Module 1: Linear Regression

To write legible answers you will need to be familiar with both Markdown and Latex

Before you turn this problem in, make sure everything runs as expected. First, restart the kernel (in the menubar, select Kernel $\rightarrow \rightarrow$ Restart) and then run all cells (in the menubar, select Cell $\rightarrow \rightarrow$ Run All).

Make sure you fill in any place that says "YOUR CODE HERE" or "YOUR ANSWER HERE", as well as your name below:

```
In [ ]: NAME = ""
STUDENT_ID = ""
```

Question 1 - Linear Regression

In this question, you will be implementing the linear regression algorithm from scratch in Python. Linear regression aims to map feature vectors to a continuous value in the range $[-\infty, +\infty]$ by linearly combining the feature values.

Model Representation

Data is represented as a dataframe or a feature matrix. Let our feature matrix be X whose dimensions are $n \times m$, θ be a weight matrix of dimensions $m \times 1$, the bias vector b a column vector of dimension $m \times 1$. Using these we can predict \hat{Y} by the following relationship:

$$\hat{Y} = X\theta + b$$

Data: Facebook posts metrics

This data contains features describing posts from a cosmetic brand's Facebook page. The authors use the following features:

- · Category,
- Page total likes: Number of people who have liked the company's page),
- Type: Type of content (Link, Photo, Status, Video),
- Post month: Month the post was published (January, February, March, ..., December),
- Post hour: Hour the post was published (0, 1, 2, 3, 4, ..., 23)
- Post weekday: Weekday the post was published (Sunday, Monday, ...,

Saturday),

Paid: If the company paid to Facebook for advertising (yes, no)

to model:

'Lifetime Post Total Reach', 'Lifetime Post Total Impressions', 'Lifetime Engaged Users', 'Lifetime Post Consumers', 'Lifetime Post Impressions by people who have liked your Page', 'Lifetime Post reach by people who like your Page', 'Lifetime People who have liked your Page and engaged with your post', 'comment', 'like', 'share', 'Total Interactions'.

There are many possible features we could try to model, but we will focus on 'Total Interactions'. Our feature space will include: Category, Page total likes, Post month, Post hour, Post weekday, and Paid. We drop "Type" simply to avoid preprocessing.

You can read more about the dataset here.

Downloading the data

```
In [ ]: !wget http://archive.ics.uci.edu/ml/machine-learning-databases/00368/Facebook_metri
import zipfile
with zipfile.ZipFile('./Facebook_metrics.zip', 'r') as zip_ref:
    zip_ref.extractall('./')
```

Reading in data

```
In [ ]:
        import pandas as pd
        import numpy as np
        np.random.seed(144)
        Shuffles the data in place
        def shuffle_data(data):
            np.random.shuffle(data)
        # Read in the data
        lr_dataframe = pd.read_csv('dataset_Facebook.csv',sep=';')
        lr_dataframe.dropna(inplace=True)
        columns_to_drop = ['Type','Lifetime Post Total Reach', 'Lifetime Post Total Impress
               'Lifetime Engaged Users', 'Lifetime Post Consumers',
               'Lifetime Post Consumptions',
               'Lifetime Post Impressions by people who have liked your Page',
               'Lifetime Post reach by people who like your Page',
               'Lifetime People who have liked your Page and engaged with your post',
               'comment', 'like', 'share']
        lr dataframe.drop(columns=columns to drop,inplace=True)
        # Normalizing all remaining columns
        def normalize_col(col):
            return (col - col.min())/(col.max() - col.min())
        lr_dataframe = lr_dataframe.apply(normalize_col)
        # Get entries as a numpy array
        lr_data = lr_dataframe.values[:, :]
        # Shuffle once for reproducibility
        shuffle_data(lr_data)
        lr_dataframe.head()
```

a) Splitting data in X and Y

In this part we will write functions to split our data into a feature matrix X (augmented by a column of 1's to account for the bias term) and a column vector we wish to predict Y and further split X and Y into X_train, X_test, y_train and y_test for training and testing.

- i) Split the dataset into X and Y. In order to vectorize our calculations, we utilize the bias trick. This simply means we need to append ones to our X matrix.
- ii) Split X and Y into the training and test sets using the provided percentage split (default is 80% training and 20% test).

```
0.00
In [ ]:
        Combines one column of all ones and the matrix X to account for the bias term
        (setting x_0 = 1) - [Hint: you may want to use np.hstack()]
        Takes input matrix X
        Returns the augmented input
        1.1.1
        def bias_trick(X):
            # YOUR CODE HERE
        Separates feature vectors and targets
        Takes raw data
        Returns X as the matrix of feature vectors and Y as the vector of targets
        def separate_data(data):
            # Split into X (remember to use bias trick) and Y
            # YOUR CODE HERE
        Takes raw data in and splits the data into
        X_train, y_train, X_test, y_test
        Returns X_train, y_train, X_test, y_test
        def train_test_split(data, train_size=.80):
            # YOUR CODE HERE
            return
```

b) Training and testing our model

Refer to the following derivation of the gradient when implementing linear regression and gradient descent below.

