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In [1]: import numpy as np
from sklearn.svm import SVC
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
import matplotlib.pyplot as plt
from sklearn.metrics import roc_curve
from sklearn.metrics import auc
from sklearn.ensemble import RandomForestClassifier
from sklearn.neighbors import KNeighborsClassifier
from sklearn.naive_bayes import GaussianNB, MultinomialNB, BernoulliNB
from sklearn.neural_network import MLPClassifier
from sklearn.metrics import confusion_matrix
```

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In [2]: features = np.genfromtxt("./Aggregated_Data.csv", delimiter=",", usecols=(1
targets = np.genfromtxt("./Aggregated_Data.csv", delimiter=",", usecols=8)

trainingFeatures, testingFeatures, trainingTargets, testingTargets = train_
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In [10]: #follows the Confusion Matrix information page on Scikit Learn to visualize
#https://scikit-learn.org/stable/auto_examples/model_selection/plot_confusi
def plotMatrix(confusionMatrix, title, color):
    classLabels = ["No Outbreak", "Outbreak"]
    plt.rcParams['figure.figsize'] = (5.0, 5.0)
    plt.imshow(confusionMatrix, cmap=plt.cm.Blues)
    plt.xticks(ticks=[0,1], labels=classLabels)
    plt.yticks(ticks=[0,1], labels=classLabels)
    plt.ylabel("Predicted Labels")
    plt.xlabel("True Labels")
    plt.title(title)
    maximum = confusionMatrix.max()/2
    plt.text(0, 0, str(confusionMatrix[0][0]), horizontalalignment="center"
    plt.text(0, 1, str(confusionMatrix[0][1]), horizontalalignment="center"
    plt.text(1, 0, str(confusionMatrix[1][0]), horizontalalignment="center"
    plt.text(1, 1, str(confusionMatrix[1][1]), horizontalalignment="center"
    plt.show()
```

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In [9]: def thresholdBasedClassification(threshold, classifier, label, testingFeatu
predictions = []
probabilities = classifier.predict_proba(testingFeatures)
for j in range(0, probabilities.shape[0]):
    if (probabilities[j][1] <= threshold):
        predictions.append(0)
    else:
        predictions.append(1)
##https://scikit-learn.org/stable/modules/generated/sklearn.metrics.confu
#uses the confusion_matrix documentation on Scikit Learn to calculate t
confusionMatrix = confusion_matrix(testingTargets, predictions)
