Assignment 4

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Datasets Used

- Iris Dataset
- Wine Dataset
- Seeds Dataset
- Breast Cancer Wisconsin Dataset
- Haberman's Survival Dataset
- Pima Indian Diabetes Dataset
- BankNote Authentication Dataset

All the datasets are taken from the well-known UCI Machine Learning repository.

Task 1 - Cluster Analysis

DBSCAN and Infomap don't work too well with the Seeds Dataset and thus we use the Wine Dataset in its place for those 2 algorithms. The Iris Dataset and the Breast Cancer Wisconsin Dataset has been used for all clustering algorithms.

K-Means Algorithm

Iris Dataset

```
Cluster 1 -> Length = 50
Cluster 2 -> Length = 53
```

Cluster 3 -> Length = 47

Cluster validation Indices (rounded to 4 decimal places):

Silhouette Index = 0.5059

Davies Bouldin Index = 0.8354

Dunne Index = 0.5303

Breast Cancer Wisconsin Dataset

```
Cluster 1 -> Length = 230
Cluster 2 -> Length = 453
```

Cluster validation Indices (rounded to 4 decimal places):

Silhouette Index = 0.5939

Davies Bouldin Index = 0.8200

Dunne Index = 0.6417

Seeds Dataset

```
Cluster 1 -> Length = 67
```

Cluster 2 -> Length = 71

Cluster 3 -> Length = 72

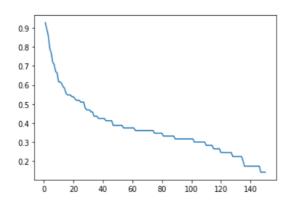
Cluster validation Indices (rounded to 4 decimal places):

Silhouette Index = 0.4506

Davies Bouldin Index = 0.9279 Dunne Index = 0.5000

DBSCAN Algorithm

Graph for determining optimum epsilon value using grid search.



Iris Dataset

Cluster 1 -> Length = 5

Cluster 2 -> Length = 50

Cluster 3 -> Length = 95

Cluster validation Indices (rounded to 4 decimal places):

Silhouette Index = 0.5336

Davies Bouldin Index = 0.5262

Dunne Index = 0.5813

• Breast Cancer Wisconsin Dataset

Cluster 1 -> Length = 183

Cluster 2 -> Length = 35

Cluster 3 -> Length = 351

Cluster validation Indices (rounded to 4 decimal places):

Silhouette Index = 0.2712

Davies Bouldin Index = 2.5451

Dunne Index = 0.0556

Wine Dataset

Cluster 1 -> Length = 23

Cluster 2 -> Length = 104

Cluster 3 -> Length = 51

Cluster validation Indices (rounded to 4 decimal places):

Silhouette Index = -0.0554

Davies Bouldin Index = 3.0640

Dunne Index = 0.0148

Infomap Algorithm

Infomap is a graph-based clustering algorithm where each data point can be treated as a node with edges connecting pairs of them. The weight of an edge corresponds to the euclidean distance between them.

For the implementation, an average value of edge weights is determined and all edges with weights less than that are dropped. The final set of edges (in a .gml file) is given as input to igraph's community_infomap.

Iris Dataset

```
Cluster 1 -> Length = 52
Cluster 2 -> Length = 97
Cluster 3 -> Length = 1
Cluster validation Indices (rounded to 4 decimal places): Silhouette Index = 0.4218
Davies Bouldin Index = 0.4220
Dunne Index = 0.4664
```

Breast Cancer Wisconsin Dataset

```
Cluster 1 -> Length = 460
Cluster 2 -> Length = 216
Cluster 3 -> Length = 6
Cluster 4 -> Length = 1
Cluster validation Indices (rounded to 4 decimal places):
Silhouette Index = 0.3982
Davies Bouldin Index = 1.0675
Dunne Index = 0.4026
```

Wine Dataset

```
Cluster 1 -> Length = 47
Cluster 2 -> Length = 131
Cluster validation Indices (rounded to 4 decimal places):
Silhouette Index = 0.6572
Davies Bouldin Index = 0.4565
Dunne Index = 0.8579
```

Clustering Algorithm developed in Assignment-1

Iris Dataset

```
Cluster 1 -> Length = 86
Cluster 2 -> Length = 50
Cluster 3 -> Length = 14
Cluster validation Indices (rounded to 4 decimal places):
Silhouette Index = 0.0309
Davies Bouldin Index = 2.2610
Dunne Index = 0.1293
```

