Q1.

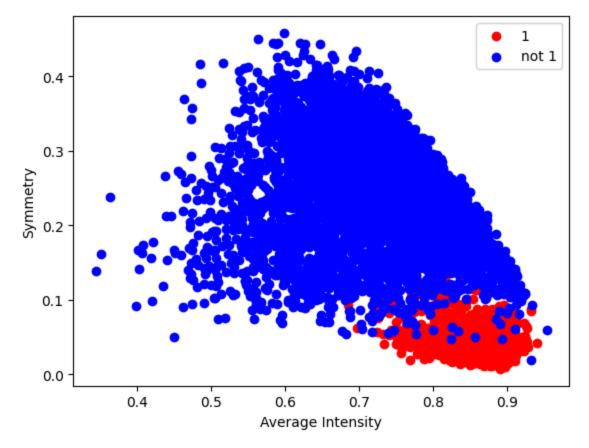
Revisit the handwritten digit recognition problem (reproduce results from chapter "Nonlinear Transformation"). Separate digit 1 from all the other digits, using intensity and symmetry as your inputs variables like you did before. Use the same data set sent to you before.

```
In [1]: import numpy as np
        import matplotlib.pyplot as plt
        import os
In [2]: data_path = "/Users/chuan/Desktop/c815_course_hws/hw2/Data"
        train_data_path = os.path.join(data_path, "DigitsTraining.csv")
        test data path = os.path.join(data path, "DigitsTesting.csv")
        train_data = np.genfromtxt(train_data_path, delimiter=',')
        # extract the first column as it is label
        train labels = train data[:,0]
        # extract the rest columns as it is features
        train features = train data[:,1:]
        # train data = (train labels, train features)
        test_data = np.genfromtxt(test_data_path, delimiter=',')
        # extract the first column as it is label
        test labels = test data[:,0]
        # extract the rest columns as it is features
        test_features = test_data[:,1:]
        # test_data = (test_labels, test_features)
        # label digit 1 as 1, others as -1
        Y train = np.where(train labels == 1, 1, -1)
        Y_test = np.where(test_labels == 1, 1, -1)
        train_features.shape, test_features.shape, Y_train.shape, Y_test.shape
Out[2]: ((7291, 256), (2007, 256), (7291,), (2007,))
In [3]: def cal avg intensity(single digit):
            return np.mean(single_digit)
        def cal symmetry(single digit):
            # reshape the single digit to 16*16
            single_digit = single_digit.reshape(16, 16)
            vertical symmetry = np.mean(np.abs(single digit - np.flip(single digit,
            horizontal_symmetry = np.mean(np.abs(single_digit - np.flip(single_digit
            return (vertical_symmetry + horizontal_symmetry) / 2
In [4]: # use intensity and symmetry as features, to create a matrix of features X t
        X train = np.zeros((train features.shape[0], 2))
        X_train[:, 0] = np.apply_along_axis(cal_avg_intensity, 1, train_features)
        X_train[:, 1] = np.apply_along_axis(cal_symmetry, 1, train_features)
```

```
# also for X_test
X_test = np.zeros((test_features.shape[0], 2))
X_test[:, 0] = np.apply_along_axis(cal_avg_intensity, 1, test_features)
X_test[:, 1] = np.apply_along_axis(cal_symmetry, 1, test_features)
```

```
In [5]: # plot the features
plt.scatter(X_train[Y_train == 1][:, 0], X_train[Y_train == 1][:, 1], c='r',
plt.scatter(X_train[Y_train == -1][:, 0], X_train[Y_train == -1][:, 1], c='t
plt.xlabel('Average Intensity')
plt.ylabel('Symmetry')
plt.legend()
```

Out[5]: <matplotlib.legend.Legend at 0x1079c3e50>



```
X_transformed[:, 0:3] = X
X_transformed[:, 3] = X[:, 1] ** 2
X_transformed[:, 4] = X[:, 2] ** 2
X_transformed[:, 5] = X[:, 1] * X[:, 2]
X_transformed[:, 6] = X[:, 1] ** 3
X_transformed[:, 7] = X[:, 2] ** 3
X_transformed[:, 8] = X[:, 1] ** 2 * X[:, 2]
X_transformed[:, 9] = X[:, 2] ** 2 * X[:, 1]
return X_transformed
```