For this question, we'll use and implement the Mean Squared Error (MSE), and gradient descent algorithm. Suppose our dataset consists of n records, each with d features:

$$X = egin{bmatrix} x_{1,1} & x_{1,2} & \cdots & x_{1,d} \ x_{2,1} & x_{2,2} & \cdots & x_{2,d} \ dots & dots & \ddots & dots \ x_{n,1} & x_{n,2} & \cdots & x_{n,d} \end{bmatrix}$$

One way to include a bias is to augment X with a column of ones:

$$X = egin{bmatrix} 1 & x_{1,1} & x_{1,2} & \cdots & x_{1,d} \ 1 & x_{2,1} & x_{2,2} & \cdots & x_{2,d} \ dots & dots & dots & \ddots & dots \ 1 & x_{n,1} & x_{n,2} & \cdots & x_{n,d} \end{bmatrix}$$

We also have n labels corresponding to the correct classification of each of the above records, $y=[y_1,y_2,\cdots,y_n]^T$, i.e.:

$$y = \left[egin{array}{c} y_1 \ y_2 \ dots \ y_n \end{array}
ight]$$

We will try to find the optimal parameter values $\theta = [\theta_0, \theta_1, \cdots, \theta_d]^T$ of our linear regression model, where θ_0 is the bias weight. To simplify our notation, let

$$\hat{y}=X heta=egin{bmatrix} X_{1,0} heta_0+X_{1,1} heta_1+\cdots+X_{1,d} heta_d\ X_{2,0} heta_0+X_{2,1} heta_1+\cdots+X_{2,d} heta_d\ dots\ X_{n,0} heta_0+X_{n,1} heta_1+\cdots+X_{n,d} heta_d \end{bmatrix}=egin{bmatrix} \hat{y}_1\ \hat{y}_2\ dots\ \hat{y}_n \end{bmatrix}$$

We seek θ such that the MSE is minimized (the 1/2 factor makes the derivation easier). Let the MSE be a function of θ , $J(\theta)$:

$$J(heta) = rac{1}{2n} \sum_{i=1}^n (\hat{y}_i - y_i)^2$$

Since the above is a convex function, it has a unique minimum value. Taking the derivative with respect to θ_i , we get:

$$egin{aligned} rac{\partial}{\partial heta_j} J(heta) &= rac{1}{2n} \sum_{i=1}^n rac{\partial}{\partial heta_j} (\hat{y}_i - y_i)^2 \ &= rac{1}{n} \sum_{i=1}^n (\hat{y}_i - y_i) rac{\partial}{\partial heta_j} (\hat{y}_i) \end{aligned}$$

Recall the chain rule from calculus, and that each \hat{y}_i is a function of the θ_i , so the above becomes:

$$rac{\partial}{\partial heta_j} J(heta) = rac{1}{n} \sum_{i=1}^n (\hat{y_i} - y_i) x_{i,j}$$

YOU SHOULD:

- i) Get training and testing set by calling train_test_split()
- ii) Define a weight (θ) vector
- ii) Implement Gradient Descent using the information above
- iii) Record the Sum Squared Error for training and test data
- iv) Return the weight matrix, train errors and test errors
- v) Plot the training and test errors and comment on the plot.

```
In [ ]:
        Takes the target values and predicted values and calculates the squared error
        between them
        def mse(y_pred, y_true):
            # YOUR CODE HERE
            return
        Implementation of the derivative of MSE.
        Returns a vector of derivations of loss with respect to each of the dimensions
        [\partial loss / \partial \theta_j]
        def mse_derivative(X,y,theta):
            # YOUR CODE HERE
            return
        Gradient descent step.
        Takes X, y, theta vector, and alpha.
        Returns an updated theta vector.
        def gradient_descent_step(X,y, theta, alpha):
            # YOUR CODE HERE
            return theta
```

```
In [ ]: # Carry out training task
# YOUR CODE HERE
        # Plot the training error and test error for different epochs (iterations of the
        # algorithm). Your plot be MSE error vs epochs.
        # YOUR CODE HERE
             cesc_errors- []
             # Define theta
             theta = np.zeros((X_train.shape[1]))
             # Carry out training loop
             for i in range(num epochs):
                 train_error = # YOUR CODE HERE
                 train_errors.append(train_error)
                 test error = # YOUR CODE HERE
                 test_errors.append(test_error)
                 # Do gradient descent on the training set
                 theta = # YOUR CODE HERE
             return theta, train_errors, test_errors
```

Data may not follow a linear relationship from the independent variable X to the dependent variable y. Fitting a linear model to this would be inaccurate and yield a high loss.