• Breast Cancer Wisconsin Dataset

Only the first 200 data points are used to reduce the extensive computation time needed by this algorithm.

```
Cluster 1 -> Length = 135
Cluster 2 -> Length = 63
```

Cluster validation Indices (rounded to 4 decimal places):

Silhouette Index = 0.4415

Davies Bouldin Index = 1.0738

Dunne Index = 0.4662

Seeds Dataset

Cluster 1 -> Length = 39

Cluster 2 -> Length = 58

Cluster 3 -> Length = 113

Cluster validation Indices (rounded to 4 decimal places):

Silhouette Index = 0.3599

Davies Bouldin Index = 1.1377

Dunne Index = 0.3642

Task-2: Classification Analysis

10 fold cross validation is done for all cases. The intermediate results are not presented here for brevity, only the final results are given for each case.

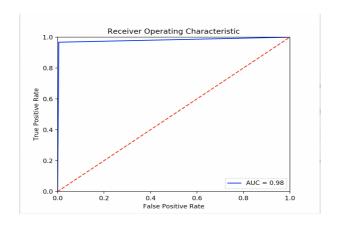
Decision Tree

• BankNote Authentication Dataset

Accuracy: 0.982497619803237 Specificity: 0.9828352222510384 Sensitivity: 0.9820866935483871 Precision: 0.9789299717514691 F-measure: 0.9803762035555337

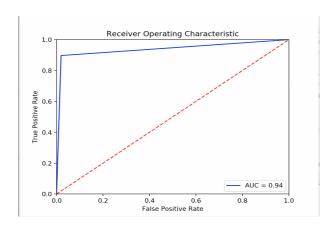
Prediction: 760 612

Actual Class label: 762 610



• Breast Cancer Wisconsin Dataset

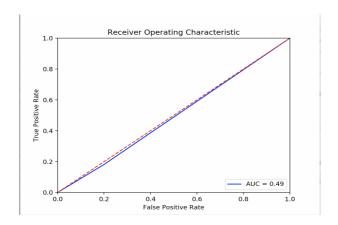
Accuracy: 0.9501705029838023 Specificity: 0.9614337520630845 Sensitivity: 0.930331616853356 Precision: 0.9305259595476987 F-measure: 0.9290235519235841 Prediction: 444 239 Actual Class label: 444 239



· Haberman's Survival Dataset

Accuracy: 0.666236559139785 Specificity: 0.7749470217044588 Sensitivity: 0.4202633477633477 Precision: 0.37238095238095237 F-measure: 0.3753861369109047

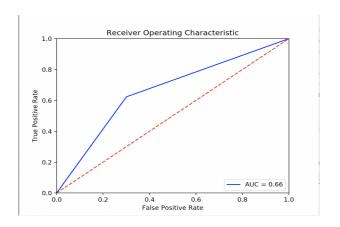
Prediction: 225 81 Actual Class label: 225 81



• Pima Indian Diabetes Dataset

Accuracy: 0.7031100478468899 Specificity: 0.7805294970986459 Sensitivity: 0.5661191805715133 Precision: 0.5792382641253608 F-measure: 0.5662043283931991

Prediction: 508 260



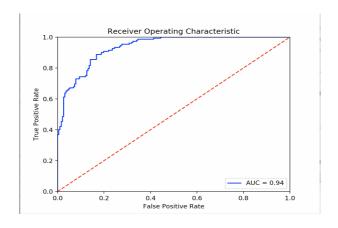
Naïve Bayes

• BankNote Authentication Dataset

Accuracy: 0.8389664656722733 Specificity: 0.8787525628364371 Sensitivity: 0.7926724168785175 Precision: 0.8386117247270881 F-measure: 0.8132050044960154

Prediction: 797 575

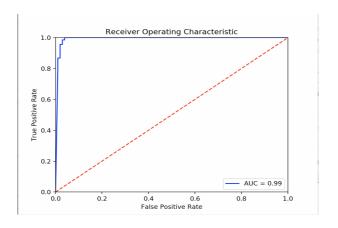
Actual Class label: 762 610



• Breast Cancer Wisconsin Dataset

Accuracy: 0.9633205456095482 Specificity: 0.9558769143796896 Sensitivity: 0.9802900137232973 Precision: 0.9233205240813935 F-measure: 0.9488355568988098

