

# Assignment 2

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## 1. Introduction

The spam detection feature of e-mail systems are crucial. Three algorithms were used in this study to compare how well they classified emails as spam or not. To compare the algorithms, we will be conducting two statistical tests, namely Friedman and Nemenyi test.

The classifiers chosen were Support Vector Machine (SVM), AdaBoost and Random Forest. Support Vector Machine's (SVM) objective is to find a hyperplane in an N-dimensional space that distinctly classifies the data points [1]. AdaBoost, short for Adaptive Boosting, is a statistical classification meta-algorithm [2] and random forest is a tree based classifier. [3].

Our data set consist of aggregated statistical summaries of e-mails in a big cohort.

## 2. Method

### 2.1. Data exploration

The predictors follow a right-skewed distribution. The target column has a 60/40-percent ratio of ham and spam making it a fairly balanced data set. Furthermore, we identified 391 duplicate instances.

### 2.2. Data transformation

Due to support vector machines being very sensitive to outliers [4] we discretized our data set so that we'd get more comparable performance results between the algorithms. Discretization was performed with the KBinsDiscretizer method from Sklearn [5] with 7 bins and kmeans binning technique. Note however that the discretized data set was only used for training the SVM classifier. All other classifiers were trained on non-transformed data.

### 2.3. Computing the Friedman and Nemenyi test

The Friedman test is a non-parametric statistical test used to determine whether there are significant differences between treatments being compared in a study. The Friedman test is a rank-based test, meaning it ranks the treatments being compared and then computes a test statistic

based on the ranks. If the test statistic passes a critical value threshold (determined by the number of treatments being compared and the significance level), then the null hypothesis  $H_o$ , that there are no significant differences between the treatments is rejected.

#### Friedman test [6]

$$(1.1) \text{ Mean rank : } \bar{R} = \frac{k+1}{2};$$

$$(1.2) \text{ Sum of squared differences : } n \sum_j (R_j - \bar{R})^2;$$

$$(1.3) \text{ Sum of squared differences : } \frac{1}{n(k-1)} \sum_{ij} (R_{ij} - \bar{R})^2;$$

$$(1.4) \text{ Friedman score - ratio between 1.2 and 1.3.}$$

The Nemenyi test is a post-hoc statistical test that is conducted after completion of a study in order to compare treatments in a group. The Nemenyi test allows us to identify which treatments are significantly different from each other. We conclude whether the difference between treatments is significant or not by seeing if the difference between means is greater than the critical difference.

#### Nemenyi test

$$(2.1) \text{ Critical difference : } CD = q_\alpha \sqrt{\frac{k(k+1)}{6n}}; [6]$$

$$(2.2) |\bar{R}_i - \bar{R}_j| > CD. [7]$$

## 2.4. Algorithms

This study has chosen the following 3 classification models to compare:

**Support vector machine:** SVMs conceptualizes each object as a point in N-dimensional space with its features being the coordinates. SVMs perform the classification task by drawing a hyperplane in such a way all instances belonging to the positive class are separated from its negative counterpart and that the distance between each category to the plane is maximized. If the data points can't be separated by a hyperplane a kernel-trick is applied so that the hyperplane can be fitted.

**AdaBoost:** a meta-algorithm that wraps over a set of weaker classifiers. The classifiers are trained on samples

of the training data. For each iteration the trained model will be tested on the other samples of the training split and for those data points, if any, that have a high error associated with them will be weighted more in the next training iteration so that the model can train on its weaker areas.

**Random forest:** an ensemble technique which uses a set of decision trees and intelligent pruning algorithms for removing splits with low information-yield and a voting system so that the all predictions from the entire set are aggregated into a final prediction.

These algorithms show great performance, are very popular [8] and don't require much-to-any hyper parameter tuning. These reasons were the motivation behind our decision.

## 2.5. Training and evaluation

We used stratified k-fold cross validation as our training and testing procedure. The paradigm is as following: we split our data into k-folds and use k-1 of these folds to train our data and the remaining one to test. We repeat this procedure k-1 more times without replacement. To ensure we get a fair representation of our target with each fold the folds preserve the percentage of samples for each class [6]. With this technique we get a more robust measure of performance by allowing our model to train and test on different parts of the data set. However this methodology is more computationally expensive [9].