If we want to model an order d polynomial relationship between X and y we can augment our initial linear model where instead of having:

$$y_i = \theta_0 + \theta_1 x_i$$

We have:

$$y_i = heta_0 + heta_1 x_i + heta_2 x_i^2 + \dots + heta_d x_i^d$$

We can use the same linear regression algorithm we if we first augment X and add extra columns (or dimensions).

$$\mathbf{X} = egin{bmatrix} x_1 & x_1^2 & \cdots & x_1^d \ x_2 & x_2^2 & \cdots & x_2^d \ dots & dots & \ddots & dots \ x_n & x_n^2 & \cdots & x_n^d \end{bmatrix}$$

Then our new higher order \hat{y} is computed same as before.

$$\hat{y} = X\theta = \begin{bmatrix} 1 & x_1 & x_1^2 & \cdots & x_1^d \\ 1 & x_2 & x_2^2 & \cdots & x_2^d \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 1 & x_n & x_n^2 & \cdots & x_n^d \end{bmatrix} \begin{bmatrix} \theta_0 \\ \theta_1 \\ \vdots \\ \theta_d \end{bmatrix} = \begin{bmatrix} \theta_0 + \theta_1 x_1 + \theta_2 x_1^2 + \cdots + \theta_d x_1^d \\ \theta_0 + \theta_1 x_2 + \theta_2 x_2^2 + \cdots + \theta_d x_2^d \\ \vdots \\ \theta_0 + \theta_1 x_n + \theta_2 x_n^2 + \cdots + \theta_d x_n^d \end{bmatrix} = \begin{bmatrix} \hat{y}_1 \\ \hat{y}_2 \\ \vdots \\ \hat{y}_n \end{bmatrix}$$

```
In [ ]:
                      import numpy as np
                      import matplotlib.pyplot as plt
                      def normalize data(data):
                                 return (data - np.min(data))/(np.max(data) - np.min(data))
                      np.random.seed(33)
                      x = np.random.uniform(-10, 10, 1000)
                      poly_coeffs = np.random.uniform(-1,1, size=(4,1))
                      y = poly\_coeffs[0] + poly\_coeffs[1]*x + poly\_coeffs[2]*(x ** 2) + poly\_coeffs[3]*(x ** 2) + po
                      x2 = np.random.uniform(-10, 10, 1000)
                      poly_coeffs = np.random.uniform(-1,1, size=(3,1))
                      y2 = poly\_coeffs[0] - 2000 + poly\_coeffs[1]*x2 + 50*poly\_coeffs[2]*(x2 ** 2) + np.
                      x = np.concatenate([x,x2])
                      y = np.concatenate([y,y2])
                      x = normalize_data(x)
                      y = normalize_data(y)
                      plt.scatter(x,y, s=10)
                      plt.show()
                      poly_data = np.hstack((x.reshape(-1,1),y.reshape(-1,1)))
                      np.random.shuffle(poly_data)
                      x = poly_data[:,0]
                      y = poly_data[:,1]
In [ ]: import numpy as np
                      from sklearn.linear_model import LinearRegression
In [ ]: reg = LinearRegression().fit(x.reshape(-1,1), y)
In [ ]: | def compute_line_from_regr(X_data, y_data, regr):
                                 l_bound = np.min(X_data)
                                 r_bound = np.max(X_data)
                                 return [1_bound, r_bound], [1_bound * regr.coef_ + regr.intercept_, r_bound * r
                      plt.scatter(x,y, s=10)
                      line_x, line_y = compute_line_from_regr(x.reshape(-1,1),y,reg)
                      plt.plot(line_x, line_y, color='r')
                      plt.show()
```

As we see above, this data doesn't follow a linear relationship, it follows some complex polynomial. In the next section you'll try to fit a higher degree polynomial to it.

