quiz3 st121411 Rom

October 19, 2020

1 QUIZ 3 | st121411 - Rom Parnichkun

The quiz is designed to be completed in 120 minutes and they are all about Classification. The quiz will start at 1 PM 19th October 2020 till 3 PM 19th October 2020.

- You may use puffer/jupyter.
- You may use online resources.
- You will not be graded if submit late.
- You are not allowed to have a conversation with a human being. (A conversation with plants and pets is fine.)
- *** Submit in PDF and jupyter notebook format ***
- *** Write your student ID and student Name on the head of this file and save the file in this naming format. quiz3 stid name***

Given 2 datasets, perform a classification on both datasets and answer the followings question.

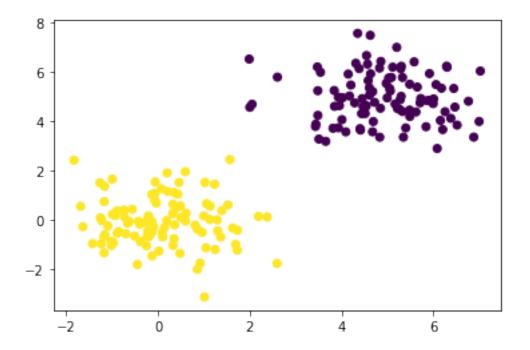
- 1. plot the data using a scatter plot. (2 pt)
- 2. Base on the data you saw, which method will you choose wo perform and why? (2 pts)
- 3. Perform the classification from scratch. (4 pts)
- 4. Is the choosen method yield a good result? If not, why? (2 pts)

2 Dataset 1 (10 pts)

[1]:

```
[2]: # 1. plot the data using a scatter plot. (2 pt)
     import matplotlib.pyplot as plt
     import numpy as np
     dataset1_np = np.array(dataset1)
    plt.scatter(dataset1_np[:,0],dataset1_np[:,1],c=dataset1_np[:,2])
                                            2
```

[2]: <matplotlib.collections.PathCollection at 0x7faa788c3940>



[3]: # 2. Base on the data you saw, which method will you choose wo perform and why? □ → (2 pts)

print("The two blobs seem to be normally distributed therefore I will use the □ → gaussian discrimant analysis to model this data")

The two blobs seem to be normally distributed therefore I will use the gaussian discrimant analysis to model this data ${\sf data}$

```
[4]: # 3. Perform the classification from scratch. (4 pts)
class GDAModel:

def parameters(self,X,y,naive = False,same_sigma = True):
    unique_y = np.unique(y)
    prior = np.zeros(len(unique_y))
    mean = np.zeros((len(unique_y),X.shape[1]))
    sigma = np.zeros((len(unique_y),X.shape[1],X.shape[1]))
    for j, yi in enumerate(unique_y):
        prior[j] = np.sum(y==yi)/y.size
        mean[j] = np.mean(X[y==yi],axis=0)
        sigma[j] = (1/np.sum(y==yi))*(X[y==yi]-mean[j]).T@(X[y==yi]-mean[j])

# sets 0 to covariance terms if using the naive assumption
    if(naive == True):
        sigma[:,~np.eye(X.shape[1],dtype=bool)] = 0
```

```
# if we're using the same sigma for all classes, we take the average of \Box
      \rightarrow them
             if(same sigma == True):
                 sigma[:] = np.mean(sigma,axis=0)
             return prior, mean, sigma, unique_y
         def joint_probability(self,x,prior,mean,sigma):
             joint_prob = np.zeros(prior.size)
             for j,mu in enumerate(mean):
                 if(sigma.ndim == 2):
                     sig = sigma
                 else:
                     sig = sigma[j]
                 gaussian = (1/(((2*np.pi)**(x.size/2))*np.sqrt(np.linalg.
      \rightarrowdet(sig)))*np.exp(-1/2*(x-mu)@np.linalg.inv(sig)@((x-mu)))
                 joint_prob[j] = prior[j]*gaussian
             return joint_prob
         def predict(self, X, prior, mean, sigma, unique_y):
             if(X.ndim == 2):
                 y_pred = []
                 for i in range(X.shape[0]):
                      joint_prob = self.joint_probability(X[i],prior,mean,sigma)
                      y_pred.append(unique_y[np.argmax(joint_prob)])
                 y pred = np.array(y pred)
             else:
                 joint_prob = self.joint_probability(X,prior,mean,sigma)
                 y_pred = unique_y[np.argmax(joint_prob)]
             return y_pred
     #6. score/error calculation
         def accuracy(self, y, y pred):
             acc = np.sum(y == y_pred)/y.size
             return acc
[5]: from sklearn.model_selection import train_test_split
     X_train, X_test, y_train, y_test = train_test_split(dataset1_np[:,[0,1]],_

dataset1_np[:,2], test_size=0.20, random_state=42)
[6]: X_test.shape
[6]: (40, 2)
```

```
[7]: GDA = GDAModel()

prior,mean,sigma,unique_y = GDA.

→parameters(X_train,y_train,naive=False,same_sigma=True)
```