(a)

Use linear regression for classification. Even though, linear regression learns a real-valued function, binary-valued functions are also real-valued $\pm 1 \in R$. Thus, you can use linear regression to compute w and approximate your binary classification wT xn \approx yn = ± 1 . Use your result for w to compute sign(wT xn) and report the value for Ein and Eout.

```
In [7]: Reg = Regression()
        # add bias term to X_train, and X_test
        X_train = np.hstack((np.ones((X_train.shape[0], 1)), X_train))
        X_test = np.hstack((np.ones((X_test.shape[0], 1)), X_test))
        Reg.fit(X_train, Y_train)
        train mse = Reg.mse(X train, Y train)
        test_mse = Reg.mse(X_test, Y_test)
In [8]: print(f"Train MSE: {train mse}")
        print(f"Test MSE: {test_mse}")
       Train MSE: 0.06089699629680428
       Test MSE: 0.0916791230692576
In [9]: # check classification accuracy
        train_acc = np.mean(Reg.predict(X_train) == Y_train)
        test_acc = np.mean(Reg.predict(X_test) == Y_test)
        print("In-sample Error Rate: ", 1-train_acc)
        print("Out-of-sample Error Rate: ", 1-test_acc)
       In-sample Error Rate: 0.015224249074201057
       Out-of-sample Error Rate: 0.02291978076731438
```

(b)

Repeat item (a) with a third-order polynomial transform Φ 3 to get a different representation of the data.

```
In [10]: # perfrom third order polynomial transformation to the features
    X_train_poly = Reg.third_order_tranform(X_train)
    X_test_poly = Reg.third_order_tranform(X_test)
In [11]: X_train_poly[1,:] # without bias term, cause the bias term will be added in
```

```
, 0.55586914, 0.34355078, 0.3089905 , 0.11802714,
Out[11]: array([1.
                 0.19096928, 0.17175828, 0.04054832, 0.10615393, 0.06560764])
In [12]: # fit the model
         Reg poly = Regression()
         Reg_poly.fit(X_train_poly, Y_train)
         train_mse_poly = Reg_poly.mse(X_train_poly, Y_train)
         test_mse_poly = Reg_poly.mse(X_test_poly, Y_test)
         print(f"Train MSE: {train_mse_poly}")
         print(f"Test MSE: {test mse poly}")
        Train MSE: 0.0565080235907283
        Test MSE: 0.07972097658196313
In [13]: # check classification accuracy
         train_acc_poly = np.mean(Reg_poly.predict(X_train_poly) == Y_train)
         test acc poly = np.mean(Reg poly.predict(X test poly) == Y test)
         print("In-sample Error Rate: ", 1-train_acc_poly)
         print("Out-of-sample Error Rate: ", 1-test acc poly)
        In-sample Error Rate: 0.014127005897682121
        Out-of-sample Error Rate: 0.019930244145490827
         (c)
         Compare Ein and Eout from (a) and (b).
In [14]: print(f"Train MSE for polynomial transformation is {train_mse_poly}")
         print(f"Train MSE for original features is {train mse}")
         print(f"Test MSE for polynomial transformation is {test mse poly}")
         print(f"Test MSE for original features is {test_mse}")
        Train MSE for polynomial transformation is 0.0565080235907283
        Train MSE for original features is 0.06089699629680428
        Test MSE for polynomial transformation is 0.07972097658196313
        Test MSE for original features is 0.0916791230692576
In [15]: print(f"In sample error rate for polynomial transformation is {1-train acc p
         print(f"In sample error rate for original features is {1-train acc}")
         print(f"Out of sample error rate for polynomial transformation is {1-test_ac
         print(f"Out of sample error rate for original features is {1-test acc}")
        In sample error rate for polynomial transformation is 0.014127005897682121
        In sample error rate for original features is 0.015224249074201057
        Out of sample error rate for polynomial transformation is 0.0199302441454908
        Out of sample error rate for original features is 0.02291978076731438
         Either compare in MSE or Accuracy for E_{in} and E_{out}, we have E_{in} < E_{out}.
         Also, result after polynomial transformation would be better, with E_{in.voly} < E_{in.original},
         and E_{out,poly} < E_{out,original}.
```