#uses the Confusion Matrix information page on Scikit Learn to find out
#https://scikit-learn.org/stable/auto_examples/model_selection/plot_con
normalizedConfusionMatrix = confusionMatrix.astype('float') / confusion
plotMatrix(confusionMatrix, label + (" Confusion Matrix (Threshold: %.4
plotMatrix(normalizedConfusionMatrix, label + (" Normalized Confusion M
```

In [5]:

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#The citations for the classifiers can be found in their respective files
decisionTree = DecisionTreeClassifier(criterion = "entropy", max_depth = 4)
decisionTree.fit(trainingFeatures, trainingTargets)

randomForest = RandomForestClassifier(n_estimators=100, max_depth=4, random
randomForest.fit(trainingFeatures, trainingTargets)

knn = KNeighborsClassifier(n_neighbors=6)
knn.fit(trainingFeatures, trainingTargets)

gaussianNaiveBayes = GaussianNB()
gaussianNaiveBayes.fit(trainingFeatures, trainingTargets)
multinomialNaiveBayes = MultinomialNB()
multinomialNaiveBayes.fit(trainingFeatures, trainingTargets)
bernoulliNaiveBayes = BernoulliNB(binarize=0.1)
bernoulliNaiveBayes.fit(trainingFeatures, trainingTargets)

network = MLPClassifier(solver='lbfgs', alpha=1e-5, hidden_layer_sizes=(3,)
network.fit(trainingFeatures, trainingTargets)

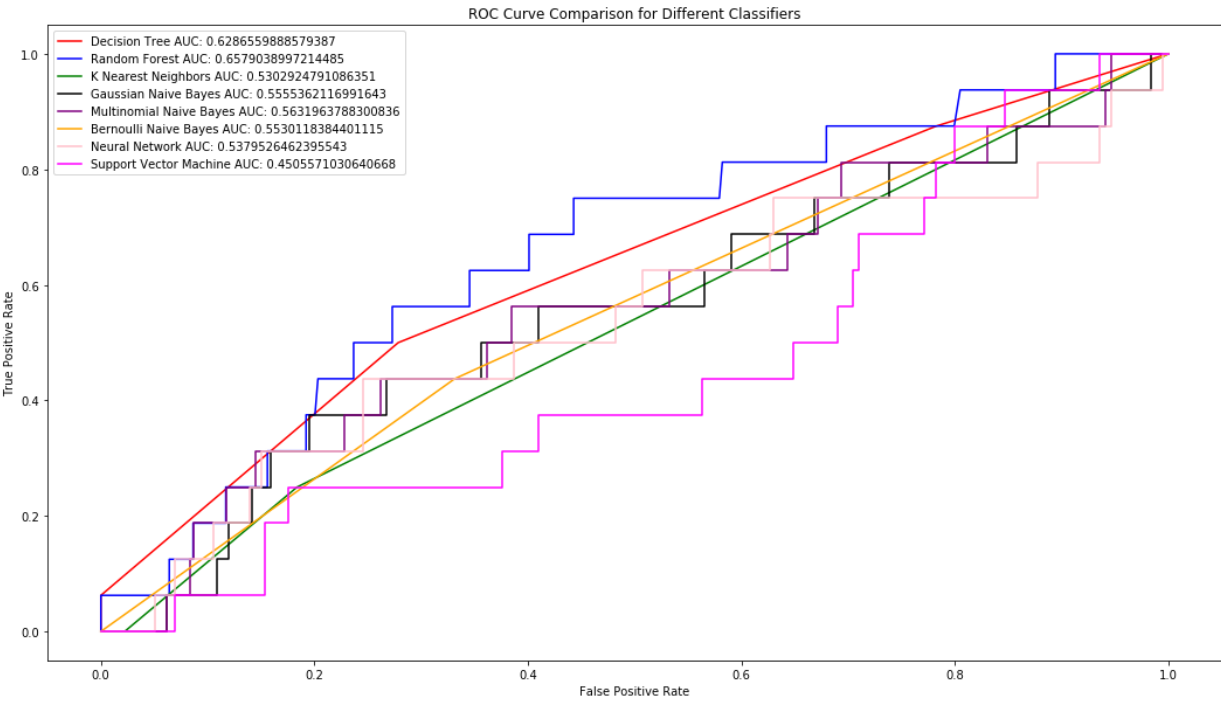
svm = SVC(kernel='poly', degree=3, random_state=0, gamma='auto', probabilit
svm.fit(trainingFeatures, trainingTargets)

classifiers = [decisionTree, randomForest, knn, gaussianNaiveBayes, multino
colors = ['red', 'blue', 'green', 'black', 'purple', 'orange', 'pink', 'mag
labels = ["Decision Tree", "Random Forest", 'K Nearest Neighbors', "Gaussia

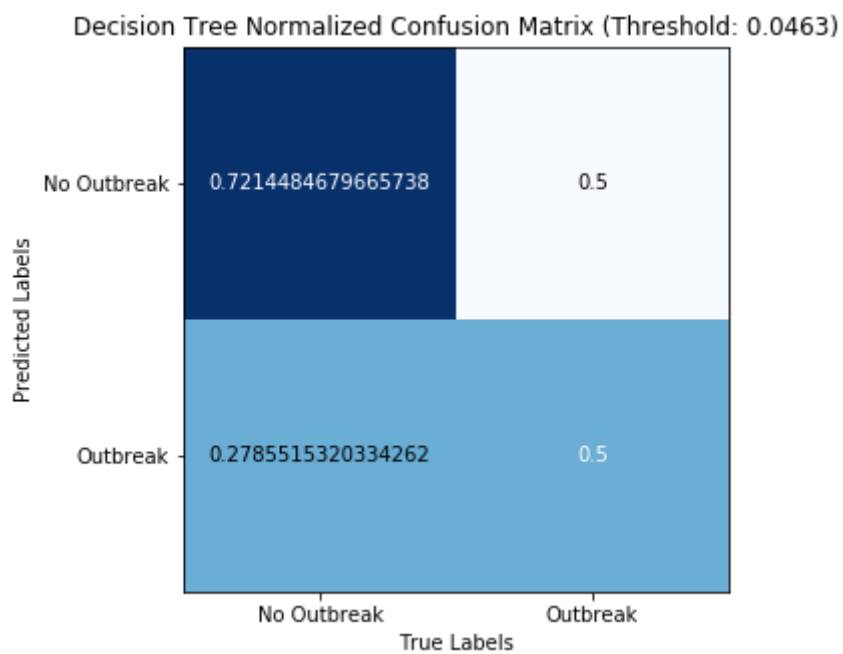
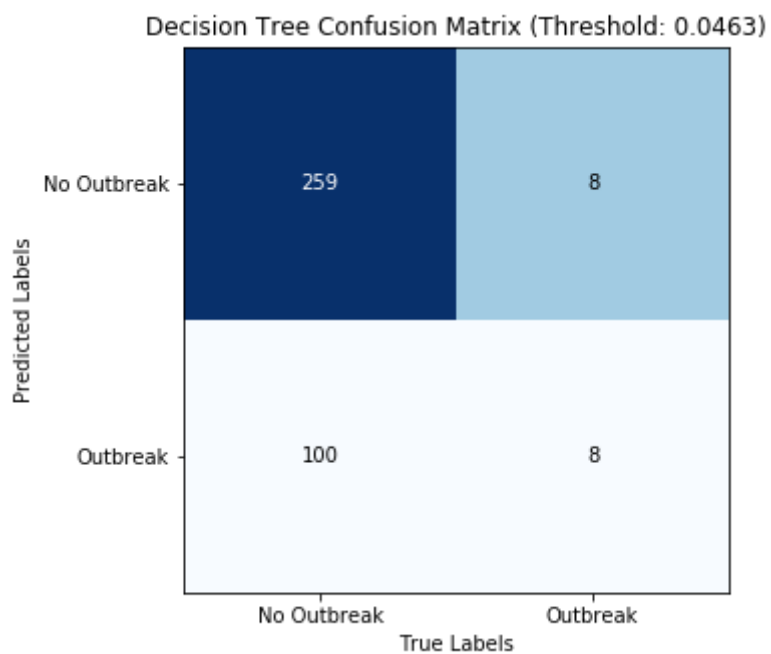
thresholds = []

plt.rcParams['figure.figsize'] = (18.0, 10.0)
for i in range(0,8):
    outbreakProbabilities = classifiers[i].predict_proba(testingFeatures)[:
    #uses the documentation on Scikit Learn for roc_curve to calculate the
    #https://scikit-learn.org/stable/modules/generated/sklearn.metrics.roc_
    fpr, tpr, tresh = roc_curve(testingTargets, outbreakProbabilities)
    thresholds.append(np.median(outbreakProbabilities))
    #uses the documentation on Scikit Learn for auc on how to calculate auc
    #https://scikit-learn.org/stable/modules/generated/sklearn.metrics.auc.
