoc process using fundamental benchmarks such as evaluating a classifier’s overall loss function, area under the curve (AUC) score, specificity, and sensitivity values. This work is aimed at addressing the difficulty of model selection by evaluating the overall classification performance between random forest and logistic regression for datasets comprised of various underlying structures. A model evaluation tool was developed in R for simulating a variety of dataset characteristics in order to evaluate performance metrics such as true positive rate, false positive rate, and accuracy under specific conditions. Our findings indicate that when increasing the variance in the explanatory and noise variables, logistic regression consistently performed with a higher overall accuracy as compared to random forest. However, the true positive rate for random forest was higher than logistic regression and yielded a higher false positive rate. In all cases a paired two sample t-test indicates there is enough evidence to suggest the false positive rate for random forest is statistically different than logistic regression. The model evaluation application developed in this work is a foundation for answering other intriguing questions related to model performance under various treatments.

1 Introduction

Datasets consist of various shapes and compositions, which poses the question, what data characteristics result in one machine learning algorithm outperforming others. Both quantitative and qualitative data can be a combination of categorical and numerical multivariates that can range from a variety of entities. When it comes to model selection, this is a constant challenge data scientist are often faced with. Training and evaluating multiple machine learning models for a given use case can be an expensive task computationally as well as time consuming. The analysis presented in this work is to provide insight into the relative performance of a learning algorithm conditioned on the particular characteristics of a dataset. To do so, random forest and logistic regression are the two models that will be analyzed for comparing binary classification ($y = 0,1$) performances with

differing dataset structures. Both classifiers have been widely implemented in various domains and their successes have been well documented. However, the particular characteristics of a dataset that make one model outperform the other is unknown as most published work compares overall performance between the two models for a single dataset. The objective of this study is to develop a statistical tool to directly observe the classification performance of each model by averaging metrics for 1000 random generations of assorted multivariate datasets. Classification metrics, such as accuracy, area under the curve, true positive rate, false positive rate, and precision are examined for performance. Finally, a pairwise two-sample t-test is conducted at the end of each simulation case study in order to provide statistical quantification as to whether a difference in model performance is conclusive enough to state the difference is significant or if the observed difference is by random chance.

Characteristics of a dataset can be comprised of missing values, outlier, highly correlated variables, concave or convex shapes, or subsets of the data that can be represented as clusters. Complex datasets that are not linearly separable, or in other words, where a linear hyperplane splits the data into two halves such that the model poorly predicts the class label of the respected observation (Figure 1). One approach to inferring underlying complexities of high-dimensional dataset is Topological Data Analysis (TDA). TDA is an evolving method that utilizes topological and geometric tools to identify relevant features in the data. TDA can be described as method that helps identify structures in noisy and incomplete datasets like clusters or other hidden shapes that can provide a more accurate representation of the dataset (Chazal et al, 2017). Models can then be trained on the new representation of the data that has been reconstructed, which has shown promising results. While TDA looks at the proximity of data points and connectivity that can be mapped to a 1-dimensional plane for representing the shape of the data (Munch,2017), our analysis is aimed at creating various complexities in the data and evaluating model performance on the raw structure in a multidimensional space. For instance, altering the variance in the explanatory and noise variables, changing the number of observations, and varying the number of continuous features included in a multivariate dataset was considered. Investigation into why models like logistic regression or random forest perform differently for simple and complex data characteristics was the motivation behind this work.

The data examined in this work is only for continuous variables that have a Gaussian distribution. However, more complex structures can exist like the toy dataset shown below in Figure 1 and Figure 2 which consists of concave and convex shape. As illustrated in these figures, both random forest and logistic regression nearly establishes an identical decision boundary for the left-hand side dataset. Alternatively, logistic regression underperforms random forest and yields a higher misclassification rate, which raises a profound question as to which data characteristics constitutes one model achieving an overall better classification score. It should be noted this work only investigates random forest and logistic regression, however generalization of the current application can be adapted

to other linear and nonlinear models. As previously alluded, performance of machine learning classifiers can yield varying results depending on the shape and structure of the data

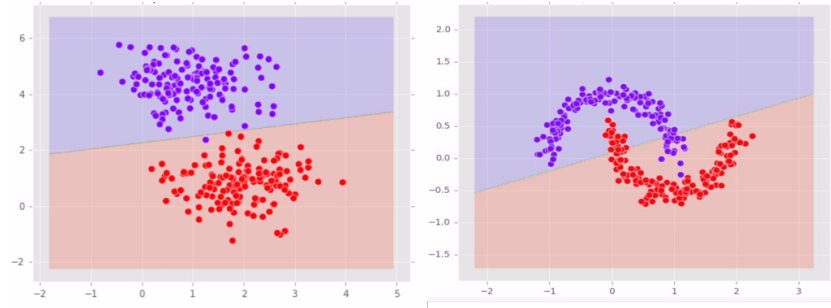


Fig. 1. Logistic Regression Decision Boundary

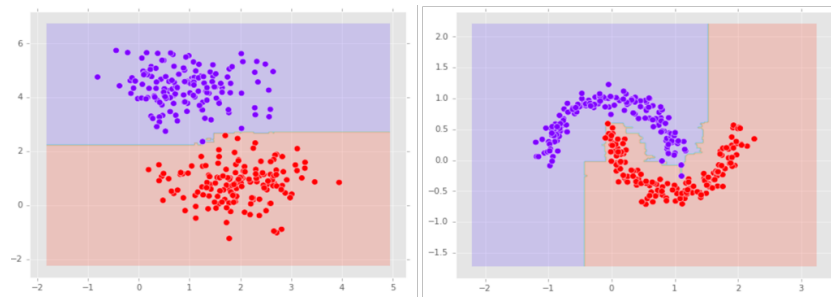


Fig. 2. Random Forest (n-Trees = 100) Decision Boundary

Numerous studies have been published that compare random forest and logistic regression algorithms however, most research experiments consisted of either a single dataset or multiple datasets from the same source. In these scenarios, sometimes logistic regression performed better while in other cases random forest performed better. For example, one experiment used several neuropsychological tests to predict dementia stated that with respect to specificity and overall classification accuracy, random forests and linear discriminant analysis rank first among all the classifiers including logistic regression (Guerreiro, 2011). Contrastingly, another article analyzing Twitter tweets surrounding the 2016 United States election concludes that when Principal Component Analysis (PCA) is applied to tweets, logistic regression provides better results than random forest (Begenilmis, 2017). The type of data and data sources used in the studies above are drastically different from each other and each algorithm performs differently

due to the type of data it was utilizing to train the classifiers. This analysis aims to provide a method of evaluating random forest and logistic regression models for a variety of data conditions.