- [8]: y_pred = GDA.predict(X_test,prior,mean,sigma,unique_y)
- [9]: from sklearn.metrics import classification_report print(classification_report(y_test, y_pred))

	precision	recall	f1-score	support
0.0	1.00	1.00	1.00	16
1.0	1.00	1.00	1.00	24
accuracy			1.00	40
macro avg	1.00	1.00	1.00	40
weighted avg	1.00	1.00	1.00	40

[10]: # 4. Is the choosen method yeild a good result? If not, why? (2 pts)
print('''The chosen method was able to yield a good result and was able to

→train very quickly because the assumption that both the blobs were normal is

→sound.''')

The chosen method was able to yield a good result and was able to train very quickly because the assumption that both the blobs were normal is sound.

3 Dataset 2 (10 pts)

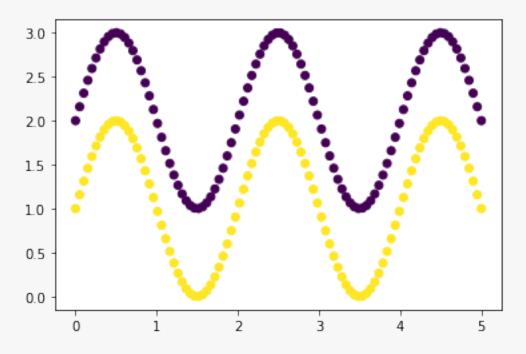
[11]:

```
[12]: # 1. plot the data using a scatter plot. (2 pt)
dataset2_np = np.array(dataset2)

print(dataset2_np.shape)
plt.scatter(dataset2_np[:,0],dataset2_np[:,1],c=dataset2_np[:,2])
```

(200, 3)

[12]: <matplotlib.collections.PathCollection at 0x7faa578f7588>



[13]: # 2. Base on the data you saw, which method will you choose wo perform and why? \hookrightarrow (2 pts)

print('from the data shown here, we can use SVM with a gaussian kernel because \hookrightarrow it will be able to accuracy classify the data. An alternative could be to \hookrightarrow use decision trees with boosting, but since decision trees ends up dividing \hookrightarrow the nodes in only 1 axis at a time, it wouldnt be as smooth as svm')

from the data shown here, we can use SVM with a gaussian kernel because it will be able to accuracy classify the data. An alternative could be to use decision trees with boosting, but since decision trees ends up dividing the nodes in only 1 axis at a time, it wouldnt be as smooth as svm

