## VIVEKANAND EDUCATION SOCIETY'S INSTITUTE OF TECHNOLOGY

Hashu Advani Memorial Complex, Collector's Colony, R C Marg, Chembur, Mumbaí-400074



## Department of Artificial Intelligence and Data Science

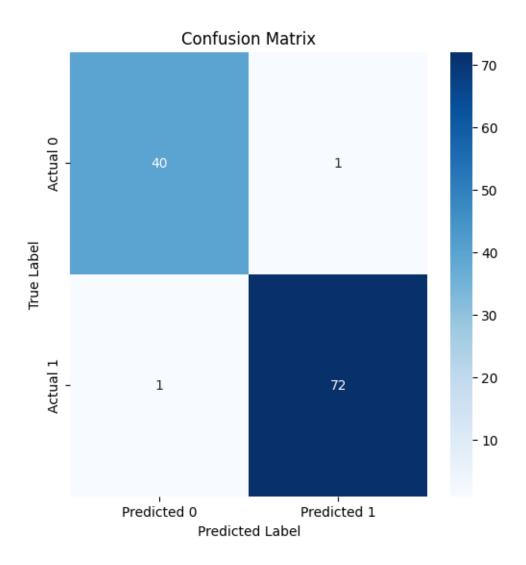
Subject:	Class: D11AD			Semester: VI	
Roll No.: 26	Name:  Dyotak Kac	hare			
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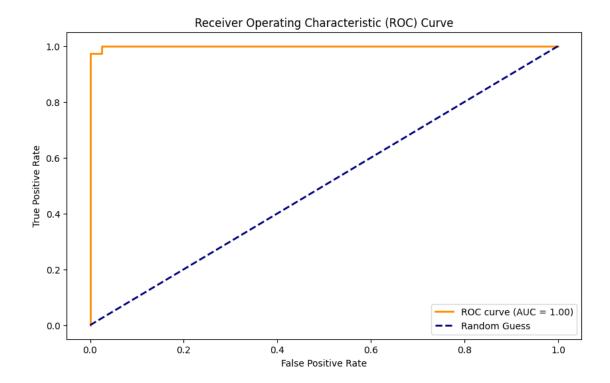
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	Kernel trick for non linear SI	VM.
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	Theory-	913.455
	us can raily reprose data	with
-)-	hyperplane by drawing a st.	raight lini
	is called linear SVM.	RAGINALIS.
	when we rannot reperate data wir	n a swaight
	line we use Non-linear SVM. In	this, we
	have kernel functions.	diameter Alb
	21 transforms data into another	aumension in
	that the data can be classified.	· Conclusion
	Kernel tricks	
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	Profunctial pernel:	min all
-	A odunomial served is a king	d of SVM
	kernel that uses a porgnomial	guy) guoi)
	map the data into a night mi	nepsion a
	Is does this by taking the dot	
	of the data points in the origin	al space
	and the polynomial function in	the new
	spare.	
	K(n,,n2): (n, <sup>†</sup> n246) <sup>d</sup>	
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Fage No.: Subject rotad \_\_\_\_ RBF Kornald transports IM The squared encledion distance is multiplied by the gama parameter and then beinding the enponent of the whole. white 1. = is the variance, & hyperparameter
2. 11 X1- X211 is the Cuiledian (1, Norm) distance between two points  $X_1$  &  $X_2$   $K(X_1, X_2) = \exp\left(-\frac{11}{2}X_1 - \frac{11}{2}X_2\right)$   $= 20^{2}$ It that the to the danshed. Conclusion Thus, we have successfully implemented Non luxer SVM. Non luser SVM. It does the by taking the dat product and the polingered hundrer in the new K(min) (m, wete) MI DOES

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[14]: from sklearn.datasets import load breast cancer
    from sklearn.svm import SVC
    from sklearn.preprocessing import StandardScaler
    from sklearn.model selection import train test split
    from sklearn.metrics import roc curve, auc, confusion matrix, classification report
    import matplotlib.pyplot as plt
    import seaborn as sns
    import numpy as np
[3]: data = load breast cancer()
    X, y = data['data'], data['target']
[4]: print(X)
    print(y)
    [[1.799e+01 1.038e+01 1.228e+02 ... 2.654e-01 4.601e-01 1.189e-01]
     [2.057e+01 1.777e+01 1.329e+02 ... 1.860e-01 2.750e-01 8.902e-02]
     [1.969e+01 2.125e+01 1.300e+02 ... 2.430e-01 3.613e-01 8.758e-02]
     [1.660e+01 2.808e+01 1.083e+02 ... 1.418e-01 2.218e-01 7.820e-02]
     [2.060e+01 2.933e+01 1.401e+02 ... 2.650e-01 4.087e-01 1.240e-01]
     [7.760e+00 2.454e+01 4.792e+01 ... 0.000e+00 2.871e-01 7.039e-02]]
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     1 1 1 1 1 1 1 0 0 0 0 0 0 1
[5]: scaler = StandardScaler()
    X scaled = scaler.fit transform(X)
[6]: X_train, X_test, y_train, y_test = train_test_split(X_scaled, y, test_size=0.2)
[22]: model = SVC(probability=True)
    model.fit(X_train, y_train)
[22]: SVC(probability=True)
[9]: print(model.score(X_test, y_test))
    0.9824561403508771
[15]: y_predict = model.predict(X_test)
```

```
[32]: for i in range(20):
          print(f"Actual value: {y_test[i]} | Predicted value: {y_predict[i]}")
     Actual value: 1 | Predicted value: 1
     Actual value: 0 | Predicted value: 0
     Actual value: 1 | Predicted value: 1
     Actual value: 0 | Predicted value: 0
     Actual value: 1 | Predicted value: 1
     Actual value: 1 | Predicted value: 1
     Actual value: 0 | Predicted value: 0
     Actual value: 0 | Predicted value: 0
     Actual value: 0 | Predicted value: 0
[19]: cm = confusion_matrix(y_test, y_predict)
      plt.figure(figsize=(6, 6))
      sns.heatmap(cm, annot=True, fmt='d', cmap='Blues', xticklabels=['Predicted 0', Predicted 0']
       plt.xlabel('Predicted Label')
      plt.ylabel('True Label')
      plt.title('Confusion Matrix')
      plt.show()
```





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