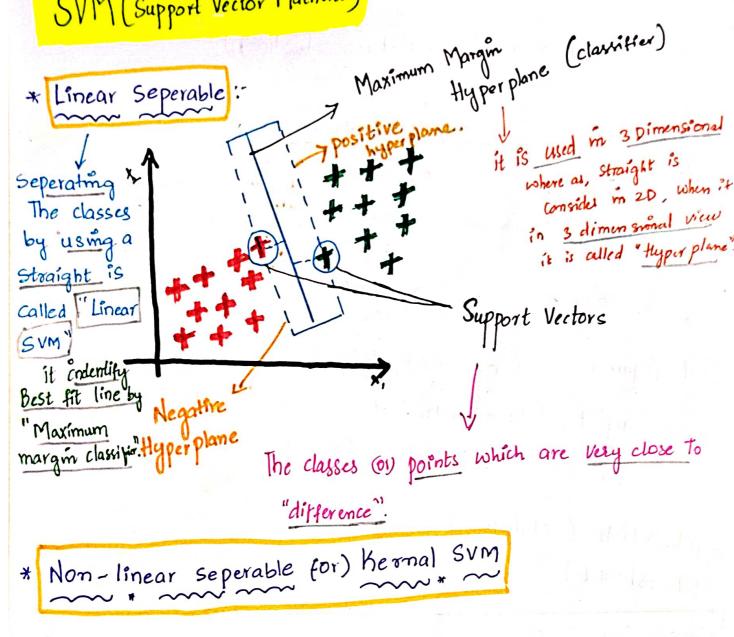
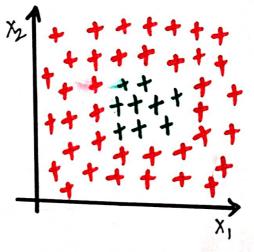
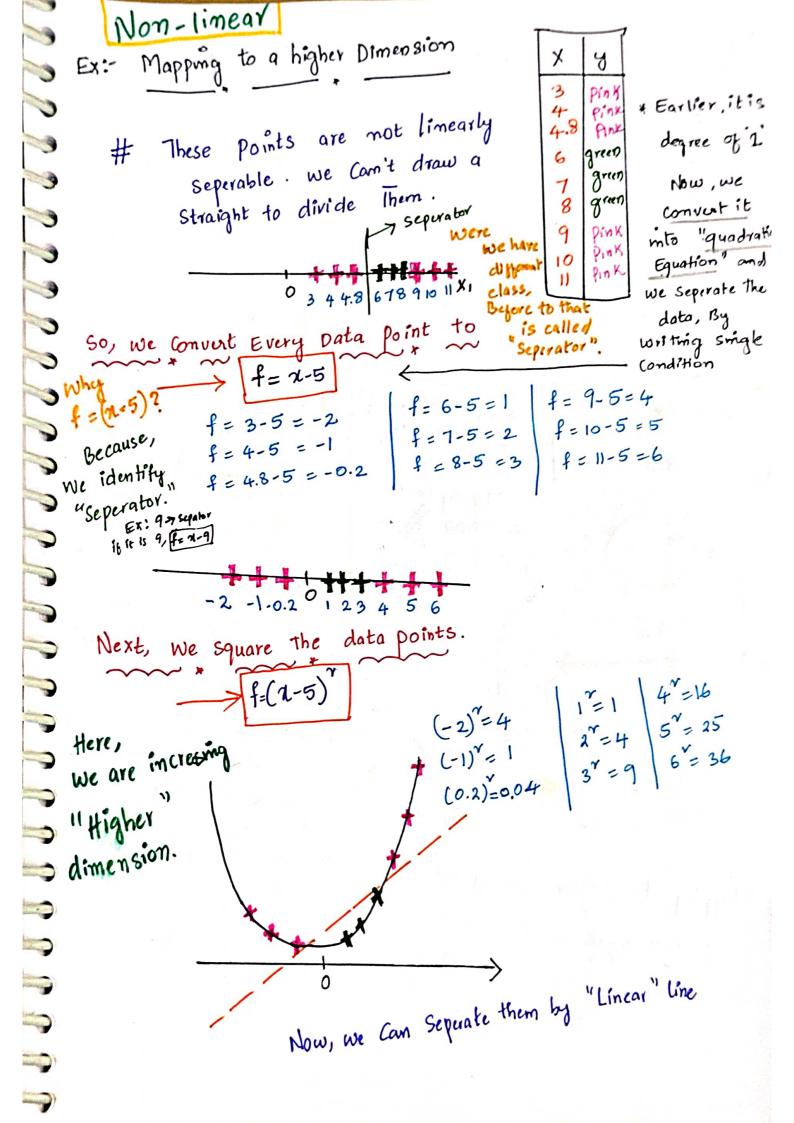
29/04/2 1.00 Pm