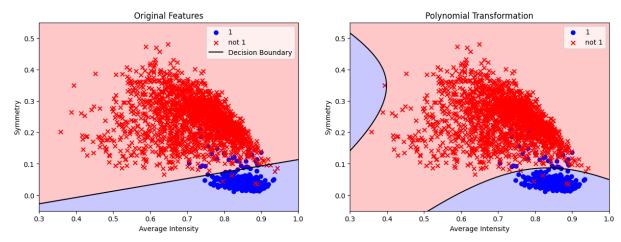
(d)

Show same plotsof Slide 13 of chapter "Nonlinear Transformation and Logistic Regression', that is, plots of the classifier boundaries for linear model and 3rd order polynomial model.

```
In [16]: X_test
Out[16]: array([[1.
                            , 0.72782227, 0.30299609],
                 [1.
                            , 0.73486719, 0.318875 ],
                 [1.
                            , 0.66407422, 0.18259766],
                 . . . ,
                            , 0.69114258, 0.33532422],
                 [1.
                            , 0.4900293 , 0.125625 ],
                 [1.
                            , 0.81102344, 0.04883203]])
                 [1.
In [17]: # plot using test data
         # 2*1 subplots for original features and polynomial transformation
         fig, axs = plt.subplots(1, 2, figsize=(15, 5))
         # original features
         axs[0].scatter(X_test[Y_test == 1][:, 1], X_test[Y_test == 1][:, 2], c='blue
         axs[0].scatter(X_test[Y_test == -1][:, 1], X_test[Y_test == -1][:, 2], c='rection{
}
         # plot the regression line
         x = np.linspace(0, 1, 100)
         y = (-Reg.weights[0] - Reg.weights[1] * x) / Reg.weights[2]
         axs[0].plot(x, y, label='Decision Boundary', color='black')
         axs[0].fill_between(x,y, 1, color='red', alpha=0.2)
         axs[0].fill_between(x,y, -0.05, color='blue', alpha=0.2)
         # set xlim and ylim to make the plot more clear
         axs[0].set_xlim(0.3, 1)
         axs[0].set xlabel('Average Intensity')
         axs[0].set_ylim(-0.05, 0.55)
         axs[0].set_ylabel('Symmetry')
         axs[0].legend()
         axs[0].set_title('Original Features')
         # polynomial transformation
         axs[1].scatter(X_test_poly[Y_test == 1][:, 1], X_test_poly[Y_test == 1][:, 2
         axs[1].scatter(X_test_poly[Y_test == -1][:, 1], X_test_poly[Y_test == -1][:,
         # plot the third order polynomial regression line
         # since the X_poly[i] is [1, x1, x2, x1^2, x2^2, x1x2, x1^3, x2^3, x1^2x2, x
         # the decision boundary is w0 + w1*x1 + w2*x2 + w3*x1^2 + w4*x2^2 + w5*x1x2
         # as a result, there may be multiple decision boundaries, so we need to plot
         y = np.linspace(-1, 1, 1000)
         X, Y = np.meshgrid(x, y)
         Z = Reg_poly.weights[0] + Reg_poly.weights[1] * X + Reg_poly.weights[2] * Y
         axs[1].contour(X, Y, Z, levels=[0], colors='black')
         # fill the area with different colors based on the contour
         axs[1].contourf(X, Y, Z, levels=[np.min(Z), 0, np.max(Z)], colors=['red', 't
         axs[1].set xlim(0.3, 1)
         axs[1].set_xlabel('Average Intensity')
         axs[1].set ylim(-0.05, 0.55)
         axs[1].set_ylabel('Symmetry')
```

```
axs[1].legend()
axs[1].set_title('Polynomial Transformation')
```

Out[17]: Text(0.5, 1.0, 'Polynomial Transformation')



Q2.