    area = auc(fpr, tpr)
    plt.plot(fpr, tpr, color=colors[i], label=labels[i]+" AUC: "+str(area))
    plt.legend()
    plt.xlabel("False Positive Rate")
    plt.ylabel("True Positive Rate")
    plt.title("ROC Curve Comparison for Different Classifiers")
plt.show()

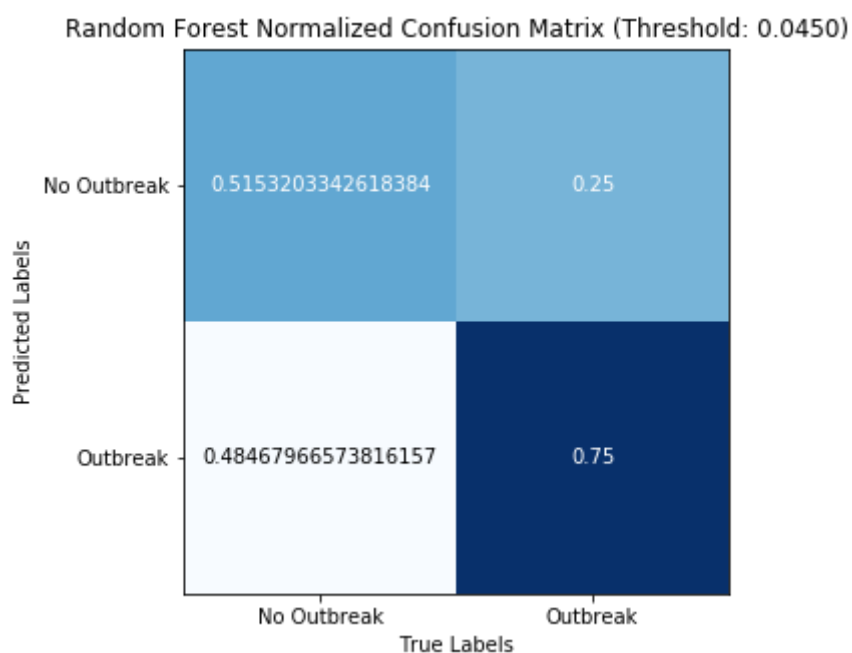
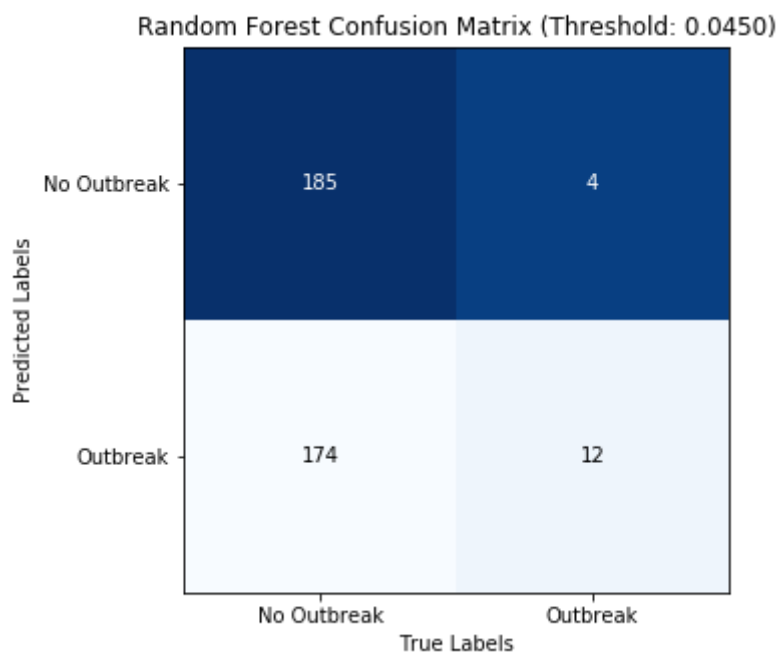
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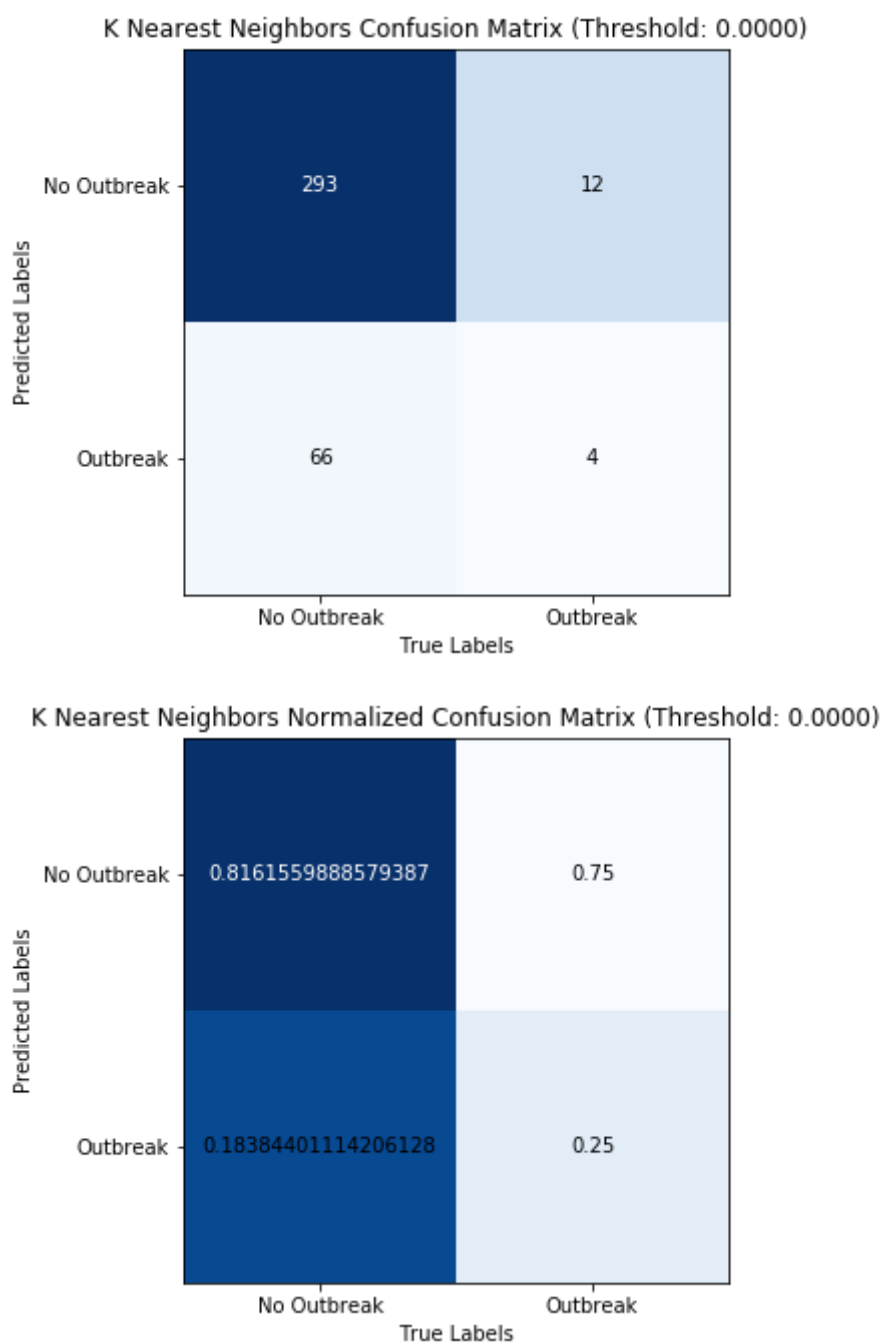