## SVM (Support Vector Machines)

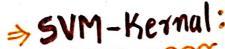


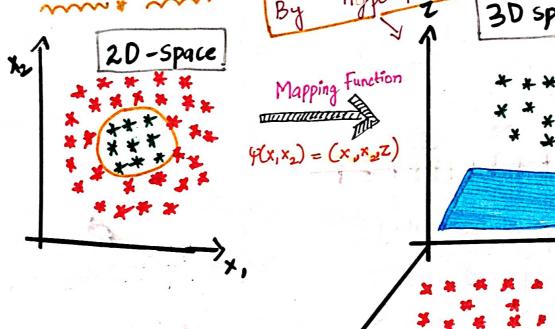


\* In This sitution, we use "Non-linear SVM" (or) Kernal SVM

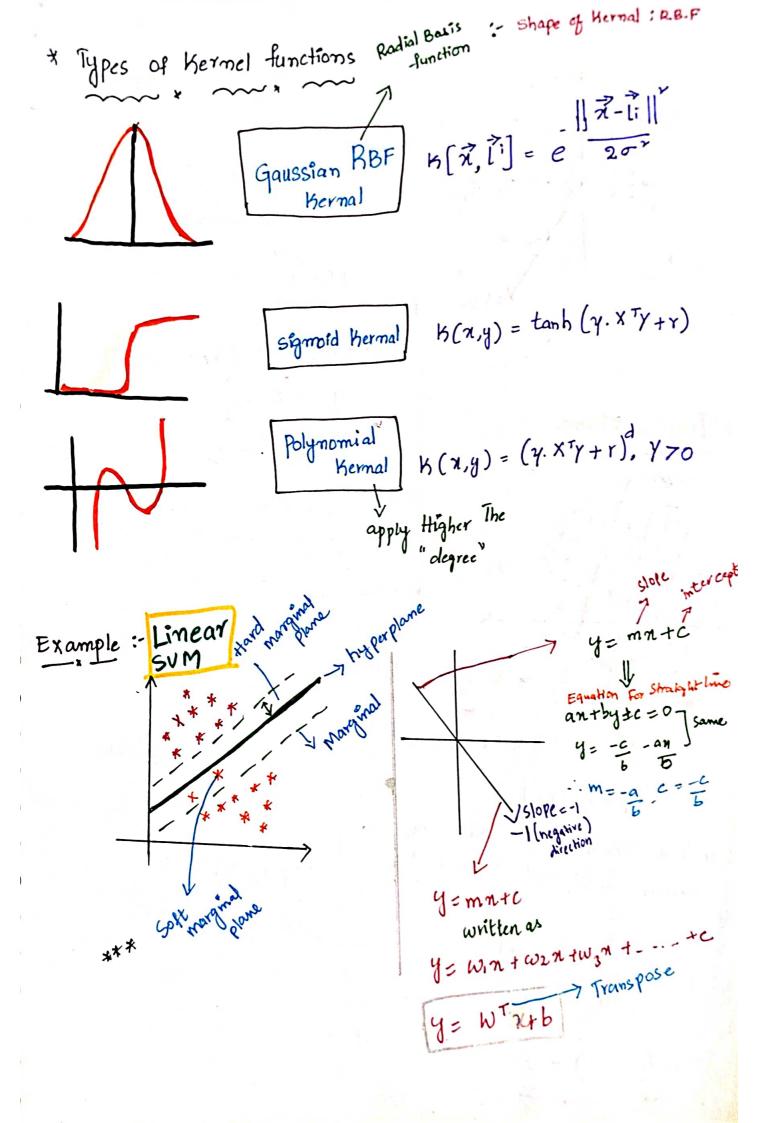


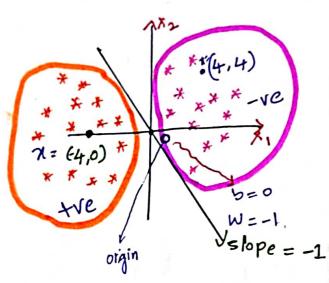
X	y	X-5	(x-5)
3 4 4.8 6 7 8 9 10 11	Penk Pink Pink green green Pink Pink	-2 -1 -1 2 3 4 5 6	410014916536