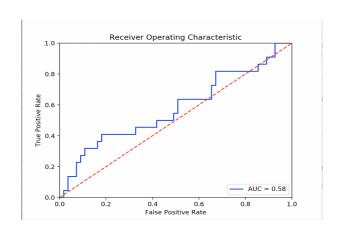
Prediction: 429 254



Accuracy: 0.7450537634408602 Specificity: 0.9385720873329569 Sensitivity: 0.211019536019536 Precision: 0.5511904761904762 F-measure: 0.28564102564102567

Prediction: 275 31

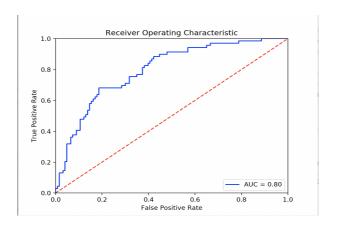
Actual Class label: 225 81



• Pima Indian Diabetes Dataset

Accuracy: 0.7592105263157896 Specificity: 0.8425872995663299 Sensitivity: 0.6082767775518201 Precision: 0.6692941963084392 F-measure: 0.6330167235470293

Prediction: 527 241



KNN

• BankNote Authentication Dataset

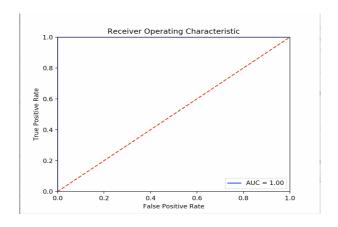
Accuracy: 0.9956310166084841 Specificity: 0.9921013090555345

Sensitivity: 1.0

Precision: 0.9904764638346727 F-measure: 0.9951569442389516

Prediction: 756 616

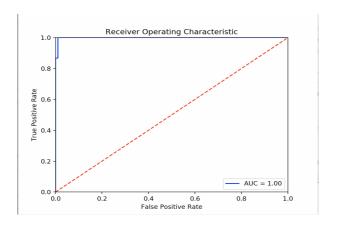
Actual Class label: 762 610



• Breast Cancer Wisconsin Dataset

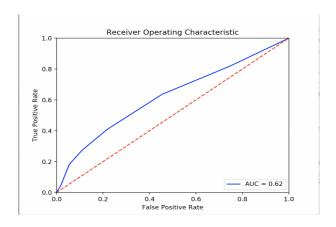
Accuracy: 0.9663682864450127 Specificity: 0.9748283800563758 Sensitivity: 0.95367297979799 Precision: 0.956188961900169 F-measure: 0.9538943469414246

Prediction: 445 238



Accuracy: 0.725268817204301 Specificity: 0.9324858886346302 Sensitivity: 0.16428571428571428 Precision: 0.44000000000000006 F-measure: 0.22576923076923078

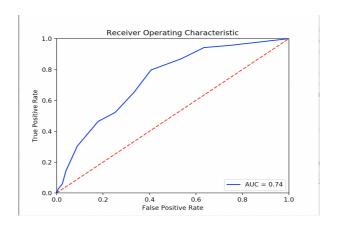
Prediction: 279 27 Actual Class label: 225 81



• Pima Indian Diabetes Dataset

Accuracy: 0.7345181134654819 Specificity: 0.8537733948309499 Sensitivity: 0.505056462056462 Precision: 0.654110233778426 F-measure: 0.5661286781167465

Prediction: 560 208

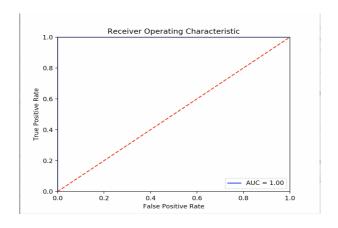


SVM

• BankNote Authentication Dataset

Accuracy: 1.0 Specificity: 1.0 Sensitivity: 1.0 Precision: 1.0 F-measure: 1.0 Prediction: 762 610

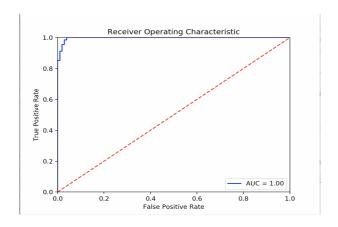
Actual Class label: 762 610



• Breast Cancer Wisconsin Dataset

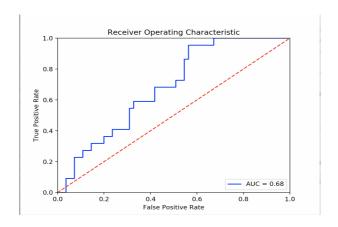
Accuracy: 0.9706947996589939 Specificity: 0.9674638205531718 Sensitivity: 0.9758705647176411 Precision: 0.94581223893066 F-measure: 0.9598474346492758