Weight regularization

When we try to fit a d-order polynomial to our data, we could end up overfitting. This happens when you try to fit a higher dimensional curve than what the distribution of our data actually exhibits. We can mitigate this by choosing an order d that matches your data closely, but often times this is not directly apparant in noisy data. Another method to avoid overfitting is $\mathbf{regularizing}$, where you modify your loss to keep weights small which flattens our polynomial. This helps us avoid learning polynomials that are too complex for our data.

To add regularization we modify our original loss function J to include our regularizing term and a new hyperparameter that we tune λ . This controls the amount of regularizing we impose on the weights. We use the loss computed from the validation set to tweak this parameter.

$$J(heta) = rac{1}{2n} \sum_{i=1}^n (h^{(i)} - y^{(i)})^2 + \lambda \sum_{j=1}^d heta_j^2$$

Our gradient computation also changes:

$$rac{\partial}{\partial heta_j} J(heta) = rac{1}{n} \sum_{i=1}^n (h^{(i)} - y^{(i)}) x_{i,j} + 2 \lambda heta_j$$

We apply this gradient the same way as before in our gradient descent algorithm:

$$heta_j = heta_j - lpha rac{\partial}{\partial heta_j} J(heta)$$

```
1.1.1
In [ ]:
        Adds columns to your data up to the specified degree.
        Ex: If degree=3, (x) \rightarrow (x, x^2, x^3)
        def add_polycols(X,degree):
            x_{col} = X[:,-1]
            for i in range(2, degree+1):
                 X = np.hstack((X,(x_col**i).reshape(-1,1)))
             return X
        Takes the target values and predicted values and calculates the absolute error
        between them
        def mse(y_pred, y_true):
            # YOUR CODE HERE
            # Feel free to use your implementation from Q1
            return
         111
        Implementation of the derivative of MSE.
        Returns a vector of derivations of loss with respect to each of the dimensions
         [\partial loss / \partial \theta_i]
        def mse derivative(X,y,theta):
            # YOUR CODE HERE
            # Feel free to use your implementation from Q1
            return
         . . .
        Computes L2 norm from theta scaled by lambda.
        Returns a scalar L2 norm.
        def 12norm(theta, lamb):
            # YOUR CODE HERE
            return
         . . .
        Computes derivative of L2 norm scaled by lambda.
        Returns a vector of derivative of L2 norms.
        def 12norm_derivative(theta, lamb):
            # YOUR CODE HERE
            # Note there is no regularization on the bias term.
            return
        Computes total cost (cost function + regularization term)
        def compute_cost(X, y, theta, lamb):
            # YOUR CODE HERE
            return
        Gradient descent step.
```

```
Takes X, y, theta vector, and alpha.
Returns an updated theta vector.
def gradient_descent_step(X, y, theta, alpha, lamb):
    # YOUR CODE HERE
    # This differs from your Q1 implementation
    return
def polynomial_regression(data, degree, num_epochs=100000, alpha=1e-4, lamb=0):
    # Get training, testing, and validation sets by calling train test validation s
    # YOUR CODE HERE
    # Record training and validation errors in lists
    train_errors = []
    val_errors = []
   # Add the appropriate amount of columns to each of your sets of data.
   X_train = add_polycols(X_train, degree)
    X val = add polycols(X val, degree)
   X_test = add_polycols(X_test, degree)
    # Define theta
    theta = np.zeros((X_train.shape[1]))
    # Carry out training loop
    for i in range(num_epochs):
        train_error = # YOUR CODE HERE
        train_errors.append(train_error)
        val error = # YOUR CODE HERE
        val_errors.append(val_error)
        # Do gradient descent on the training set
        theta = # YOUR CODE HERE
        # This prints the validation loss
        if i % (num_epochs//10) == 0:
            print(f'({i} epochs) Training loss: {train error}, Validation loss: {va
    print(f'({i} epochs) Final training loss: {train_error}, Final validation loss:
    # Compute the testing loss
    test error = # YOUR CODE HERE
    print(f'Final testing loss: {test_error}')
    return theta, train_errors, val_errors
```

As we mentioned above, we use the validation set's loss to tweak our hyperparameters. Please carry out the training task while monitoring the validation loss and varying the polynomial order d and regularization constant λ . Your answer should get close to minimizing the validation and testing losses.

```
In []: # degree d
    polynomial_order =

# regularization constant lambda
    regularization_param =
    theta, train_errors, val_errors = polynomial_regression(poly_data, polynomial_order)

In []: # Call plot_results() to see how your polynomial fits.
    def plot_results(theta, X, Y):
        y_hat = sum([t*X**i for i,t in enumerate(theta)])
        plt.scatter(X, y_hat, s=10, color='r')
        plt.scatter(X, Y, s=10)
        plt.show()
```