\* Mapping to higher dimensional space can be Highly Compute/ intensive. More RAM needed.





$$y = \begin{bmatrix} y = W^{T}x + b \end{bmatrix}$$

$$y = \begin{bmatrix} -1 \\ 0 \end{bmatrix} \begin{bmatrix} 4 & 4 \end{bmatrix}$$

=-4+0=-4 Any point above the line is negative Value.

 $W^T \chi + b = +1[\kappa]$ 

Our Aim Should increase distance, to Perform model well Maximize)

$$W^{T}n_{1}+b=+1$$
  
 $W^{T}n_{2}+b=-1$ 

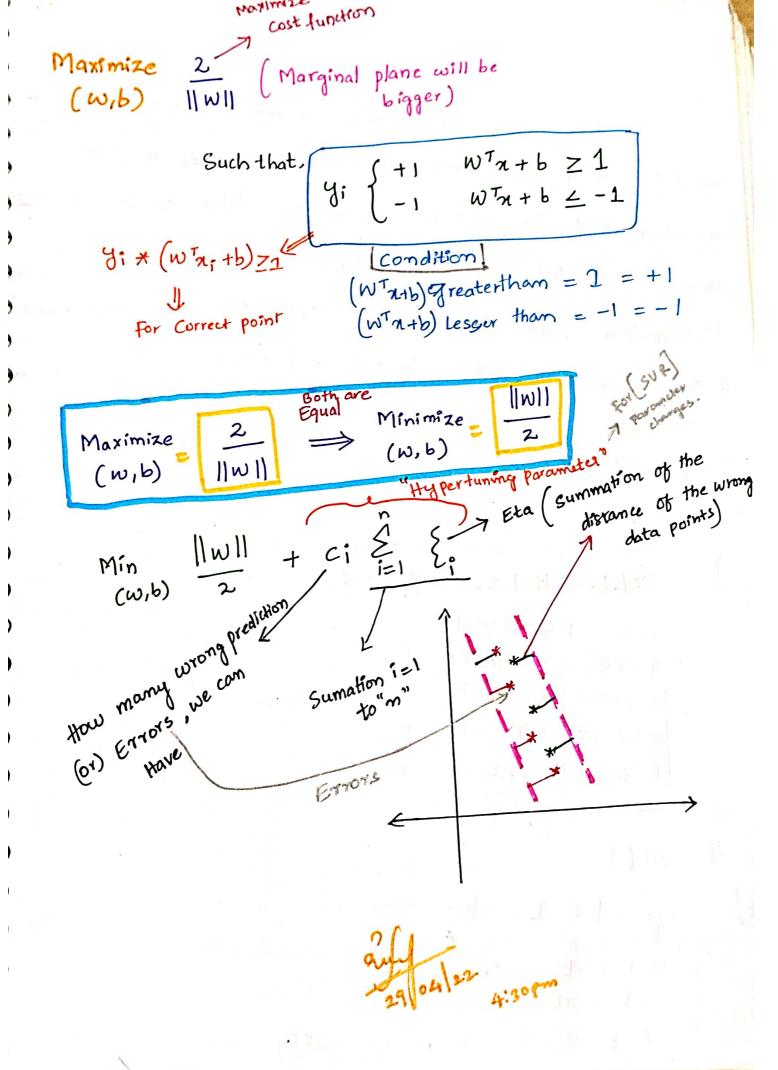
Vectors + magnitude 11 w 1

$$WT(x_1-x_2) = 2$$

$$\|W\|$$

 $W^{T}(\eta_{1}-\eta_{2})=2$  (difference between) above & below plane)

> WTA+b=0



24/2/3006

CODE !