Fit auto-regressive (AR) models with regular and stochastic gradient de- scents. Refer to the two-tap predictor example in the slides of chapter 5a. Consider an AR(2) model:

$$x(n)=-w_1x(n-1)-w_2x(n-2)+\epsilon(n),$$

where x(n) is a time series data, x(n-1) and x(n-2) are the lag 1 and lag 2 series of x(n), and $\epsilon(n)$ is a Gaussian noise with zero mean.

```
In [18]: import numpy as np
         import matplotlib.pyplot as plt
         import os
In [19]: # read data
         base_dir = "/Users/chuan/Desktop/c815_course_hws/hw4/data_hw4"
         path_data1 = os.path.join(base_dir, "data1.csv")
         path_data2 = os.path.join(base_dir, "data2.csv")
In [20]: data1 = np.genfromtxt(path_data1, delimiter=',')
         data2 = np.genfromtxt(path_data2, delimiter=',')
         data1.shape, data2.shape
Out[20]: ((1000,), (1000,))
In [21]: # create lag 1 and lag 2 data for both datasets, and set the first row to 0
         data1_lag1 = np.zeros(data1.shape)
         data1_lag1[1:] = data1[:-1]
         data1_lag2 = np.zeros(data1.shape)
         data1_{lag2[2:]} = data1[:-2]
         data2_lag1 = np.zeros(data2.shape)
         data2_{lag1[1:]} = data2[:-1]
```

```
data2_lag2 = np.zeros(data2.shape)
data2_lag2[2:] = data2[:-2]
```

```
In [22]: class GD:
             def __init__(self, learning_rate=0.1, max_iter=10000, stop_criteria=1e-6
                 self.learning_rate = learning_rate
                  self.max iter = max iter
                  self.stop_criteria = stop_criteria
                  self.weights = None
                  self.bias = None
                  self.training errors = []
                  self.early_stop = early_stop
             # Standard Gradient Descent
             def original gradient(self, X, Y):
                 # initialize weights
                 self.weights = np.zeros(X.shape[1])
                 m = X.shape[0] # Number of samples
                 for i in range(self.max_iter):
                     # compute predictions
                     predictions = X.dot(self.weights)
                     # calculate the gradient
                     gradient = (2 / m) * (X.T.dot(predictions - Y))
                     # update weights
                     self.weights -= self.learning_rate * gradient/2
                     # calculate the training error
                     training_error = np.mean((Y - predictions) ** 2)
                     self.training errors.append(training error)
                     # check for convergence
                     if self.early stop:
                          if np.linalq.norm(gradient) < self.stop criteria:</pre>
                              print(f"Converged at iteration {i}")
                              break
                  return self.weights
             # Stochastic Gradient Descent
             def stochastic_gradient(self, X, Y):
                 # initialize weights
                  self.weights = np.zeros(X.shape[1])
                 m = X.shape[0] # number of samples
                 for i in range(self.max_iter):
                     #for j in range(m):
                     j = np.random.randint(0, m)
                     # compute predictions
                     prediction = X[j].dot(self.weights)
                     # calculate the gradient
                     gradient = 2 * X[j] * (prediction - Y[j])
                     # update weights
```

```
self.weights -= self.learning_rate * gradient

# calculate the training error
training_error = np.mean((Y - X.dot(self.weights)) ** 2)
self.training_errors.append(training_error)

# check for convergence
if self.early_stop:
    if np.linalg.norm(gradient) < self.stop_criteria:
        print(f"Converged at iteration {i}")
        break
return self.weights</pre>
```

(a)

Estimate w1 and w2 for the two datasets using regular gradient descent. Write the final values for w1 and w2 for each dataset.

```
In [23]: original_gd = GD()
         # fit the model, where X is the lag 1 and lag 2 data, and one bias term, and
         X = np.vstack((np.ones(data1.shape), data1 lag1, data1 lag2)).T
         Y = data1
         weights = original qd.original gradient(X, Y)
         # print w1, w2, not w0
         print("w1:", weights[1])
         print("w2:", weights[2])
        Converged at iteration 3187
        w1: -0.4459449377412381
        w2: -0.15594109730889713
In [24]: # also fit for data2
         X = np.vstack((np.ones(data2.shape), data2 lag1, data2 lag2)).T
         Y = data2
         weights = original_gd.original_gradient(X, Y)
         print("w1:", weights[1])
         print("w2:", weights[2])
        Converged at iteration 1146
        w1: -0.11396238532127233
        w2: -0.8018278712897495
```

(b)