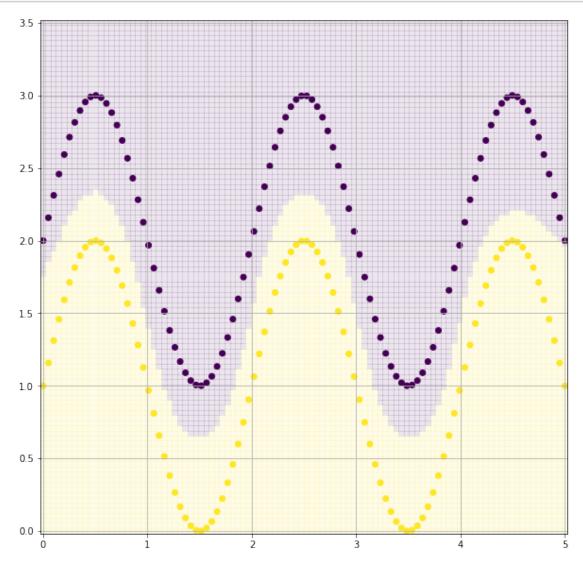
```
[14]: # 3. Perform the classification from scratch. (4 pts)
      from numpy import linalg
      import cvxopt
      import cvxopt.solvers
      import pylab as pl
      class SVMModel:
          def linear_kernel(self,x1, x2):
              return np.dot(x1, x2)
          def polynomial_kernel(self,x, y, p=2):
              return (1 + np.dot(x, y)) ** p
          def gaussian_kernel(self,x, y, sigma=0.9999):
              return np.exp(-linalg.norm(x-y)**2 / (2 * (sigma ** 2)))
          def fit(self, X, y, kernel, C):
              n_samples, n_features = X.shape
              # Gram matrix
              # initialize kernel matrix
              K = np.zeros((n_samples, n_samples))
              # Kernel matrix
              for i in range(n_samples):
                  for j in range(n_samples):
                      if kernel == 'linear_kernel':
                          K[i,j] = self.linear_kernel(X[i], X[j])
                      elif kernel == 'polynomial_kernel':
                          K[i,j] = self.polynomial_kernel(X[i], X[j])
                      else:
                          K[i,j] = self.gaussian_kernel(X[i], X[j])
              P = cvxopt.matrix(np.outer(y,y) * K)
              q = cvxopt.matrix(np.ones(n_samples) * -1)
              A = cvxopt.matrix(y, (1,n_samples))
              b = cvxopt.matrix(0.0)
              if C is None:
                  G = cvxopt.matrix(np.diag(np.ones(n_samples) * -1))
                  h = cvxopt.matrix(np.zeros(n_samples))
              else:
                  tmp1 = np.diag(np.ones(n_samples) * -1)
```

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tmp2 = np.identity(n_samples)
           G = cvxopt.matrix(np.vstack((tmp1, tmp2)))
           tmp1 = np.zeros(n_samples)
           tmp2 = np.ones(n_samples) * C
           h = cvxopt.matrix(np.hstack((tmp1, tmp2)))
       # solve QP problem
       solution = cvxopt.solvers.qp(P, q, G, h, A, b)
       # Lagrange multipliers
       a = np.ravel(solution['x'])
       # Support vectors have non zero lagrange multipliers
       sv_idx = a > 1e-5
      ind = np.arange(len(a))[sv_idx]
      a = a[sv_idx]
      sv = X[sv_idx]
       sv_y = y[sv_idx]
      print("%d support vectors out of %d points" % (len(a), n_samples))
       # Intercept
      b = 0
       for n in range(len(a)):
          b += sv y[n]
           b -= np.sum(a * sv_y * K[ind[n],sv_idx])
      b /= len(a)
       # Weight vector
      if kernel == 'linear_kernel':
           w = np.zeros(n_features)
           for n in range(len(a)):
               w += a[n] * sv_y[n] * sv[n]
       else:
           w = None
      return sv, sv_y, a, w, b
  def project(self,X, kernel,sv, sv_y, a, w, b):
       if w is not None:
           return np.dot(X, w) + b
       else:
           y_predict = np.zeros(len(X))
           for i in range(len(X)):
               for a_val, sv_y_val, sv_val in zip(a, sv_y, sv):
                   if kernel == 'polynomial_kernel':
                       s += a_val * sv_y_val * self.polynomial_kernel(X[i],_
→sv_val)
```