The data Shown here Simulates a medical Study in which Data :injected with a virus were given various doses of two medicines and then checked 2 weeks later to see if they were still injected. Given this data, Our goal is to create a classification model Than Predict Cgiven two dosage measurments If they mouse will still be injected with virus. # load data, with import libraries.

# df = Pd. read\_csv ( "mouse\_viral\_study. Csv")

# df. head()

out :

med_1_mL	Med-2-mL	Virus_PreSent
6.508231 4.126116 6.427870 3.672953	8. 58 2531 3.073459 6.369 758	0

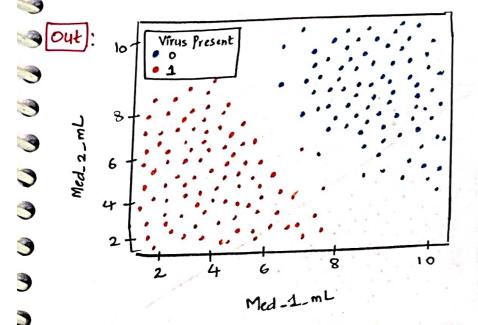
# df. info ()

Out: Range Index 400 entires 0 to 399 Float 64 o Med-1-mL 400 non null Float 64 med\_2\_ml 400 non null int 64 2 VIIII - Present 400 non null

EDA

# Sns. Scatterplot (x="Med-1-ml", y="Med-2-ml", hue="Virus preson) data = df

Plt. show ()



2 = df. drop ("Virus Present", axis=1) y = df ("Virus present"]

MODELLING

from Sklearn. Svm import SVC model = svc (Kernel = "linear")

model . fit (x,y)

SVC [Kernal = "Linear")

```
Seperating Hyper plane
#Sms. Scatterplot (n = "Med_1-ML", y = "Med_2-mL", hue = Virus Pres.
                   Palette = "Seismic", dota = df)
  # we want to some how automatically Create a Seperating hyperplane
       (a line in 20)
 # n = np. linspace (0, 10, 100)
   M = -1
    b = 41
 # y= m*71 +b
# plt.plot (2,4,"k")
                         4
  # User defined function
            Plot-svm-boundary (model, n,y):
               n = n. Values
               y = y. Values
                plt. scatter (n[:,0], n[:,1], c=y, s=30,
            # scatter plot
                                  Cmap = "seismic")
            # plot The decision function.
                an = Plt. gca ()
                xlim = an. get-xlim()
                ylim = an. get-ylim()
```

30/4/22

```
# Creating grid to Evaluate model.
         nn = np. linspace (πlim [o], nlim [1], 30)
         yy = np. lmspace (ylim[0], ylim [1], 30)
          YY, XX = np. meshgrid (yy, xx)
          My = mp. Vstack ([xx. ravel(), YY. ravel()]). T
           Z = model · decision-function (xy). reshape (xx. Shape
         # plot decision boundary and margins.
          an. Contour (XX, YY, Z, colors = "K", Levels = [-1,0,1
                       alpha = 0.5, lime styles = ["--", "-", "-"
         an. scatter (model. Support_vectors-[:,0], model.
         # plot Support vectors.
                 Support-vectors-[:,1], s=100, linewidth=1,
                  face colours = "mone", edge colors = "k")
         Plt. show ()
# plot-svm-boundary (model, n,y)
[out]:
```

# Our goal with SVM is to create the best seperating hyperplane with Maximum margin classifier.

# In 2 dimensions. The hyperplane is Simply a line.

In 3 dimensional -> plane

Straight

Hyper Parameters

out ?

#C: Regularization parameter.

# The Strength of the originalization is inversely proportional

# The Strength of the original zation is inversely proportional

to 'C". Must be strictly positive. The penality is a

to 'C". Must be strictly positive. The penality is a

(less sensitility).