2 Analytical Tool

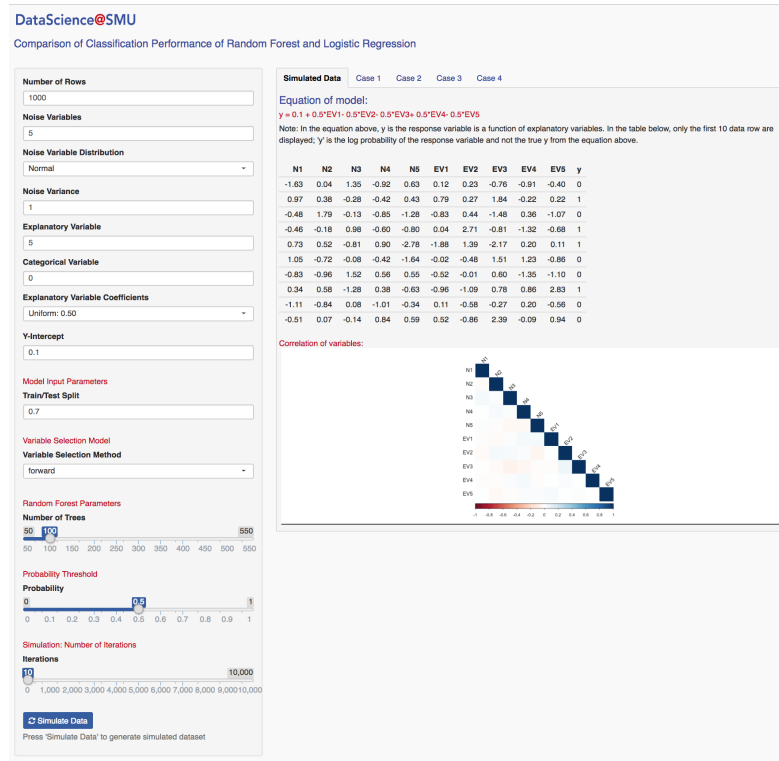


Fig. 3. R Shiny Application - Data Simulator

To conduct the statistical analysis, an interactive web application was developed using RShiny which allows end users to rapidly generate simulated datasets and evaluate performance metrics between the machine learning models. This application is shown in Figure 3 and allows several input options to be configured prior to generating a multivariate dataset of an arbitrary length such as specifying number of observations in the dataset, variances in the features, amount of noise and explanatory variables, and the distribution of input features as either Gaussian or Poisson. Moreover, the user has the ability to choose how the parameter estimates are weighted (e.g. uniform weights or unbalanced), allowing for a subset of the explanatory variables to be a more significant predictor of the

response variable than others. The left column of the tool also has configurations for the machine learning models such as setting the number of trees for random forest or specifying the percentage of the training and testing set splits.

Simulated Data										
Case 1 Case 2 Case 3 Case 4										
Equation of model:										
$y = 0.1 + 0.5 \cdot EV1 + 0.5 \cdot EV2 + 0.5 \cdot EV3 + 0.5 \cdot EV4 + 0.5 \cdot EV5$										
Note: In the equation above, y is the response variable is a function of explanatory variables. In the table below, only the first 10 data row are displayed; 'y' is the log probability of the response variable and not the true y from the equation above.										
N1	N2	N3	N4	N5	EV1	EV2	EV3	EV4	EV5	y
1.24	0.73	1.84	2.11	-0.55	-2.25	-0.41	-0.64	0.86	-1.48	1
0.12	-0.63	1.07	-0.32	0.30	-1.06	1.85	0.32	-1.71	0.40	0
1.01	0.64	0.11	1.23	0.07	-0.94	-2.64	0.29	-0.02	-0.64	0
-2.13	0.47	-0.38	-0.46	-1.99	-0.44	0.35	-0.19	-0.42	1.01	0
1.62	0.45	-1.33	-0.24	1.89	-0.10	-0.36	0.19	0.65	0.45	1
-0.57	-0.05	-1.12	0.46	-0.02	-0.11	-0.06	1.08	-1.55	-1.18	0
1.28	1.57	1.13	-0.55	-1.21	1.46	0.53	-0.29	1.15	-0.09	1
0.43	-0.78	-0.19	0.40	0.53	2.06	0.52	-1.19	1.48	1.55	0
-1.80	0.29	-0.59	-0.45	0.16	-2.29	0.07	-0.18	-1.41	-0.45	0
-1.76	-0.41	-0.41	0.06	0.35	-0.67	0.01	-0.47	-1.36	0.23	0

Fig. 4. Equation of the response and first 10 rows of the simulated dataset

For performing numerical simulations, creating synthetic datasets is pivotal for the analysis. SimStudy is an open source package in the R programming language. This was leveraged in this work as the method for producing datasets of various structures. Given this work is aimed at model performance for binary classification, the response variable 'y' is a function of only the explanatory variables 'x' included in the model equation (Eq. 1) and displayed in the tool in Figure 4. Figure 4 is a screenshot of the first tab in the center content of the tool. It displays the equation as well as the first 10 rows of the dataset with all columns of 'EV', 'N', and the response 'y'. The binary response variables take on the value of either a 1 or 0; thus the formula represents the log of odd or probability of the response being a 1 or 0. As previously stated, the explanatory variable 'EV' is related to the binary response, while the noise variables 'N' are not. The parameter estimates explain the relationship between independent variables 'X' and the dependent variable 'Y', and the 'Y' scale is known as the logit, or log of odds. For each simulation case study explored in the work, the default parameter estimate, beta, is set to a uniform 0.50 and the input features, both noise and explanatory variables, are all continuous and normally distributed.

$$\log(y) = N_0 + \beta_1 X_1 + \dots \beta_n X_n + N_n \quad (1)$$

The rest of the tabs in the tool are for the case studies that we explored in this work. On each tab, there is a description of what each case is simulating and the results of running random forest and logistic regression predictions on the simulated data. The average of the evaluation metrics are summarized in a table and line charts and a spread of the evaluation metrics are seen in a boxplot. Figure 5 is an example of the summary, table of average evaluation metrics for each model, and a two sample t-test for the difference in models.

Simulated Data

Case 1

Case 2

Case 3

Case 4

Case Description

In this case, we simulate the dataset and set the number of explanatory variables to 1, 5, 10, 20, and 50 and display the results of each simulation.

