ECE421: Introduction to Machine Learning Programming Assignment 1

Assigned: Jan 16, 2023; Due: Feb 3, 2023 @ 11:59 p.m.

Objectives

In this assignment, you will be implementing the following algorithms: Pocket Algorithm, and Linear Regression; using two different methods of coding approaches. In the first approach, you will implement these algorithms using Python and functions in the NumPy library only. In the second approach, you will use scikit-learn to gauge how well your initial implementation using NumPy functions fares in comparison to off-the-shelf modules available in scikit-learn. You will also be asked to answer several questions related to your implementations. To avoid any potential installation issue, you are encouraged to develop your solution using Google Colab notebooks.

Requirements

In your implementations, please use the function prototype provided (i.e. name of the function, inputs and outputs) in the detailed instructions presented in the remainder of this document. We will be testing your code using a test function that which evokes the provided function prototype. If our testing file is unable to recognize the function prototype you have implemented, you can lose significant portion of your marks. In the assignment folder, the following files are included in the starter_code folder:

- PerceptronImp.py
- LinearRegressionImp.py

These files contain the test function and an outline of the functions that you will be implementing. You also need to submit a separate PAl_qa.pdf file that answer questions related to your implementations.

1 Pocket Algorithm

In the PerceptronImp.py file, you will implement the Pocket Algorithm to classify two different classes in the IRIS dataset. You do not need to download the dataset as it is included in the scikit-learn library. For evaluation, we provide you the test function test_Part1() (Note: keep this function as it is in your submission). This function loads the IRIS dataset, runs your implementation and outputs the confusion matrix on the test set (for each coding approach). Follow the below instructions to get started.

Part 1a: Pocket Algorithm using NumPy and Python

You will be implementing the Pocket Algorithm using the NumPy library functions in the PerceptronImp.py file. You will be computing parameters of a linear plane that best separates input features belonging to two classes. Specifically, you will be implementing four functions which are detailed in the following:

- Function 1: def fit_Perceptron(X_train, y_train)
 - Inputs: X_train, y_train

 The first input X_train represents the matrix of input features that belongs to \mathbb{R}^{NXd} where N is the total number of training samples and d is the dimension of each input feature vector.

 The second input y_train is an N dimensional vector where the i^{th} component represents the output observed in the training set for the i^{th} row in X_train matrix which corresponds to the i^{th} input feature. Each element in y_train takes the value +1 or -1 to represent the first class and second class respectively.

- Output: w

The output of this function is the vector w that represents the coefficients of the line computed by the pocket algorithm that best separates the two classes of training data points. The dimensions of this vector is d+1 as the offset term is accounted in the computation.

- Function implementation considerations:

This function computes the parameters w of a linear plane which separates the input features from the training set into two classes specified by the training dataset. As the pocket algorithm is used, you will set the maximum number of epochs (the maximum number of passes over the training data) to 5000. Useful functions in NumPy for implementing this function are: zeros (for initializing the weight vector with 0s), shape (for identifying the number of rows and columns in a matrix), hstack (to add an additional column to the front of the original input matrix), ones (for setting the first column of the input feature matrix to 1), dot function to take the dot product of two vectors of the same size. You will also use the function errorPer that you will implement next to compute the average number of misclassifications for the plane you are currently considering with respect to the training dataset.

- Function 2: def errorPer(X_train, y_train, w)
 - Inputs: X_train, y_train and w

The inputs to this function are X_train and y_train. X_train is defined with the additional column of ones and y_train is defined in a manner similar to Function 1. w represents the coefficients of a linear plane and is of d+1 dimensions.

- Output: avgError

The output of this function is the average number of points that are misclassified by the plane defined by w.

Function implementation considerations:
 You will use the pred function that you will implement in Function 3 to find the output of the classifier defined by w.

- Function 3: def pred(X_i,w)
 - Inputs: X_i, w

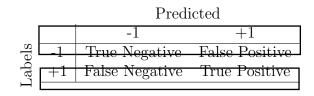
The first input is X_i which is the feature vector of d+1 dimensions of the i^{th} test datapoint. w is defined in a manner similar to Function 2.

- Output: Class label

The class predicted for linear classifier defined by w for the input datapoint X_i is computed. This output can take one of the following values: +1 and -1.

- Function implementation considerations:
 - If the dot product of the vector w and the input point X_{-i} is **strictly** positive, then you can consider the input point to map to a point that is above the line (i.e. belongs to class 1). Otherwise, it will belong to class -1.
- Function 4: def confMatrix(X_train,y_train,w)
 - Inputs: X_train, y_train and w
 These inputs are defined in the same manner as Function 1.
 - Output: A two-by-two matrix composed of integer values
 - Function implementation considerations:

This function will populate a two-by-two matrix. Using the zero-index, the (0,0) position represents a count of the total number of points correctly classified to be class -1 (True Negative). The (0,1) position contains a count of total number of points that are in class -1 but are classified to be class +1 by the classifier (False Positive). The (1,0) position contains a count of total number of points that are in class +1 but are classified to be class -1 by the classifier (False Negative). The (1,1) position represents a count of the total number of points correctly classified to be class +1 (True Positive). Refer to the table below.