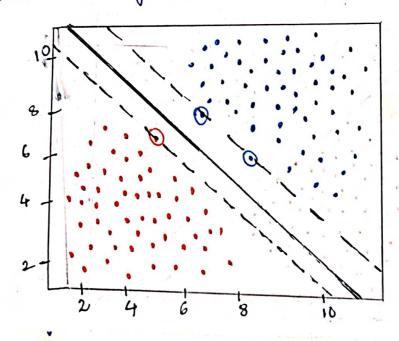
Squared 12 penalty. # Higher The "c" value - lesser The oniss classification

# C \ (# lower The "c" value - in correct answer

# model = SVC (Kernel = "Linear", C = 1000)

# model fit (n,y)

plot\_svm\_boundary (model, 2,y)

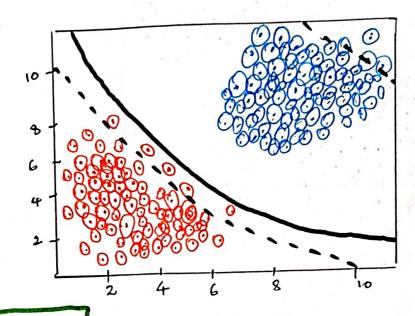


```
# model = SVC (Kernel = "linear", C = 0.05)
       model. fit (x,y)
         plot - sym_ (model , x,y)
    #
            boundary
    out:
                       000000
           8
           6
           2
3
     # Kernal
3
3
        1. RBF
     model = SVM (Kernel = "rbf", C=1)
3
3
model. fit (n,y)
      plot-sym-boundary (model, n,y)
  out]:
          10
          8
                                   10
```

3

### 2. Sigmoid

#### out:



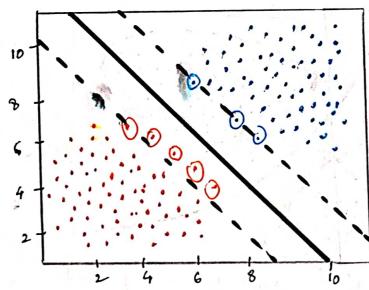
### 3. Polynomial

model = SVC (Kernel = "poly", C=1, degree = 1

model . fit (71,4)

plot - sum - boundary (model, n, y)

### out:



# model = svc (Kernel = "poly", , c = i, clegree = 2) # model · fit (7,4) Quadratic # plot-SVm-boundary (model, 1,y) curve. out 10 8 6 4 gamma: { scale, auto } or float # Default = "scale" Kernel Coefficient for "rbf", "poly" & "sigmo" # gamma "gamma = Scale" > Trum. of feature x x. Var () # model = SVC (Kernel = "rbf", C= 1, gamma = OA) 1 n-features. # model . fit (n,y) # plot-svm\_boundary (model, n, y) out i 6

# Hyper parameter Tuning

1.0

out :

```
from Skleam, model_selection import Grid Search CV
# estimator = svc()
# param - grid = { "c": [0,0.01,0.1], "Kernel": ["Inear", "rbf"]
                     , "gamma": [0.01,0.02,0.]}
# grid = Grid Search CV (estimator, param-grid, CV=5)
# grid . fit (n,y)
       Grid Search CV (cv=5, estimator = SVC (), Param-grid =
                {"c": [0,0.01,0.1], "Kermel": ("linear", "15f"), gamma
                  [0.01, 0.02, 0.1]}
           'c": 0.01, "gamma": 0.01, "kernel": "rbp")
# grid. best-params -
                                If we don't write
   grid. best_score -
                                      garnina,
                                           "lineal
                                      greed best prediction
```

and accurany.

. : Only for This grolden

```
tram test Split
from Sklearn. model_selection import train_test-split
# n_tram, n_test, y_tram, y_test = tram-test-split (x,y,
                      test-size = 0.25, random_state = 29)
Scaling
from Sklearn. Preprocessing import Standard Scaler

# Sc = Standard Scaler ()
# n_train = Sc. fit - transform (n_train)
# n_test = Sc. fit_transform (n_test)
model
# model = SVC (Kernel = "rbf", C = 0.01, gamma = 0.01)
# model. fit (n.tram, y.tram)
out SVC (c=0.01, gamma = 0.01, kernel = "Linear")
# ypred_tram = model. (n_tram)
# 4 pred-test = model. Predict (n-test)
```

accuracy from Sklearn. metrics importacuracy\_score # accuracy-score (y-test, y pred-test) out : Que: hlhat is "model Selection"? Ans: Identify The best Algorithm with best Hyperpan By applying hyperparameter "tunning"