Logistic Regression Results

The table below displays the results of running logistic regression 10 times with 5 noise variables and 1, 5, 10, 20, and 50 explanatory variables in the dataset.

algorithm	fpr	tpr	rec	prec	acc	auc	num_ev
Logistic Regression	0.59	0.61	0.54	0.61	0.52	0.56	1.00
Logistic Regression	0.41	0.59	0.58	0.59	0.59	0.63	5.00
Logistic Regression	0.41	0.69	0.65	0.69	0.65	0.67	10.00
Logistic Regression	0.31	0.72	0.71	0.72	0.70	0.78	20.00
Logistic Regression	0.34	0.61	0.65	0.61	0.64	0.69	50.00

Random Forest Results

The table below displays the results of running random forest 10 times with 5 noise variables and 1, 5, 10, 20, and 50 explanatory variables in the dataset.

algorithm	fpr	tpr	rec	prec	acc	auc	num_ev
RandomForest	0.54	0.69	0.58	0.69	0.59	0.59	1.00
RandomForest	0.40	0.69	0.64	0.69	0.65	0.63	5.00
RandomForest	0.58	0.77	0.61	0.77	0.62	0.70	10.00
RandomForest	0.45	0.68	0.61	0.68	0.62	0.68	20.00
RandomForest	0.42	0.68	0.62	0.68	0.64	0.68	50.00

Paired Two Sample T-Test Results

fpr	tpr	rec	prec	acc	auc	metric
0.19	0.073	0.662	0.073	0.932	0.659	p-value

Fig. 5. Example case study tab

3 Methods and Experiments

The two machine learning algorithms studied in this work consist of random forest and logistic regression. Both models have been widely implemented in various disciplines for classification and regression purposes. Not only are these algorithms known for their success, but also their simplicity to implement and relatively straightforward to interpret. The functionality of logistic regression,

a parameter based model, and random forest, a non-parametric model, are discussed in the following section.

3.1 Random Forest

Random forest is an ensemble-based learning algorithm which is comprised of 'n' collection of de-correlated decision trees (Hastie, 2009). It is built off the idea of bootstrap aggregation, which is a method for resampling with replacement in order to reduce variance. The algorithm uses multiple trees to average (regression) or compute majority votes (classification) in the terminal leaf nodes when making a prediction. Presented by Leo Breiman and built off the idea of decision trees, random forest models resulted significant improvements in prediction accuracy as compared to a single tree by growing 'n' number of trees where each tree in the training set is sampled randomly without replacement (Breiman, 1966). Decision trees consist simply of a tree-like structure where the top node is considered the root of the tree that is recursively split at a series of decision nodes from the root until the terminal node or decision node is reached.

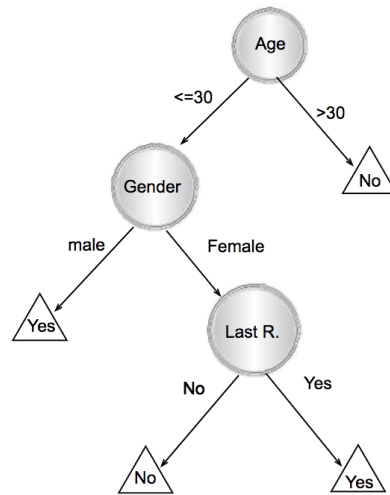


Fig. 6. Example of Decision Tree (Lior Rokach et al)

As illustrated in the tree structure, the decision tree algorithm is a top down greedy approach by partitioning the dataset into smaller subsets. The result is a tree with a series of decision nodes and leaf node. The decision node has two or more branches where the features with the highest information gain is split. The predictor variable that yields the highest information gain is the root node. The leaf node is represented as a prediction, and in this case, classifying either

1 or 0. Decision trees can handle both categorical and numerical data. First, in order to determine the information gain which is based on the entropy after splitting on an attribute, entropy is computed. Information gain is based on the principles from information theory that uses entropy to compute impurity of datasets. Entropy measures the homogeneity of the subset data; if entropy equals one then the class labels are equally divided while an entropy of zero means the sample is completely homogeneous.

$$Entropy = -p \log_2(p) - q \log_2(q) \quad (2)$$

Advantages of using a tree-like learning algorithm allow for training models on large datasets in addition to quantitative and qualitative input variables. Additionally, tree-based models can be immune to redundant variables or variables with high correlation which may lead to overfitting in other learning algorithms. Trees also have very few parameters to tune for when training the model and performs relatively well with outliers or missing values in a dataset. However, trees are prone to poor prediction performance; decision trees themselves are prone to overfitting noise in a training set which ultimately leads to results with high variance. In other words, this means the model could accurately predict the same data it was trained on but may not possess the same performance on datasets without the similar patterns and variations in the training set. Even fully grown decision trees are notorious for overfitting and do not generalize well to unseen data; random forest solves the overfitting conundrum by using a combination or "ensemble" of decision trees where the values in the tree are a random, independent, sample.

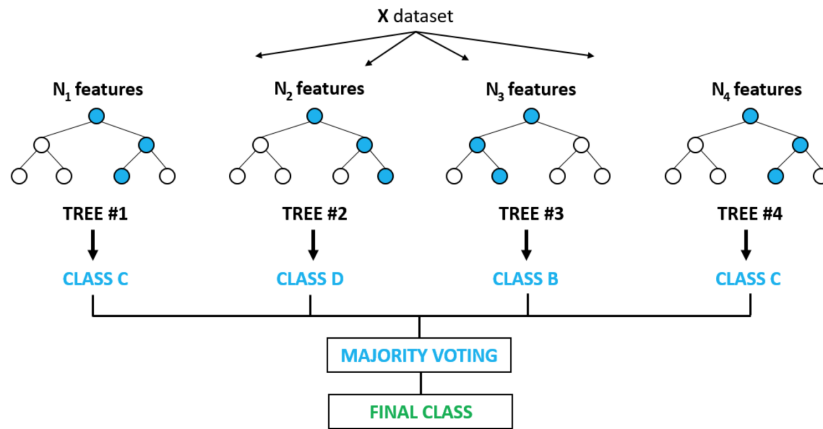


Fig. 7. Example of Decision Tree (Balazs Holczer)

The idea of randomly sampling the without replacement is known as bagging and this results in a different tree being generated to train on; averaging the results from the 'n' number of trees will result in decreasing the variance and establishing a smoother decision boundary (Hastie, 2009). For instance, while using random forest for classification, each tree will give an estimate of the probability of the class label, the probabilities will be averaged over the 'n' trees and the highest yields the predicted class label. In addition to bagging or bootstrap aggregation, in order to further reduces the variance in the decision boundary further, the trees must be completely uncorrelated and the method of bootstrapping alone is not enough. Breiman introduced the idea of randomly sampling 'm' number of features at each decision split in the tree as a way to de-correlate the trees in the random forest algorithm.