We measure model performance with the following three metrics:

**F-measure:** also known as F1 score, is a performance metric used to evaluate a classification model. It is computed as the harmonic mean between precision and recall. F-measure ranges between 0 to 1, with higher values meaning better performance. It is a balanced metric that considers both precision and recall of the model.

$$F1 = 2 * \frac{precision * recall}{precision + recall}, F1 \in [0, 1]$$

**Precision:** is the fraction of true positive (TP) predictions made by the model among all positive predictions.

$$precision = \frac{TP}{TP + FP}, precision \in [0, 1]$$

**Recall:** is the fraction of true positive predictions made by the classifier among all actual positive predictions.

$$recall = \frac{TP}{TP + FN}, recall \in [0, 1]$$

**Accuracy:** the percentage of predictions that are correct.

$$accuracy = \frac{TP + TN}{TP + FP + TN + FN}, accuracy \in [0, 1]$$

**Computational time:** time taken to train algorithm.

## 3. Results

The algorithms performance, group statistic and rank for each fold is presented in the below tables. Note that these are rounded values. For complete values check our python notebook.

	SVM	AdaBoost	Random forest
1	0.9349 (3)	0.9349 (2)	0.9588 (1)
2	0.9326 (3)	0.9413 (2)	0.9565 (1)
3	0.9283 (3)	0.9370 (2)	0.9522 (1)
4	0.9609 (2)	0.9630 (1)	0.9522 (3)
5	0.9304 (3)	0.9348 (2)	0.9587 (1)
6	0.9435 (3)	0.9457 (2)	0.9500 (1)
7	0.9457 (2)	0.9391 (3)	0.9609 (1)
8	0.9457 (3)	0.94783 (2)	0.9652 (1)
9	0.9391 (2)	0.9261 (3)	0.9457 (1)
10	0.9217 (3)	0.9304 (2)	0.9522 (1)
Average	0.9383	0.9400	0.9552
Std	0.0112	0.0104	0.0058
Average rank	2.7	2.2	1.2

TABLE 1. ACCURACY

	SVM	AdaBoost	Random forest
1	0.9153 (3)	0.9180 (2)	0.9359 (1)
2	0.9132 (3)	0.9264 (2)	0.9448 (1)
3	0.9081 (3)	0.9192 (2)	0.9392 (1)
4	0.9508 (2)	0.9544 (1)	0.9479 (3)
5	0.9070 (3)	0.9148 (2)	0.9337 (1)
6	0.926 (3)	0.9319 (2)	0.9359 (1)
7	0.9315 (2)	0.9218 (3)	0.9392 (1)
8	0.9292 (3)	0.9322 (2)	0.9524 (1)
9	0.9222 (2)	0.9056 (3)	0.9438 (1)
10	0.8960 (3)	0.9080 (2)	0.9307 (1)
Average	0.9199	0.9232	0.9404
Std	0.0155	0.0141	0.007
Average rank	2.7	2.1	1.2

TABLE 2. F1-SCORE

	SVM	AdaBoost	Random forest
1	0.6522 (2)	0.4799 (1)	0.8543 (3)
2	0.4660 (1)	0.4678 (2)	1.0299 (3)
3	0.4923 (2)	0.4803 (1)	0.9217 (3)
4	0.4581 (1)	0.4777 (2)	0.9284 (3)
5	0.4696 (2)	0.4502 (1)	0.9095 (3)
6	0.4982 (1)	0.4997 (2)	0.8108 (3)
7	0.4859 (2)	0.4617 (1)	0.8169 (3)
8	0.4772 (1)	0.4879 (2)	0.8161 (3)
9	0.5296 (2)	0.4937 (1)	0.8134 (3)
10	0.4527 (1)	0.5061 (2)	0.8488 (3)
Average	0.4982	0.4805	0.8750
Std	0.0586	0.01730	0.0716
Average rank	1.5	1.5	3.0

TABLE 3. TRAINING TIME

The computed Friedman statistic for each table is approximately: 15.4 for accuracy, 11.4 for F1-score and 20.0 for time, which compared to the critical value, being 7.8 for  $k = 3$  and  $n = 10$  at the  $\alpha = 0.05$  [6], the Friedman statistic of all experiments is larger, resulting in all of them rejecting the null-hypotheses. To figure out which algorithms differed significantly from each other for each respective table, we conducted a Nemenyi test. We calculated the critical distance with formula (2.1): resulting in  $\approx 1.149$ . Next we computed equation (2.2) for each algorithm and table. In the accuracy experiment we find SVM and Random forest displaying a significant difference. Further, we looked into the F1-table, which gave us the same result as in accuracy table. Finally, we took a look at the training time table, where AdaBoost and Random Forest showed a difference greater than the critical difference.

## 4. Conclusion

This study showed the process of applying statistical tests on classification problems. There lies great value in being able to deduce the highest performing treatment, however with each benefit comes a cost. In decision making the benefit-to-cost ratio has to be considered. If one can afford having a more computationally expensive algorithm then, according to our tests, the random forest classifier serves this scenario best. If computational resources are scarce than an algorithm such as AdaBoost might lend you the best results with respect to cost.

## References

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