Estimate w1 and w2 for the two datasets using stochastic gradient descent, and compare the result with part (a). Write the final values for w1 and w2 for each dataset.

```
In [25]: # for data1
    stochestic_gd = GD()
    X = np.vstack((np.ones(data1.shape), data1_lag1, data1_lag2)).T
    Y = data1
    weights = stochestic_gd.stochastic_gradient(X, Y)
```

```
print("w1:", weights[1])
print("w2:", weights[2])

w1: -0.4163773715798901
w2: -0.21296943798480322

In [26]: # for data2
    stochestic_gd = GD()
    X = np.vstack((np.ones(data2.shape), data2_lag1, data2_lag2)).T
    Y = data2
    weights = stochestic_gd.stochastic_gradient(X, Y)
    print("w1:", weights[1])
    print("w2:", weights[2])

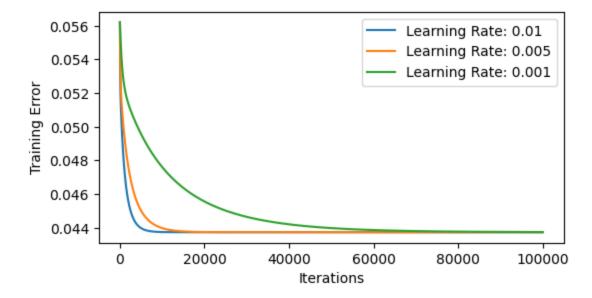
w1: -0.2324134501136457
w2: -0.7805733656040625
```

(c)

Plot the training error as a function of the iteration step using 3 different learning rates on dataset 1 while employing gradient descent

```
In [27]: learning_rates = [0.01, 0.005,0.001]
         \max iter = 100000
In [28]: # for data1
         X = np.vstack((np.ones(data1.shape), data1 lag1, data1 lag2)).T
         Y = data1
         learning errors = []
         for lr in learning rates:
             gd = GD(learning_rate=lr, max_iter=max_iter, early_stop=False)
             weights = gd.original_gradient(X, Y)
             learning errors.append(gd.training errors)
         # plot the learning curve, if stopped early, fill the curve with the last va
         max_len = max([len(errors) for errors in learning_errors])
         plt.figure(figsize=(6, 3))
         for i, lr in enumerate(learning_rates):
             if len(learning_errors[i]) < max_len:</pre>
                 learning errors[i] += [learning errors[i][-1]] * (max len - len(lear
             plt.plot(range(len(learning_errors[i])), learning_errors[i], label=f"Led
         plt.xlabel('Iterations')
         plt.ylabel('Training Error')
         plt.legend()
```

Out[28]: <matplotlib.legend.Legend at 0x11af5fc50>

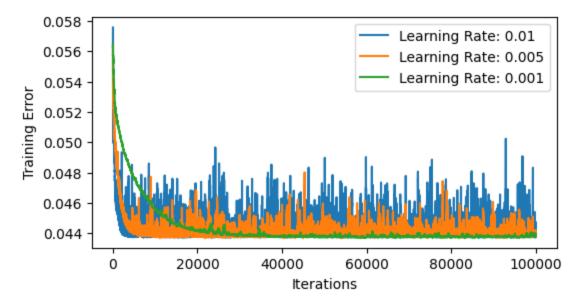


(d)

Plot that training error as a function of the iteration step using 3 different learning rates on dataset 1 while employing stochastic gradient descent.

```
In [29]: # for data1
         learning errors = []
         for lr in learning_rates:
             gd = GD(learning rate=lr, max iter=max iter, early stop=False)
             weights = gd.stochastic_gradient(X, Y)
             learning_errors.append(gd.training_errors)
         # plot the learning curve, if stopped early, fill the curve with the last va
         max_len = max([len(errors) for errors in learning_errors])
         plt.figure(figsize=(6, 3))
         for i, lr in enumerate(learning_rates):
             if len(learning_errors[i]) < max_len:</pre>
                 learning_errors[i] += [learning_errors[i][-1]] * (max_len - len(lear
             plt.plot(range(len(learning_errors[i])), learning_errors[i], label=f"Lea
         plt.xlabel('Iterations')
         plt.ylabel('Training Error')
         plt.legend()
```

Out[29]: <matplotlib.legend.Legend at 0x1080cfdd0>



Explain

Through comparison, it is found that unlike the regular gradient descent, the stochastic gradient descent (SGD) shows more fluctuation in terms of error trends, while both of the graphs show certain level of convergence, implying a successful learning.

The reason for this noisy error interpretation for SGD implies is that SGD only update an individual point for each iteration, which will bring larger violitility because not every sample will be a perfect representation for the learning.