The following is the mark breakdown for Part 1a:

- Test file successfully runs all four implemented functions: 8 marks
- Outputs of all four functions are close to the expected output: 12 marks
- Code content is organized well and annotated with useful comments: 10 marks

Part 1b: Pocket Algorithm using scikit-learn

In this part, you will use the scikit-learn library to train the binary linear classification model. You will then compare the performance of your implementation in Part 1a with the one available in the scikit-learn library. You will implement one function in this part in the PerceptronImp.py file. You can refer to the scikit-learn demo covered in the lecture to aid you with this completing this part or refer to this page: https://scikit-learn.org/stable/modules/generated/sklearn.linear_model.Perceptron.html.

- Function: def test_SciKit(X_train, X_test, Y_train, Y_test)
 - Inputs: X_train, X_test, Y_train, Y_test

 The first input X_train represents the matrix of input features that belongs to \mathbb{R}^{NXd} where N is the total number of training samples and d is the dimension of each input feature vector. The second input X_test represents the matrix of input features that belongs to \mathbb{R}^{MXd} where M is the total number of testing samples and d is the dimension of each input feature vector. The third input Y_train is an N dimensional vector where the i^{th} component represents the output observed in the training set for the i^{th} row in X_train matrix which corresponds to the i^{th} input feature. The similar counterpart to X_test is Y_test.
 - Output: A two-by-two matrix composed of integer values
 This function will output the result obtained from the confusion matrix function imported from sklearn.metrics library to report the performance of the model fitted using the Perceptron algorithm available in the sklearn.metrics library.
 - Function implementation considerations:

 Perceptron and confusion_matrix functions imported from sklearn.linear_model,
 sklearn.metrics will be utilized to fit the linear classifer using the Perceptron learning
 algorithm and evaluate the performance of this algorithm. As presented in the scikit-learn
 demo in the lecture, you will initiate an object of the Perceptron type, you will run the fit
 function to train the classifier, you will use the predict function to perform predictions using
 the trained algorithm and finally you will use the confusion_matrix function to identify how
 many points have been classified correctly and incorrectly in the test dataset. Refer to the
 below questions in order to set the parameters for your Perceptron model.

Answer the following question(s), write and save your answer in a separate PA1_qa.pdf file. Remember to submit this file together with your code.

- 1. Refer to the documentation, what is the functionality of the tol parameter in the Perceptron class? (2 marks)
- 2. If we set max_iter=5000 and tol=1e-3 (the rest as default), does this guarantee that the algorithm will pass over the training data 5000 times? If not, which parameters (and values) should we set to ensure that the algorithm will pass over the training data 5000 times? (2 marks)
- 3. How can we set the weights of the model to a certain value? (2 marks)

4. How close is the performance (through confusion matrix) of your NumPy implementation in comparison to the existing modules in the scikit-learn library? (2 marks)

The following is the mark breakdown for Part 1b:

- Test file successfully runs implemented function: 4 marks
- Output is close to the expected output from the test file: 5 marks
- Code content is organized well and annotated with comments: 3 marks
- Questions are answered correctly: 8 marks

2 Linear Regression

In the LinearRegressionImp.py file, you will implement the Linear Regression algorithm and test its performance using the diabetes dataset. You do not need to download the dataset as it is included in the scikit-learn library. For evaluation, we provide you the test function test_Part2()(Note: keep this function as it is in your submission). This function loads the diabetes dataset, runs your implementation and outputs the mean-squared-error on the test set (for each coding approach). Follow the below instructions to get started.

Part 2a: Linear Regression using NumPy and Python

You will be implementing in LinearRegressionImp.py the exact computation of the solution for linear regression using the NumPy library functions via the least squares method. You will be computing the parameters of a linear plane that best fits the training dataset. Specifically, you will be implementing three functions which are detailed in the following:

- Function 1: def fit_LinRegr(X_train, y_train)
 - Inputs: X_train, y_train

 The first input X_train represents the matrix of input features that belongs to \mathbb{R}^{NXd} where N is the total number of training samples and d is the dimension of each input feature vector. The second input y_train is an N dimensional vector where the i^{th} component represents the output observed in the training set for the i^{th} row in X_train matrix which corresponds to the i^{th} input feature. Each element in y_train takes a value in \mathbb{R} .
 - Output: w The output of this function is the vector w that represents the coefficients of the line computed using the least square method that best fits the training data points. The dimensions of this vector is d+1 as the offset term is accounted in the computation.
 - Function implementation considerations:

 This function computes the parameters w of a linear plane which best fits the training dataset. Useful functions in NumPy for implementing this function are: shape (for identifying the number of rows and columns in a matrix), hstack (to add an additional column to the front of the original input matrix), ones (for setting the first column of the input feature matrix to 1), dot function to take the dot product of two vectors of the same size, transpose function for taking the transpose of a vector, and linalq.inv function for finding the inverse of a square matrix.
- Function 2: def mse(X, y, w)
 - Inputs: X, y and w
 The inputs to this function, X and y, are defined in a manner similar to X_train and y_train in Function 1. w represents the coefficients of a linear plane and is of d+1 dimensions.