3.2 Logistic Regression

Linear models are composed of one or multiple independent variables that describes a relationship to a dependent response variable. Mapping qualitative or quantitative input features to a target variable that is attempted to being predicted such as financial, biological, or sociological data is known as supervised learning in machine learning terminology if the labels are known. One of the most common utilized linear statistical models for discriminant analysis is logistic regression.

$$\pi_i = \beta_0 + \beta_1 X_1 + \beta_n X_n \quad (3)$$

Simplicity and interoperability of logistic regression can occasionally lead to outperforming other sophisticated nonlinear models such as ensemble learners or support vector machines. However, in the event the response variable is drawn from a small sample size, then logistic regression models become insufficient and performs poorly for binary responses A number of learning algorithms could be applied to modeling binary classification data types, however the focal point of this work is to examine one linear model, logistic regression.

Unlike the response variable for logistic regression which is quantitative, the target variable for logistic regression is the posterior probability of being classified in the ith group of a binary or multi-class response (Hastie, 2009). Logistic regression makes several assumptions such as independence, responses (logits) at every level of a subpopulation of the explanatory variable are normally distributed, and constant variance between the responses and all values of the explanatory variable. Intuitively, a transformation to the response variable is applied to yield a continuous probability distribution over the output classes bounded between 0 and 1; this transformation is called to “logistic” or “sigmoid” function where ‘z’ corresponds to log odds divided by the logit (Ng, 2008). The parameter estimates inform whether there is an increase or decrease in the predicted log odds of the response variable that would be predicted by one unit increase or decrease in one of the explanatory variables (e.g. x1), while holding all other explanatory variables constant.

$$\sigma(Z) = \frac{1}{1 + \exp^{-z}} \quad (4)$$

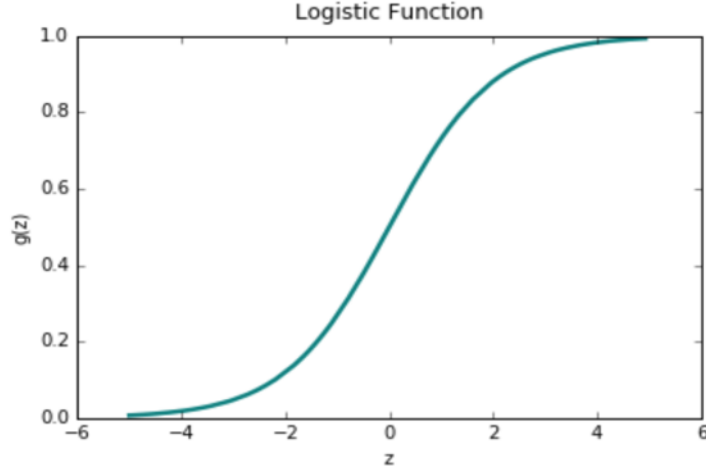


Fig. 8. Logistic Function

For a binary response, the logistic regression model can be expressed by summing over the linear combinations of input features and a corresponding weight plus a bias term for each instance as shown below in equation (3) and (4).

$$p(y^{(i)} = 1|x^{(i)}, w) = \frac{1}{1 + \exp^{-(w^T x^{(i)} + b)}} \quad (5)$$

$$p(y^{(i)} = 0|x^{(i)}, w) = \frac{1}{1 + \exp^{(w^T x^{(i)} + b)}} \quad (6)$$

The objective is to find a set of weights such that the negative log likelihood is minimized over the defined training set using optimization techniques such as gradient descent or stochastic gradient descent [3]. Minimizing the negative log likelihood also means maximizing the likelihood or probability the parameter estimate, π , of selecting the correct class. The loss function that measures the difference between the ground truth label and the predicted class label is referred to as the cross-entropy. If the prediction is very close to the ground truth label, the loss value will be low. Alternatively, if the prediction is far from the true label, the resulting log loss will be higher.

$$J(\theta) = -\frac{1}{m} \sum p_i \log(y_i) + (1 - p_i) \log(1 - y_i) \quad (7)$$

4 Criteria for Model Comparison

When comparing overall model performance, accuracy, true positive rate, false positive rate, precision, recall, and AUC were considered as the core metrics.

For each simulation case, the dataset was randomly partitioned into 70% being utilized to train the model while the remainder 30% is for testing on out-of-sample data. First metric is accuracy, the percentage of correct classification. If classified as a success, how often does the model predict success and if it was a failure, how often is a failure predicted. Next, true positive rate and sensitivity are calculated as the portion of positives or successes that are correctly identified. On the other hand, false positive rate is the portion that was incorrectly identified as positive or success but is actually negative. Depending on the application and domain, one may care about incorrectly classifying a positive more than incorrectly classifying a negative. For example, when dealing with anything medical or health related, such as predicting if a patient will have dementia, it is extremely important to have a low false positive rate because telling someone they have dementia when they do not can cause a lot of emotional stress. False positive rate is also important in quality fields where cost is high. To test a silicon wafer for defects for instance, a machine goes through and places a dot where there could be a defect in the material or conductivity. If there are too many defects, a tester will often throw away the entire wafer. However, one wafer could cost \$10,000 or more so throwing away a wafer that could be perfectly fine because of a false defect can be very costly. When dealing with an automated event or airport security, it may be okay to have a higher false negative rate because it is a relatively cheap and non-life-threatening task to confirm an automated alert as actually positive.

These data points can be graphically represented using the receiver operating characteristic curve or ROC curve. The ROC curve is a graph with the x axis from 0 to 1 of the false positive rate, and the y axis from 0 to 1 of the true positive rate at various threshold settings. A perfect predictor would have a false positive rate of 0 and a true positive rate of 1. When graphed over a series of thresholds, the area under the curve (AUC) can provide a single value for providing insight into how well the model is classifying the labels. For interpretation, the higher the AUC, the better the model performs. The AUC is more descriptive than accuracy because it is a balance of accuracy and false positive rate.

Both recall and precision are often reported for classification performances. Recall is the ability to find all relevant instances while precision is the proportion of the data points the model considers relevant that are actually relevant. Precision is the number of true positives divided by the number of true positives and false positives. This provides an indication of the ability of a classification model to identify only relevant data. For example, if running a preliminary test to predict if a patient has a disease or not, precision would be equal to the number of patients who have the disease and were predicted correctly divided by the number of patients who have the disease and were predicted correctly plus the patients incorrectly predicted as having the disease. Recall is the number of true positives divided by the number of true positives and false negatives. Recall tells

us the ability of a model to find all relevant cases within a dataset. Using the example above, recall would be equal to the number of patients who have the disease and were predicted correctly divided by the number of patients who have the disease and were predicted correctly plus the patients incorrectly predicted as not having the disease. Thus, a high recall is desired to find all patients who actually have the disease and can allow a lower precision if the cost of a follow up check is low.