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In [11]: thresholdBasedClassification(thresholds[0], classifiers[0], labels[0], testi
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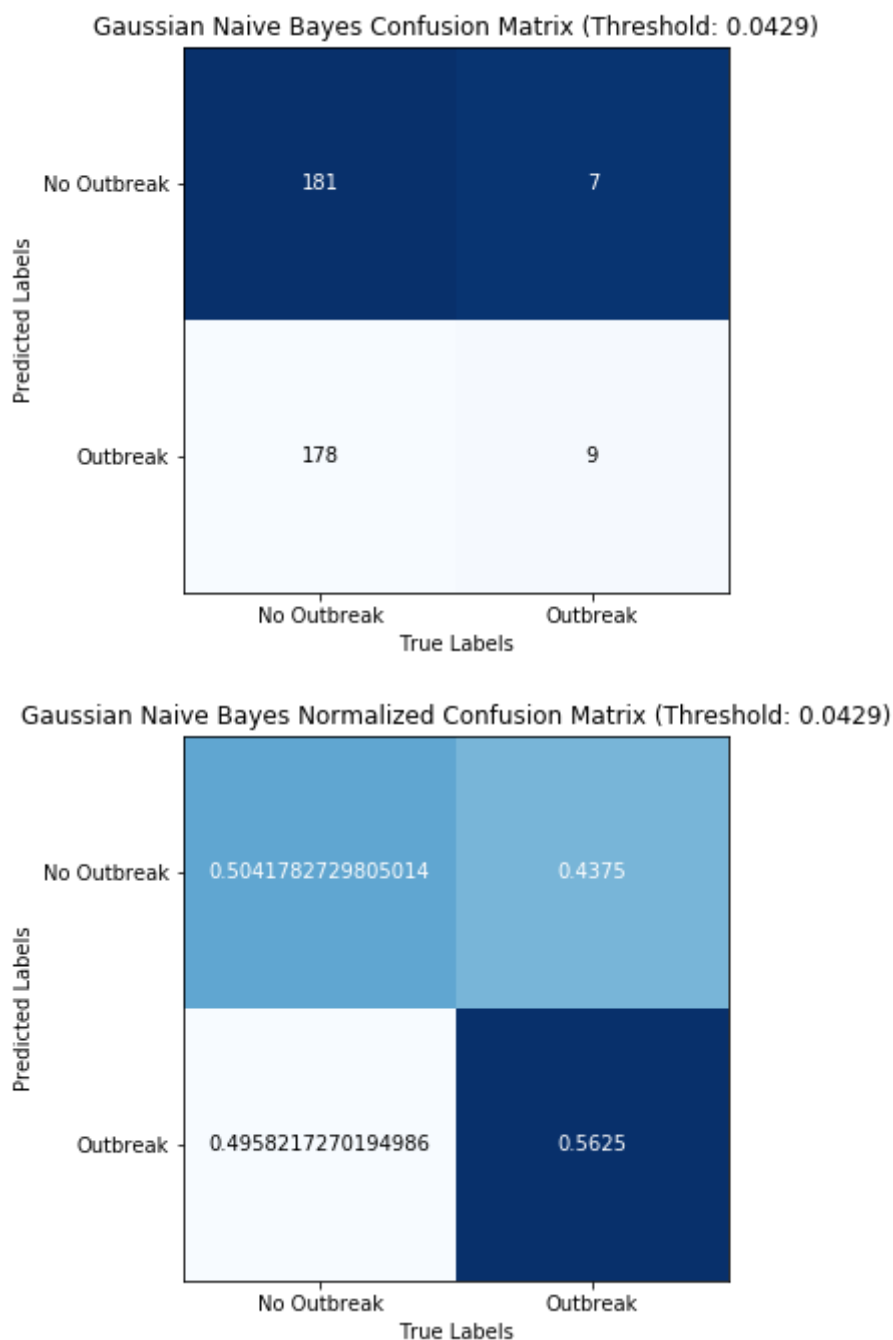
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In [12]: thresholdBasedClassification(thresholds[1], classifiers[1], labels[1], testi
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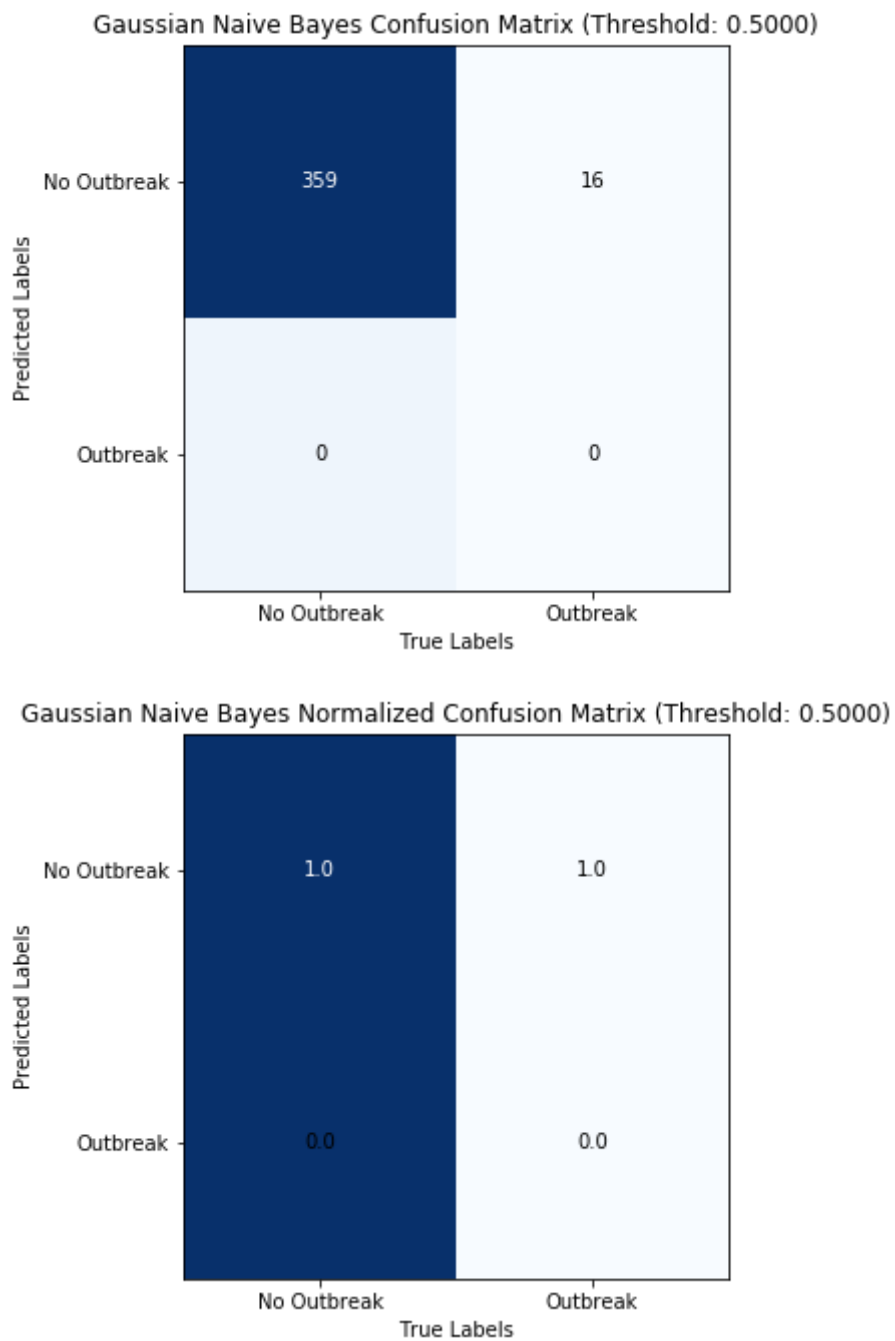
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In [13]: thresholdBasedClassification(thresholds[2], classifiers[2], labels[2], testi