- Output: avgError
 The output of this function is the mean squared error introduced by the linear plane defined by w.
- Function implementation considerations:
 You will use the pred function that you will implement in Function 3 to find the output of the linear plane defined by w. Functions from NumPy that will be useful are: shape (for identifying the number of rows and columns in a matrix), hstack (to add an additional column to the front of the original input matrix), ones (for setting the first column of the input feature matrix to 1), and dot function to take the dot product of two vectors of the same size.
- Function 3: def pred(X_i,w)
 - Inputs: X_i , w

 The first input is X_i which is the feature vector of d+1 dimensions of the i^{th} test datapoint. w is defined in a manner similar to Function 2.
 - Output: Predicted value The output predicted by the linear regression model defined by w for the input datapoint X_i is computed. This output can take values in the Real space \mathbb{R} .
 - Function implementation considerations:
 The dot product function in NumPy will be useful for this function implementation.

For this implementation, we also provide the test function subtestFn(). This function loads a toy dataset, runs your NumPy implementation and return a message indicating whether your solution works when X_train is not a full-column rank matrix, i.e. the input features are not linearly independent.

Answer the following question(s), write and save your answer in a separate PA1_qa.pdf file. Remember to submit this file together with your code.

- When we input a singular matrix, the function linalg.inv often returns an error message. In your fit_LinRegr(X_train, y_train) implementation, is your input to the function linalg.inv a singular matrix? Explain why. (2 marks)
- As you are using linalg.inv for matrix inversion, report the output message when running the function subtestFn(). We note that inputting a singular matrix to linalg.inv sometimes does not yield an error due to numerical issue. (1 marks)
- Replace the function linalg.inv with linalg.pinv, you should get the model's weight and the "NO ERROR" message after running the function subtestFn(). Explain the difference between linalg.inv and linalg.pinv, and report the model's weight. (2 marks)

The following is the mark breakdown for Part 2a:

- Test file successfully runs all three implemented functions: 8 marks
- Outputs of all four functions are close to the expected output: 12 marks
- Code content is organized well and annotated with useful comments: 5 marks
- Questions are answered correctly: 5 marks

Part 2b: Linear Regressions using scikit-learn

In this part, you will use the scikit-learn library to train the linear regression model. You will then compare the performance of your implementation in Part 2a with the one available in the scikit-learn library. You will implement one function in this part in the LinearRegressionImp.py file. You can refer to the scikit-learn demo covered in the lecture to aid you with this completing this part.

• Function: def test_SciKit(X_train, X_test, Y_train, Y_test)

- Inputs: X_train, X_test, Y_train, Y_test

 The first input X_train represents the matrix of input features that belongs to \mathbb{R}^{NXd} where N is the total number of training samples and d is the dimension of each input feature vector. The second input X_test represents the matrix of input features that belongs to \mathbb{R}^{MXd} where M is the total number of testing samples and d is the dimension of each input feature vector. The third input Y_train is an N dimensional vector where the i^{th} component represents the output observed in the training set for the i^{th} row in X_train matrix which corresponds to the i^{th} input feature. The similar counterpart to X_test is Y_test.
- Output: error This function will output the mean squared error on the test set, which is obtained from the mean_squared_error function imported from sklearn.metrics library to report the performance of the model fitted using the linear regression algorithm available in the sklearn.metrics library.
- Function implementation considerations:
 LinearRegression and mean_squared_error functions imported from sklearn.linear_model,
 sklearn.metrics will be utilized to fit the linear classifier using the linear regression algorithm and evaluate the performance of this algorithm. As presented in the scikit-learn
 demo in the lecture, you will initiate an object of the LinearRegression type, you will run
 the fit function to train the model, you will use the predict function to perform predictions
 using the trained algorithm and finally you will use the mean_squared_error function to
 compute the mean squared error of the trained model.

How close is the performance of your implementation in comparison to the existing modules in the scikit-learn library? Place this comment at the end of the code file.

The following is the mark breakdown for Part 2b:

- Test file successfully runs implemented function: 6 marks
- Output is close to the expected output from the test file: 8 marks
- Code content is organized well and annotated with comments: 6 marks