5 Ethics

The disclaimer for utilizing this tool when selecting which machine learning model to implement in a production setting should be done so with caution. Currently, the tool does not have certain functionalities to mimic real-world datasets, such as hyper-parameter tuning, handling extreme outliers or imputing missing values. Failure to incorporate these types of characteristics in the simulated dataset can lead to inaccurate conclusions as to which algorithm yielded a better performance. Therefore, any conclusions drawn from the RShiny tool may not generalize for dataset in a production environment and hence the authors of this paper are not responsible for such conclusions. Moreover, the tool only considers random forest and logistic regression. As a result, other algorithms such as support vector machines or neural networks could produce higher prediction accuracies and could be a better model to implement with datasets where the decision boundary is non-linearly separable.

Additionally, since all data will be simulated via the analytical tool developed in this work, there is no concerns of the availability, integrity, usability, and security of the data. For future work, functionality that will allow users to upload their own dataset to this tool and run an evaluation may be considered. In this event, the authors will not store or publish any of the users' data inside the application. Because data is not currently being stored, security scans will not be performed on the application, but best practices will be applied to avoid any known vulnerabilities. If any issues are reported, the authors will also notify users of potential deficiencies as well as correct these deficiencies if possible. Finally, in the context of the synthetic data generated in the application, there is no legal violations or security concerns. It should be clearly stated that the users should only consider the tool for educational purposes only as this application is still in the development phase. Any decisions drawn from the tool are not endorsed by the authors of this paper. The application should be used for education purposes only.

6 Analysis and Results

The analytical tool has four case studies that runs on simulated data based on the user's customized inputs. Figure 9 is an example of what the output of the results looks like. On each tab of the tool, a table of the evaluation metrics for each model as well as line graphs of those evaluation metrics. For accuracy, true

and false positive rates, a boxplot is displayed of the spread of all values for each simulation run. For example, if I set the number of simulations to 1000, the table and line chart will show me the average value of all 1000 simulations and the boxplots take into account the values for each of the 1000 simulations.

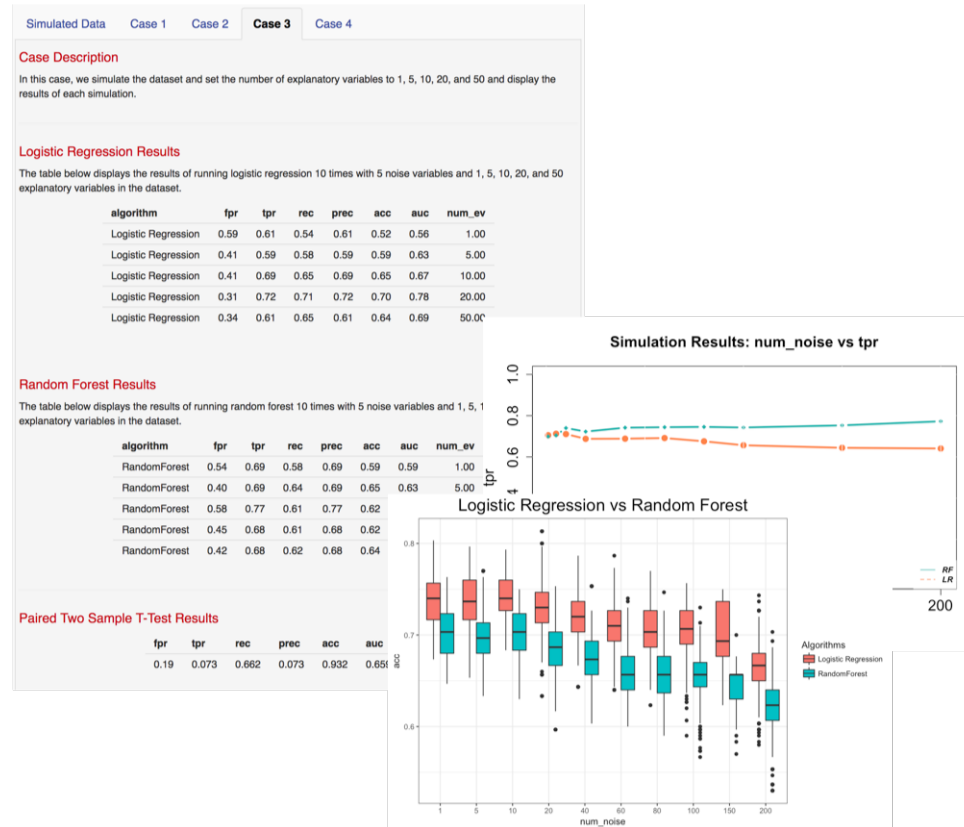


Fig. 9. Results of running each model of simulated data in each case study

6.1 Case 1

The first case investigated was comparing model performance with respect to change in variance in the explanatory and noise variables. The hypothesis was that an increase in variance would strengthen the accuracy for both models. For this simulation case, the application was configured to run 1000 simulations for 1000 observations. In the top row of Figure 10, the results display the accuracy for varying levels of variance in 10 noise and 5 explanatory variables. There is both visual evidence from Figure 10 and statistical evidence from the paired

t test (p-value ≤ 0.05) to suggest that, on average, the accuracy of the logistic regression model is greater than that of the random forest model.

The bottom row of Figure 10 displays the results of the true positive rate on the left and false positive rate on the right. The true positive rate for both models are nearly the same at each variance level. However, one can see that the false positive rate for random forest is not significantly higher than logistic regression (p-value = 0.63). Even though both models have the same performance in terms of correctly classifying a true value as true, the false positive rate for random forest is higher than logistic regression. This causes logistic regression to outperform random forest in terms of overall accuracy at each level of variance.