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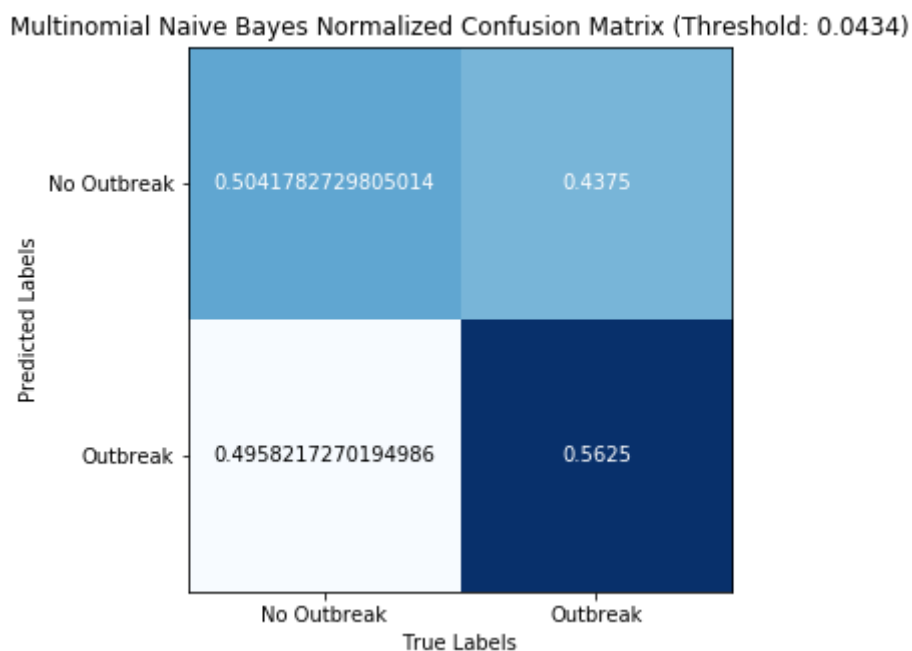
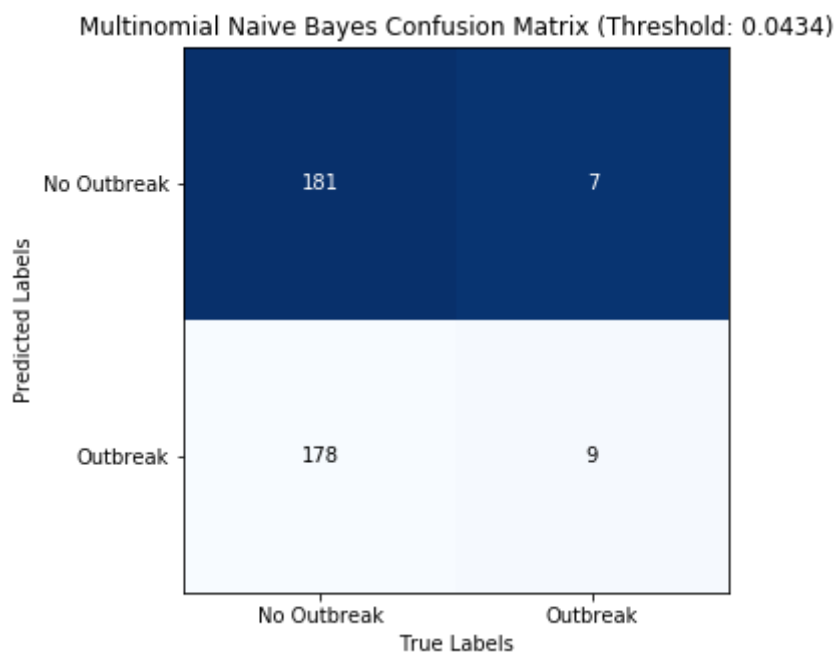
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In [14]: thresholdBasedClassification(thresholds[3], classifiers[3], labels[3], testi
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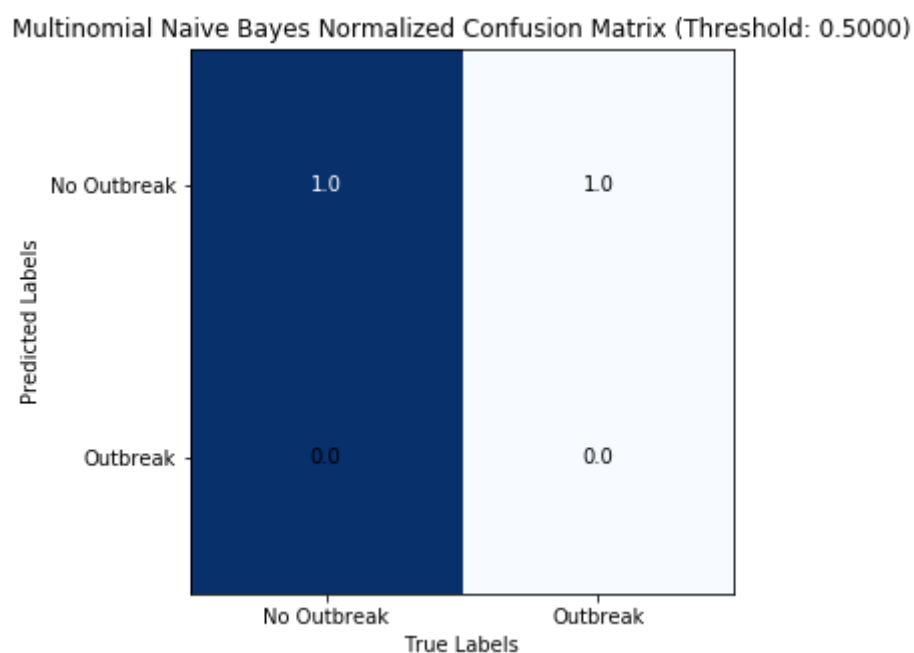
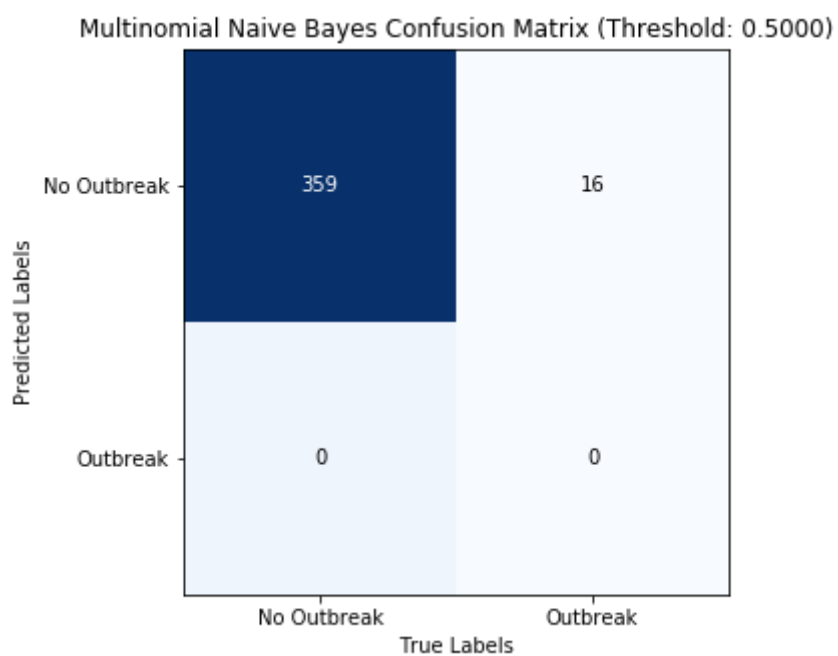