Prediction: 436 247



Accuracy: 0.738279569892473 Specificity: 0.9280952380952382 Sensitivity: 0.21119047619047623 Precision: 0.551666666666665 F-measure: 0.293005883005883

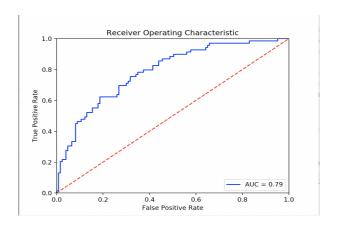
Prediction: 273 33 Actual Class label: 225 81



• Pima Indian Diabetes Dataset

Accuracy: 0.7578092959671907 Specificity: 0.8717053464237511 Sensitivity: 0.5382652975089014 Precision: 0.6884103608868742 F-measure: 0.6008505361904428

Prediction: 558 210



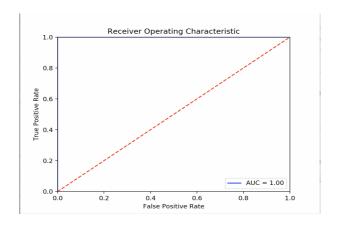
Bagging

• BankNote Authentication Dataset

Accuracy: 0.9890722521950703 Specificity: 0.9896917509541462 Sensitivity: 0.9877610134963074 Precision: 0.9864901343593644 F-measure: 0.9870368301341568

Prediction: 761 611

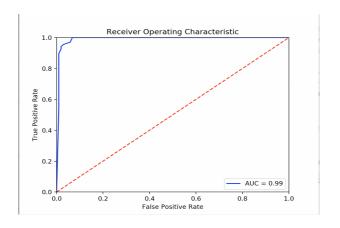
Actual Class label: 762 610



• Breast Cancer Wisconsin Dataset

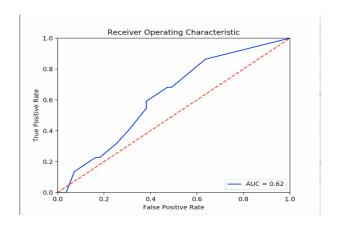
Accuracy: 0.954688832054561 Specificity: 0.9653364376651545 Sensitivity: 0.9291794364314386 Precision: 0.9425795315795316 F-measure: 0.9340692263711133

Prediction: 445 238



Accuracy: 0.6762365591397849 Specificity: 0.8227567441084066 Sensitivity: 0.2894993894993895 Precision: 0.3866666666666666666 F-measure: 0.30789904539904545

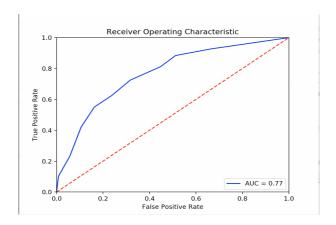
Prediction: 244 62 Actual Class label: 225 81



• Pima Indian Diabetes Dataset

Accuracy: 0.7435577580314423 Specificity: 0.8529084524070729 Sensitivity: 0.5348681311816619 Precision: 0.6604950540602714 F-measure: 0.5862814191475204

Prediction: 549 219



For each classifier, I have computed the following performance measure metrics using 10-fold cross validation.

- 1. Accuracy
- 2. Specificity
- 3. Sensitivity
- 4. Precision
- 5. Recall
- 6. F-Measure

Also, I have included the number of samples predicted as belonging to each class versus actually belonging to each class in the displayed output. And I have drawn the ROC curves for each case as well.

Four different datasets have been used for testing each of the five classifiers.

- 1. BankNote Authentication Dataset
- 2. Breast Cancer Wisconsin Dataset
- 3. Haberman's Survival Dataset
- 4. Pima Indian Diabetes Dataset

On the basis of all the analysis performed, one general conclusion that I can draw is that the BankNote Authentication Dataset and the Breast Cancer Wisconsin Dataset seem relatively easily to classify or cluster because all algorithms achieve nearly perfect results for these 2 cases. Whereas for the other datasets, the algorithms only have average performance.