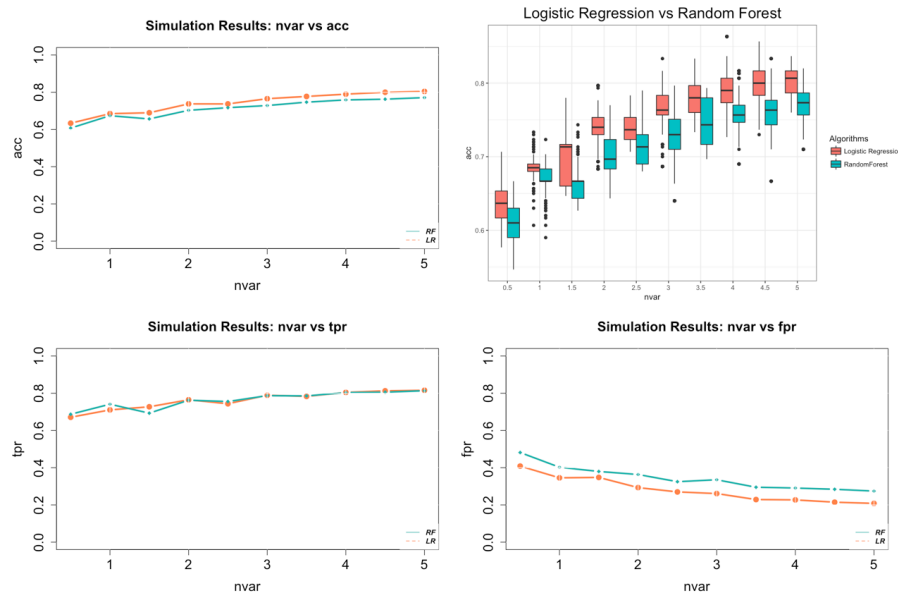


Fig. 10. Case 1 Simulation Results: 5 noise and 10 explanatory variables

The simulation is conducted again, but now adding more noise variables (noise = 100) and the results of the accuracy is shown in the top row of Figure 11. By looking at Figure 10 and Figure 11, a similar trend is observed. With a p-value less than 0.05, there is strong evidence to suggest that a significant difference in accuracy between the two models exists. The boxplots for this simulation is also comparable with 10 noise variables where minimal overlap in the boxplots for each model at each level of variance is observed. However, with 100 noise variables, the boxplots are much more consistent in that they are all about the same size for each level of variance. In Figure 11, the boxplots for each model are noticeably different sizes.

The bottom row of Figure 11 displays the true positive rate on the left and false positive rate on the right with 100 noise variables and 5 explanatory variables over increasing levels of variance in the variables. Interestingly, for variance 0.5 to 2.5 and a lot of noise, random forest has a higher true positive rate. At around variance = 3.0 and higher, random forest still has a higher true positive rate, but it is not as large of a difference from logistic regression than variance ≤ 2.5 . The false positive rate for random forest is again higher than logistic regression so the gap in higher true positive rate is not enough to make overall accuracy higher for random forest.

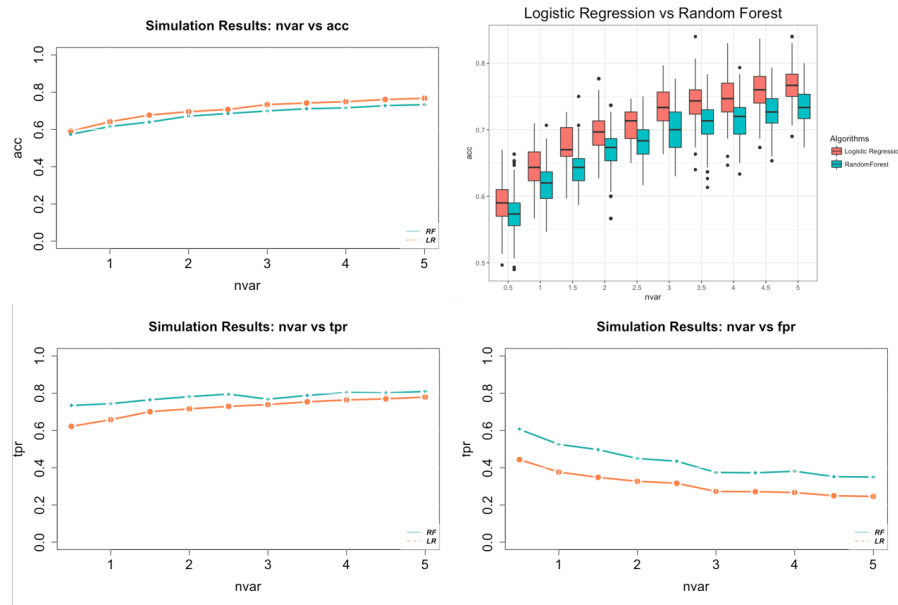


Fig. 11. Case 1 Simulation Results: 100 noise and 5 explanatory variables

6.2 Case 2

In case 2, we compared model performance with respect to change in the amount of noise in the dataset. We first did this by running 1000 simulations on a dataset with the number of noise variables = 1, 5, 10, 20, and 50. In Figure 12, we see the results of the accuracy for each model when the number of explanatory variables is 5 with 1000 observations. As we expected, as the amount of noise in the dataset increases, we see the accuracy start to decline for both models. However, we were not able to get the full picture by stopping at 50 noise variables, so we ran this again increasing the noise further. Figure 13 shows the results of accuracy when setting the number of noise variables to 1, 5, 10, 20, 40, 60, 80,

100, 150, 200. Accuracy is still slightly declining as we increase the noise past 50 noise variables and logistic regression is still performing with a higher accuracy (p-value = 9.559e-07).

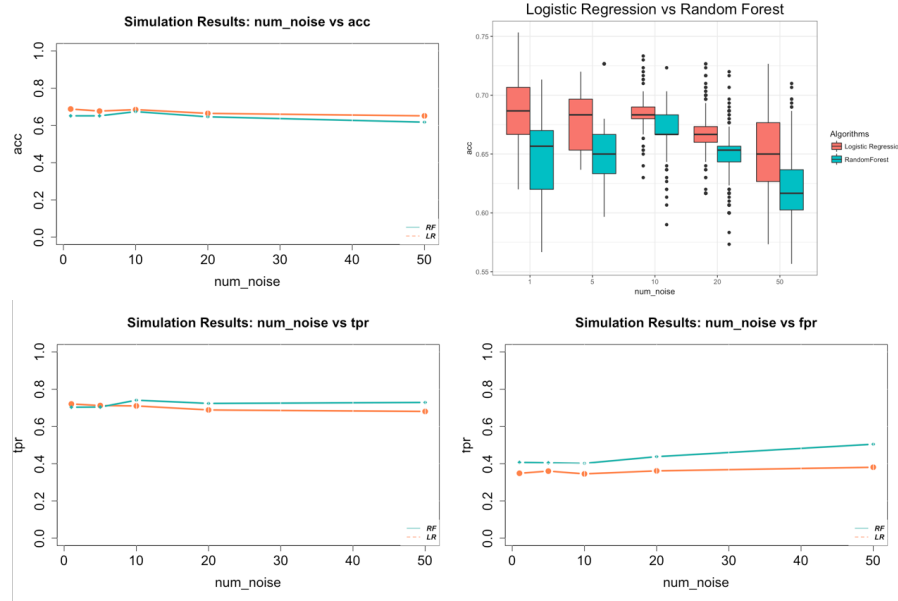


Fig. 12. Case 2 Simulation Results: 1 to 50 noise variables

The bottom rows of Figures 12 and 13 show the true positive rate on the left and false positive rate on the right. For true positive rate, when the number of noise variables is the less than or equal to the number of explanatory variables in the dataset, logistic regression is higher. However, once the number of noise variables exceeds the number of explanatory variables, random forest begins to have a higher true positive rate than logistic regression. As the amount of noise in the data increases, the false positive rate for both models also increase. However, the rate of increase in false positive rate for random forest is greater than the rate of increase in false positive rate for logistic regression as noise increases. Logistic regression does not have much change in true or false positive rate as noise increases past 50, but random forest false positive rate noticeably increases past 50 noise variables.