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In [22]: thresholdBasedClassification(0.5, classifiers[3], labels[3], testingFeatures
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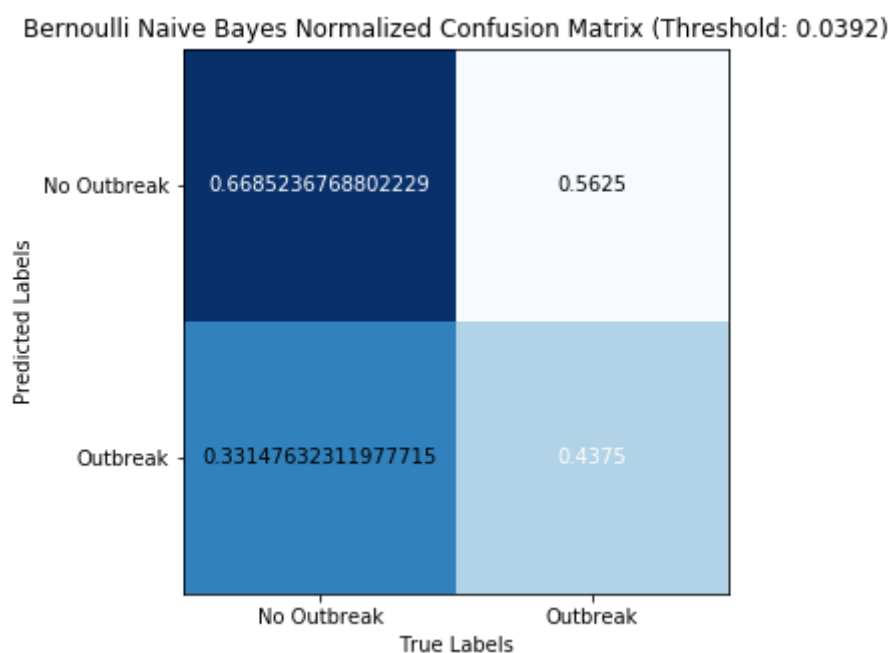
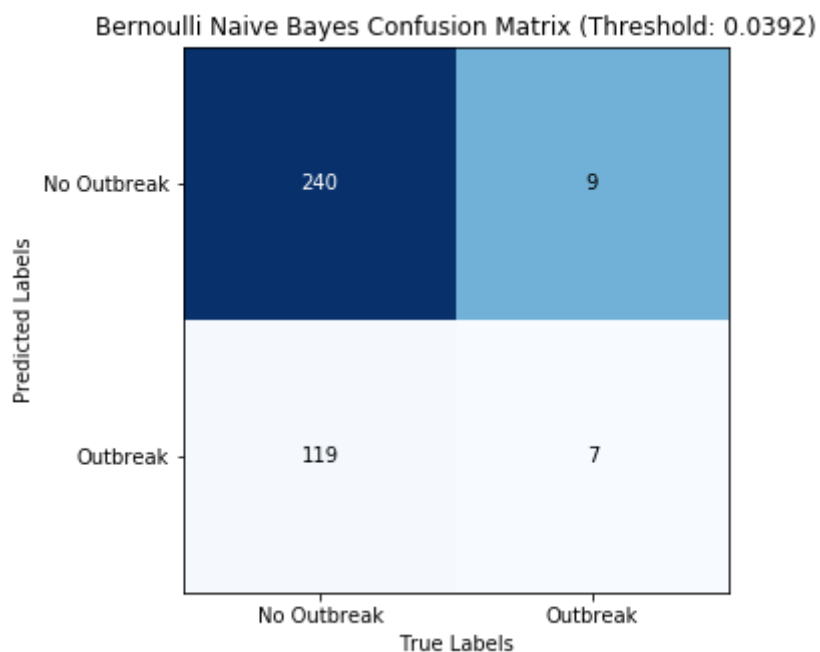

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In [15]: thresholdBasedClassification(thresholds[4], classifiers[4], labels[4], testi
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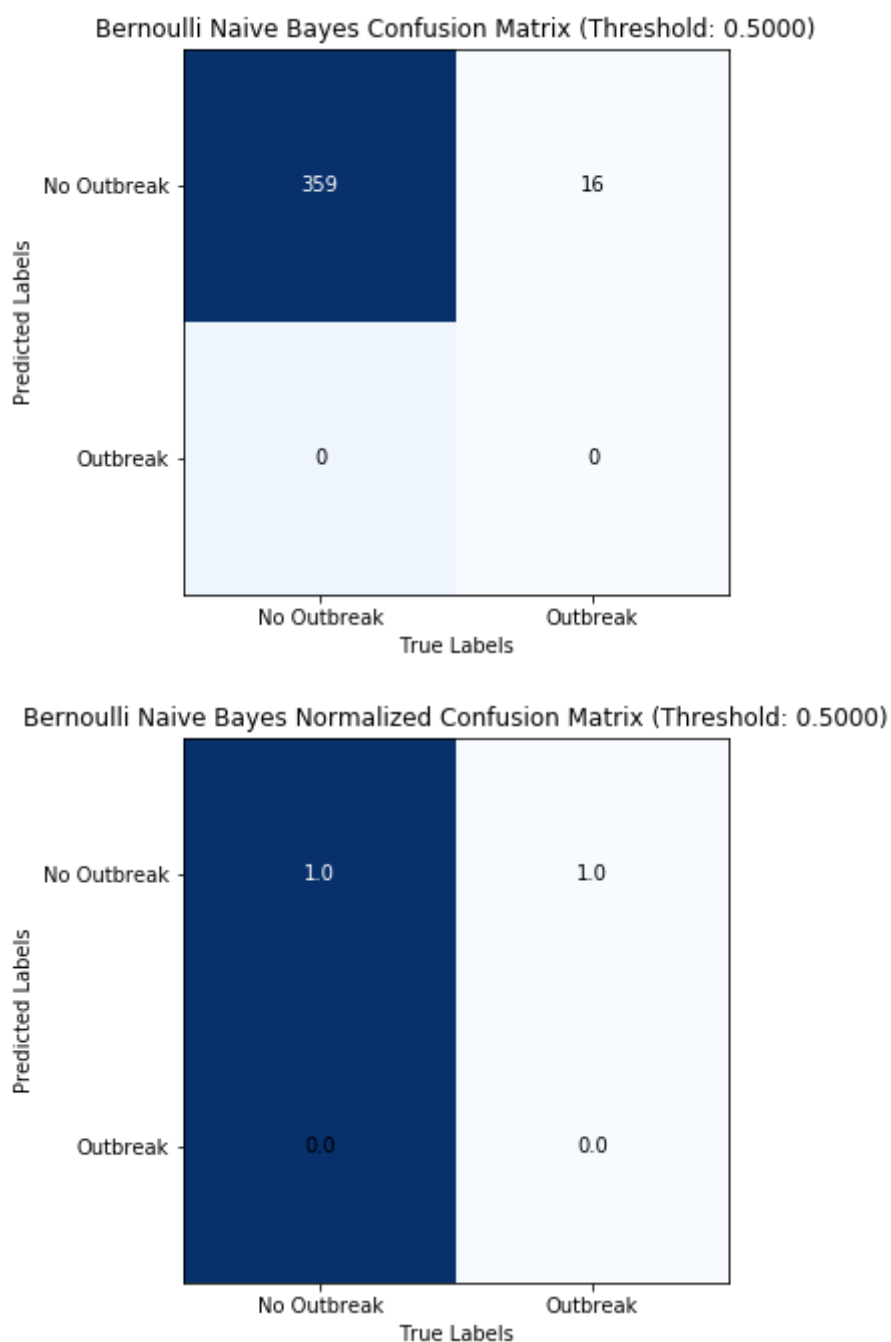