```
else:
                              s += a_val * sv_y_val * self.gaussian_kernel(X[i],_
       ⇒sv_val)
                      y predict[i] = s
                  return y_predict + b
          def predict(self, X, kernel, sv, sv_y, a, w, b):
              return np.sign(self.project(X, kernel,sv, sv_y, a, w, b))
          def plot_contour(self,X1_train, X2_train, kernel, sv, sv_y, a, w, b):
              pl.plot(X1_train[:,0], X1_train[:,1], "ro")
              pl.plot(X2_train[:,0], X2_train[:,1], "bo")
              pl.scatter(sv[:,0], sv[:,1], s=100, c="g")
              # here we choose the range between -7 and 7 as we have choosen
              # the mean to be between -4 and 4 while generating data with the
       \rightarrow variance of 0.8
              X1, X2 = np.meshgrid(np.linspace(-7,7,50), np.linspace(-7,7,50))
              X = np.array([[x1, x2] for x1, x2 in zip(np.ravel(X1), np.ravel(X2))])
              Z = self.project(X, kernel,sv, sv_y, a, w, b).reshape(X1.shape)
              pl.contour(X1, X2, Z, [0.0], colors='k', linewidths=1, origin='lower')
              pl.contour(X1, X2, Z + 1, [0.0], colors='grey', linewidths=1, __
       →origin='lower')
              pl.contour(X1, X2, Z - 1, [0.0], colors='grey', linewidths=1, __
       →origin='lower')
              pl.axis("tight")
              pl.show()
[15]: X_train, X_test, y_train, y_test = train_test_split(dataset2_np[:,[0,1]],__
       →dataset2_np[:,2], test_size=0.20, random_state=42)
      y_train[y_train==0] = -1
      y_test[y_test==0] = -1
[16]: SVM = SVMModel()
      kernel = 'gaussian kernel'
      sv, sv_y, a, w, b = SVM.fit(X_train,y_train,kernel=kernel,C=None)
          pcost
                      dcost
                                         pres
                                                dres
                                  gap
      0: -6.9496e+01 -1.9281e+02 5e+02 1e+01 2e+00
      1: -1.7109e+02 -2.9223e+02 2e+02 6e+00 9e-01
      2: -2.7725e+02 -3.8134e+02 1e+02
                                         2e+00 4e-01
      3: -3.1719e+02 -3.9262e+02 8e+01 5e-01 9e-02
      4: -3.3379e+02 -3.5527e+02 2e+01 1e-01 2e-02
      5: -3.4160e+02 -3.5062e+02 9e+00 4e-03 6e-04
```

```
6: -3.4564e+02 -3.4767e+02 2e+00 7e-04 1e-04
     7: -3.4659e+02 -3.4706e+02 5e-01
                                   2e-05 3e-06
     8: -3.4689e+02 -3.4699e+02 1e-01
                                   4e-06 6e-07
     9: -3.4692e+02 -3.4698e+02 6e-02 2e-06 2e-07
    10: -3.4697e+02 -3.4697e+02 2e-03 3e-08 5e-09
    11: -3.4697e+02 -3.4697e+02 2e-05 3e-10 5e-11
    Optimal solution found.
    25 support vectors out of 160 points
[17]: | y_pred = SVM.predict(X_test,kernel,sv,sv_y,a,w,b).astype(int)
     print("y_test:",y_test.astype(int))
     print("y_pred:",y_pred)
     from sklearn.metrics import classification_report
     print(classification_report(y_test, y_pred))
    -1 1 1 1 -1 -1 -1 -1 1 -1 -1 1 1 1]
    -1 1 1 1 -1 -1 -1 -1 -1 1 -1 1 1 1 1
                precision recall f1-score
           -1.0
                     1.00
                              1.00
                                      1.00
                                                 17
            1.0
                     1.00
                              1.00
                                      1.00
                                                 23
                                      1.00
                                                 40
        accuracy
                     1.00
                              1.00
                                      1.00
                                                 40
       macro avg
                                      1.00
    weighted avg
                     1.00
                              1.00
                                                 40
[18]: resolution = 100
     x_series = np.linspace(0,5,resolution)
     y_series = np.linspace(0,3.5,resolution)
     x_mesh,y_mesh = np.meshgrid(x_series,y_series)
     x_mesh = x_mesh.reshape(-1,1)
     y_mesh = y_mesh.reshape(-1,1)
     mesh = np.append(x_mesh,y_mesh,axis=1)
     y_pred = SVM.predict(mesh,kernel,sv,sv_y,a,w,b).astype(int)
     x_mesh = x_mesh.reshape(resolution,resolution)
     y_mesh = y_mesh.reshape(resolution,resolution)
     y_pred = y_pred.reshape(resolution, resolution)
```

```
[19]: plt.figure(figsize=(10,10))
    plt.scatter(dataset2_np[:,0],dataset2_np[:,1],c=dataset2_np[:,2])
    plt.pcolormesh(x_mesh,y_mesh,y_pred,cmap='viridis',shading='auto',alpha=0.1)
    plt.grid(True)
```



[20]: # 4. Is the choosen method yeild a good result? If not, why? (2 pts)
print('Yes the method I used yielded a good result because SVMs with a gaussian

→kernel can classify any data as long as they arent overlapping, although

→SVMs can have disadvantages such as slow training time, for this small

→dataset, it is able to train very quickly')

Yes the method I used yielded a good result because SVMs with a gaussian kernel can classify any data as long as they arent overlapping, although SVMs can have disadvantages such as slow training time, for this small dataset, it is able to

train very quickly