6.3 Case 3

In case 3, we compared model performance with respect to change in the number of explanatory variables in the dataset. In other words, the number of variables that relate to the response variable we are predicting. We did this by running

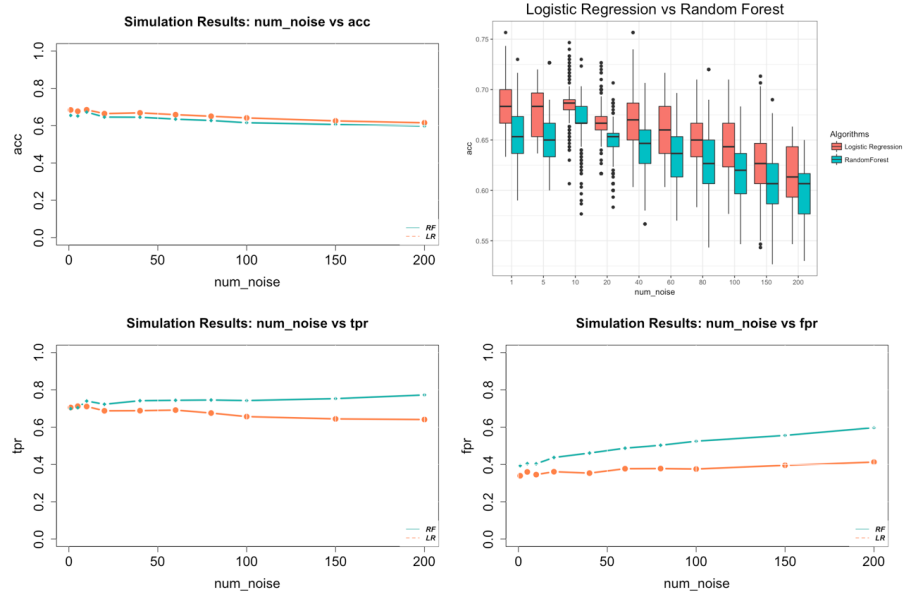


Fig. 13. Case 2 Simulation Results: 1 to 200 noise variables

1000 simulations on a dataset with the number of explanatory variables = 1, 5, 10, 20, 30, 40, 50. In the top row of Figure 14, we see the results of the accuracy for each model when the number of noise variables is 50 with 1000 observations. With 30 or less explanatory variables, as the number of explanatory variables in the dataset increases, the accuracy increases as well. When the number of explanatory variables is above 30, random forest begins to taper off whereas logistic regression continues to increase in overall accuracy.

The bottom row of Figure 14 displays the true positive rate on the left and false positive rate on the right. When we look at the true positive rate for the models, below 30 explanatory variables, random forest had a higher true positive rate. At 30 explanatory variables, logistic regression crosses over and continually increases to have a higher true positive rate than random forest. When we look at false positive rate, logistic regression decreases as we add more explanatory variables. Random forest false positive rate initially decreases from 1 to 10 explanatory variables but from 10 to 50 explanatory variables, there is not much change. The crossover point in true positive rate at 30 explanatory variables and the continued decrease in false positive rate for logistic regression as explanatory variables increases is evident in the overall accuracy plots. From the accuracy plots, we can see the drastic gap in performance of the two models after 20 to 30 explanatory variables.

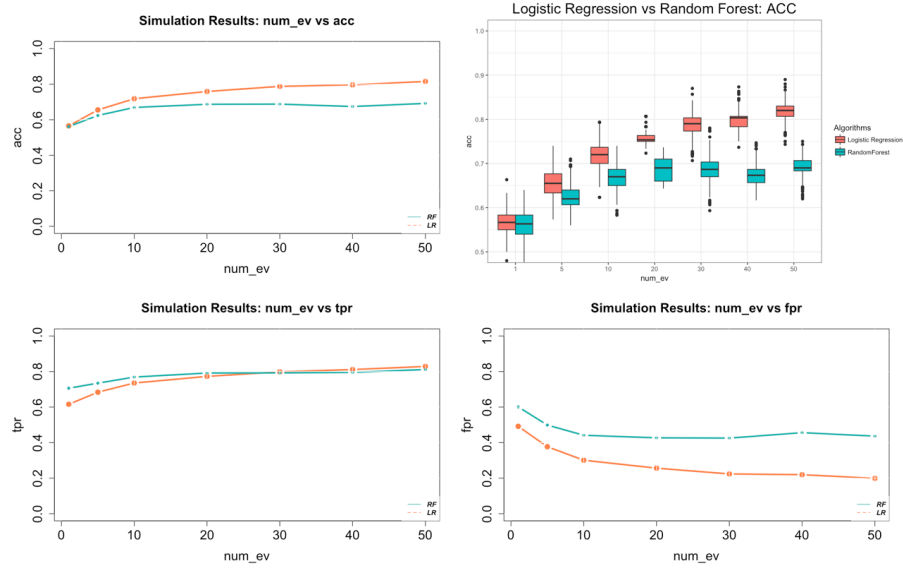


Fig. 14. Case 3 Simulation Results

6.4 Case 4

This simulation case study looked at iteratively increasing the number of observations in the dataset from 10 to 10000 while holding the number of explanatory variables in the model constant. A total of four different subcases were evaluated in which the number of explanatory variables ranged from 1,10,20, and 50; each case comprised of 10 noise variables. Given the computational complexities and completion time to train and validate a model for 1000 simulation as described in the previous cases and increasing the overall size of the dataset, the total number of simulations for specific case study is 10. Hence the moderate variance observed in Figure 15 and Figure 16.

One of the interesting findings in this simulation case is random forest and logistic regression perform nearly the same up until approximately 1000 observations in the dataset before diverging or in some instances crossing over as shown in Figure 16. Secondly, in Figure 15, the overall accuracy for logistic regression is consistently higher than random forest with 100 trees as both the number of explanatory and observations increases. With 10 and 20 explanatory variables included in the model respectively, the difference in overall accuracy is at a minimal compared to the case where 50 explanatory variables are included.

Additional analysis will need to be performed to determine if these observations are consistent when random forest is trained with a larger number of trees in addition to increasing the number of simulations to have a better understanding of the true accuracies.