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In [23]: thresholdBasedClassification(0.5, classifiers[4], labels[4], testingFeatures
```



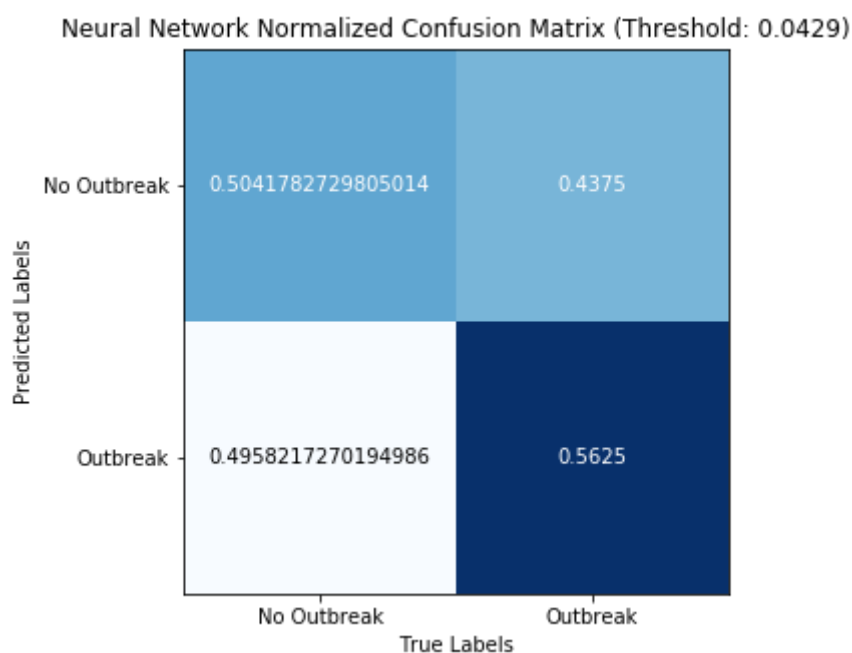
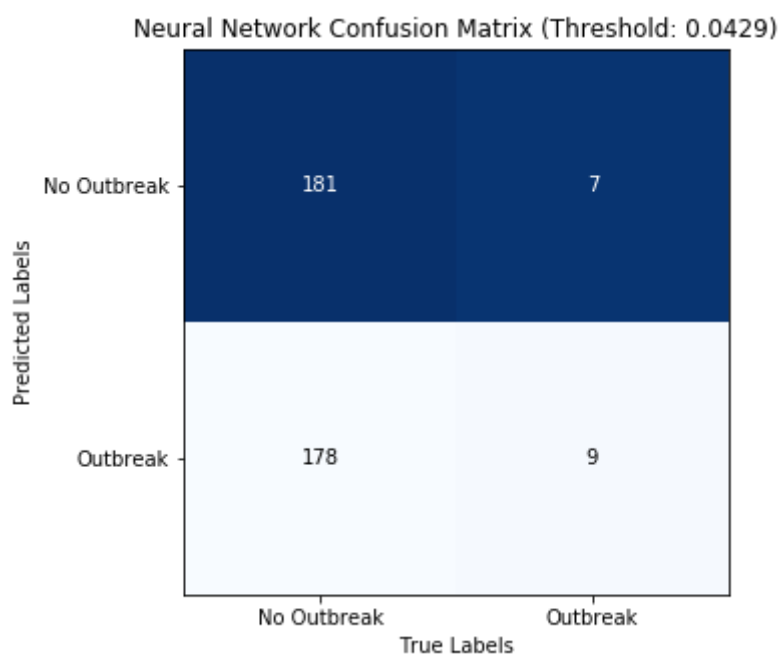
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In [16]: thresholdBasedClassification(thresholds[5], classifiers[5], labels[5], test_i
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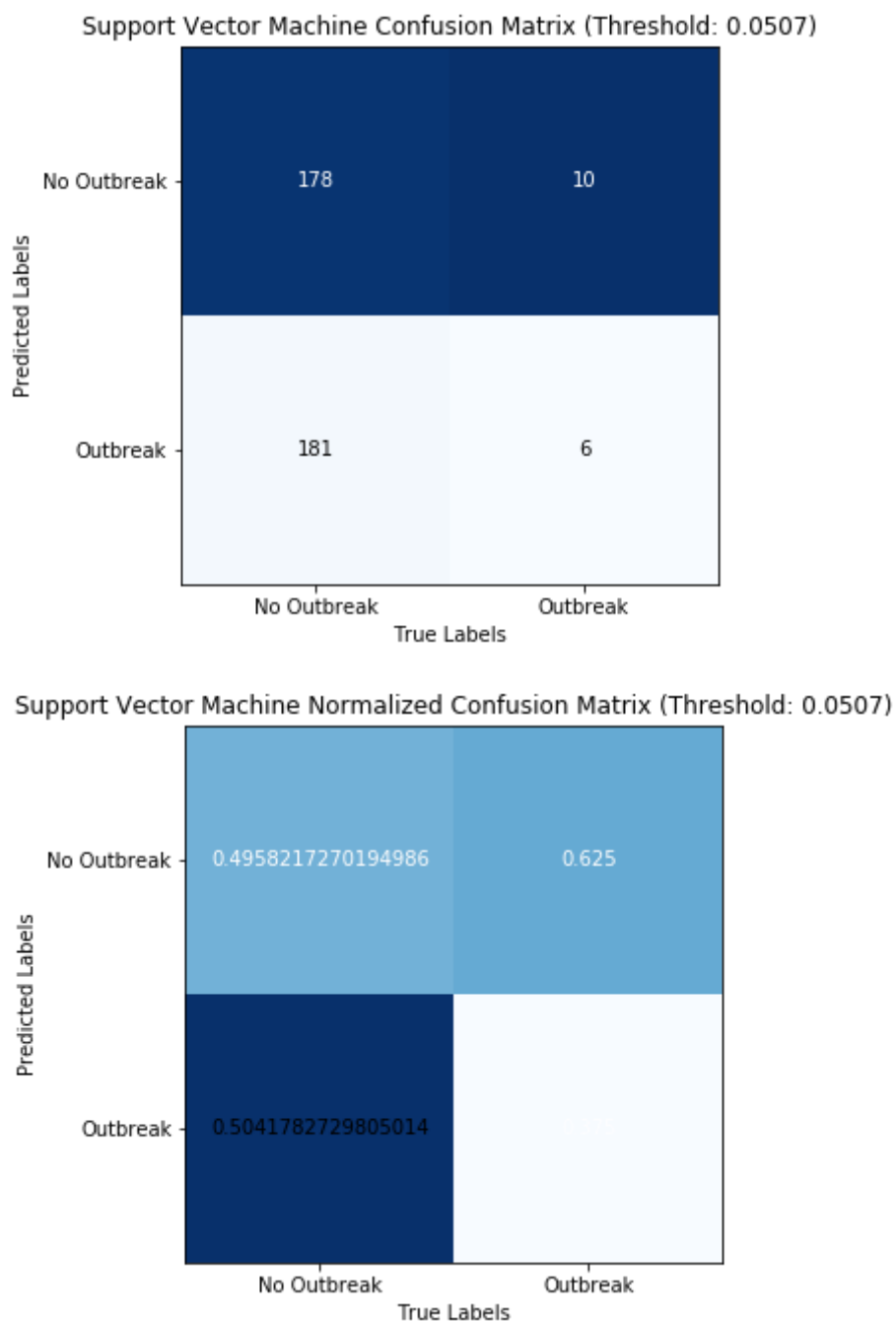
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In [19]: thresholdBasedClassification(0.5, classifiers[5], labels[5], testingFeatures
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In [17]: thresholdBasedClassification(thresholds[6], classifiers[6], labels[6], testi
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In [18]: thresholdBasedClassification(thresholds[7], classifiers[7], labels[7], testi
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In [ ]:
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