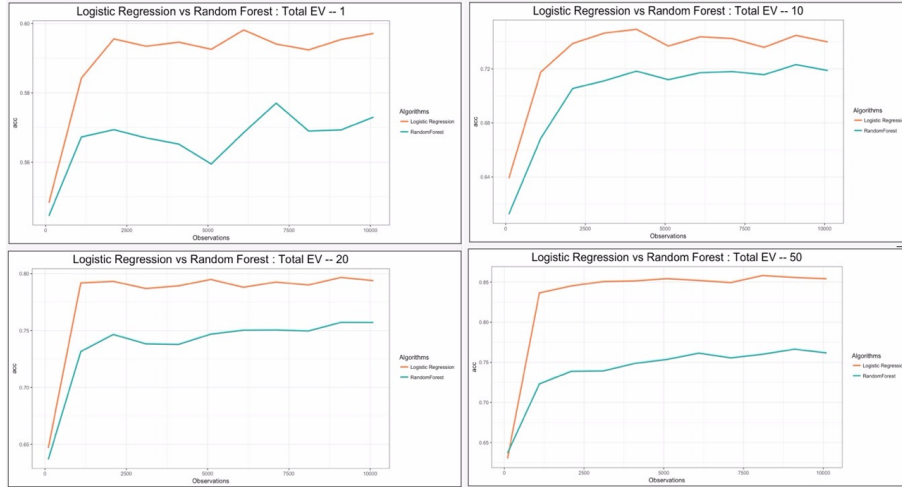


Fig. 15. Case 4 Simulation Results - Accuracy

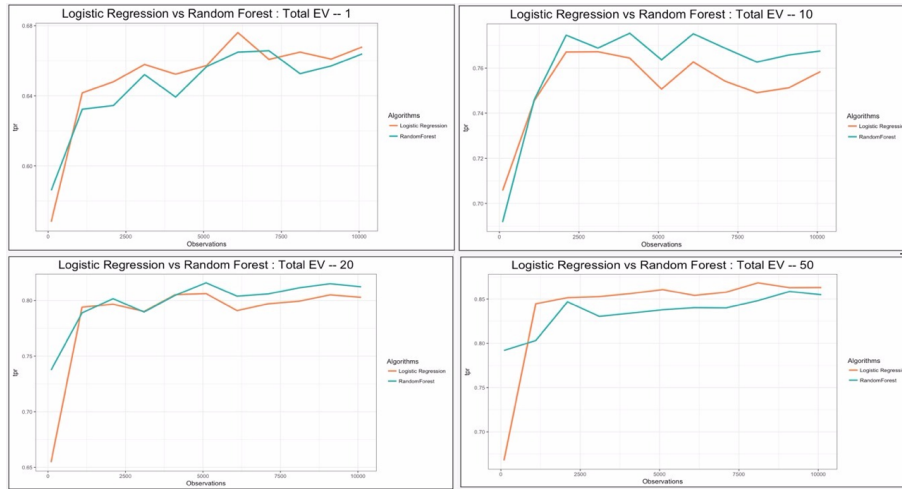


Fig. 16. Case 4 Simulation Results - True Postive Rate

7 Conclusions

We are waiting to add hard conclusions here. Unfortunately, our results are based off of running random forest set at 100 decision trees. In the next five weeks we will re-run simulations are come to conclusions.

In the meantime, we have the following general conclusions:

- Logistic regression performs better when the number of noise variables is less than or equal to the number of explanatory variables
- Random forest has a higher true and false positive rate as the number of explanatory variables increases in a dataset
- Logistic regression and random forest are comparable for smaller datasets as in datasets with less than 1000 observations

8 Future Work

In the current development version, the application provides the ability to answer in-depth statistical questions and evaluate classification performance of only two machine learning models. Future development is to incorporate other algorithms such as Naive Bayes, XGBoost, and Artificial Neural Networks. Also, the application can be expanded beyond binary classification to multi-labeled datasets and evolving to include regression.

Specifying the for the number of trees in the random forest model is an input the user will need to tune for in attempt to improve performance. Rather than hard coding the number of trees, the user could select an apply grid search option, which is an exhaustively optimization method that scans all possible parameter combinations in order to find the best estimators that yields the highest accuracy or other specified metrics.

Lastly, one of the motivations behind this work is having the ability to open the door for other questions of interest to help address, like what is the average Type I or Type II error for logistic regression using forward selection criteria under similar data structures outlined in the results section of this paper. Just one example, but this application is intended to provide a foundation and the ability to expand well beyond the scope of work presented in this analysis.

References

1. Beğenilmis, Erdem; Üsküdarlı, Suzan. Organized Behavior Classification of Tweet Sets using Supervised Learning Methods. eprint arXiv:1711.10720. 11/2017.
2. Bertrand Michel. A Statistical Approach to Topological Data Analysis. Statistics [math.ST]. UPMC Université Paris VI, 2015.
3. Frédéric Chazal; Bertrand Michel An introduction to Topological Data Analysis: fundamental and practical aspects for data scientists. 2017
4. Graham Dunn. (2007) Regression Models for Method Comparison Data. *Journal of Biopharmaceutical Statistics* 17:4, pages 739-756
5. Guerreiro, Manuela; Maroco, João; de Mendonça, Alexandre; Rodrigues, Ana; Santana, Isabel; Silva, Dina. Data mining methods in the prediction of Dementia: A real-data comparison of the accuracy, sensitivity and specificity of linear discriminant analysis, logistic regression, neural networks, support vector machines, classification trees and random forests. *BMC Research Notes* 20114:299. <https://doi.org/10.1186/1756-0500-4-299>.
6. Hastie, T., Tibshirani, R., Friedman, J. (2009). *The elements of statistical learning: data mining, inference and prediction*. Springer.
7. Olson, Randal S.; Moore, Jason H. Identifying and Harnessing the Building Blocks of Machine Learning Pipelines for Sensible Initialization of a Data Science Automation Tool. eprint arXiv:1607.08878. 07/2016.
8. Ng, Andrew. CS229 Lecture Notes. Stanford University. 2012
9. Munch, Elizabeth. A User's Guide to Topological Data Analysis. University at Albany. 2017
10. Ruiz-Gazen, Anne; Villa, Nathalie Storms Prediction: Logistic Regression vs Random Forest for Unbalanced Data.
11. Thanh Lam, Hoang; Thiebaut, Johann-Michael; Sinn, Mathieu; Chen, Bei; Mai, Tiep; Alkan, Oznur. One button machine for automating feature engineering in relational databases. eprint arXiv:1706.00327. 06/2017.
12. Zoran Bursac, C Heath Gauss, David Keith Williams, David W Hosmer: Purposeful Selection of Variables in Logistic Regression. *Source Code for Biology and Medicine*. 2008

Yes, we know we need more. We have 5 weeks and many papers to go through.