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# LINEAR REGRESSION

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# LINEAR REGRESSION

## Applied Mathematics And Statistics

### 1. INFORMATION

#### Lecturer:

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- Le Thanh Tung
- Phan Thi Phuong Uyen

**Class:** 20CLC08

#### Student:

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#### Completion

Requirement	Detail	Completion
1	Use all 10 features the topic provides	100%
2	Build a model using only 1 feature, find the model that gives the best results	100%
3	Build the freedom model, find the model that gives the best results	100%
4	Explain all functions and the implement to the report	100%

### 2. CONTENTS

#### 2.1.Library Support

```
import pandas as pd
import numpy as np
# Import thêm dữ thư viện nếu cần
import math
```

- **pandas library** is used for reading two files namely 'train.csv' and 'test.csv'
- **numpy library** is used for matrix processing

- **math library**: I use it to calculate RMSE by square root of 2 of MSE by `math.sqrt()`

## 2.2.Function

Four methods in class **OLSLinearRegression** (**fit**, **get\_params**, **predict**, **mse**) I refer from **lab04.ipynb** on moodle of lecturer. Other functions, do it by myself

### 2.2.1. *fit(self, X, y)*

**Input**: matrix X (mxn), matrix y (m)

**Output**: Object with X,y,w properties

**Approach**: Calculate the w value by fomular  $w=(X^T.X)^{-1}.X^T.y$

### 2.2.2. *get\_params(self)*

**Input**: None

**Output**: w value in object

**Approach**: Return self.w

### 2.2.3. *predict(self, X)*

**Input**: matrix X (mxn)

**Output**: matrix y (m)

**Approach**: Use the w value calculated in previous to calculate the predict y value

### 2.2.4. *mse(y, y\_hat)*

**Input**: matrix y result, matrix y\_hat result that predicted by calculating with w value

**Output**: the MSE value

**Approach**: Calculate MSE value by fomular

$$MSE = \frac{1}{n} \sum_{i=1}^n (y_i - \hat{y}_i)^2$$

### 2.2.5. *rmse(y, y\_hat)*

**Input**: matrix y result, matrix y\_hat result that predicted by calculating with w value

**Output**: the MSE value

**Approach:** Calculate RMSE value by square root of 2 of MSE

### 2.2.6. *FrameToNumpy(X\_train, y\_train, X\_test, y\_test)*

I refer the function `to_numpy()` from [Python Examples \(\\*\\*\)](#)

**Input:** 4 matrix with frame datatype

**Output:** 4 matrix with np.array datatype

**Approach:** Use function `to_numpy()` (\*\*) to convert it

### 2.2.7. *Init\_Data()*

**Input:** None

**Output:** 2 matrix clone and the number of set (after divided the total data set by 5)

**Approach:**

- Copy from the original matrix and make a shuffle ( I will describe later)
- Divide total data set by 5

**Implement:**

- Use `.copy()` to copy data from X,y\_train to X,y\_train\_clone
- Make a shuffle by `Shuffle_Data(X_train_clone, y_train_clone)` function
- Get len of X\_train\_clone and divided by 5 to save in numberSet

### 2.2.8. *Shuffle\_Data(X\_train\_clone, y\_train\_clone)*

The idea of code I refer from DelftStack (\*)

**Input:** 2 matrix

**Output:** 2 matrix after shuffle data

**Approach:** Use the random array index to change the position of each element in matrix ( The index of two matrixes are the same )

**Implement:**

- Use `np.arange(len(X_train))` to get an array index in order with size = X\_train's length
- Use `np.random.shuffle(rand_result)` to shuffle array index
- `X_train_clone[rand_result]` to reassign value at each index with new index that defined in array random index `rand_result`

### 2.2.9. *cross\_validation(X\_train\_feature, y\_train\_feature, numberSet)*

This function implement the 5-fold cross validation method to find the best model

**Input:** 2 matrix feature, the numer of set

**Output:** RMSE of the model

**Approach:**

- Divide the total sets by 5. In this project, there is 5 groups of set
- Use 1 group ( include X\_train\_feature and y\_train\_feature ) for validating. The others are use for training
- Changing validating group hay training group in loop until each group becomes validating group one time. For example, the first time use the 1<sup>st</sup> group for validating and 2<sup>nd</sup>, 3<sup>rd</sup>, 4<sup>th</sup>, 5<sup>th</sup> group for training so that the second time use the 2<sup>nd</sup> group for validating and 1<sup>st</sup>, 3<sup>rd</sup>, 4<sup>th</sup>, 5<sup>th</sup> group for training,....
- Sum the RMSE at each time in loop. After 5 times the training and validating, divided the RMSE by 5 to get the average RMSE in this model

**Implement:**

- Use loop from 0 to 5 (i=0; i<5) to do with 5 times validating and training data
- There are 4 matrix to store data at each loop **X\_j\_val** and **y\_j\_val** ( for validate ), **X\_j\_train** and **y\_j\_train** ( for training )
- Assign data to **X\_j\_val**

```
X_j_val = X_train_feature[numberSet*j:numberSet*(j+1),:]
```

**X\_j\_val** will get data from **X\_train\_feature** with all data in column and specific rows in each loop ( mutiple 217 with j ). [For example:](#)

**The first loop:** **X\_j\_val** = X\_train\_feature[0:numberSet,:] = X\_train\_feature[0:217,:]

**The second loop:** **X\_j\_val** = X\_train\_feature[217:434,:]

.....

- Assign data to **y\_j\_val** is the same with **X\_j\_val** without choosing column (':') because **y\_j\_val** just has 1 column and n row. So we need to get all data in specific rows

```
y_j_val = y_train_feature[numberSet*j:numberSet*(j+1)]
```

- Assign data to **X\_j\_train**

**X\_j\_train** will get data from **X\_train\_feature** except **X\_j\_val**.

The above code I merge 2 matrix before (from 0 to j) and after (from j+1 to end) **X\_j\_val**

```
X_j_train = np.concatenate([X_train_feature[0:numberSet*j,:],
X_train_feature[numberSet*(j+1):len(X_train_feature),:]])
```

matrix by **np.concatenate**. For example

**Supposed that, X\_j\_val** = X\_train\_feature[0:217,:]. So that **X\_j\_train** will merge two matrix with get all columns data and specific rows: **X\_train\_feature[0:0,:]** and **X\_train\_feature[217:1085,:]**

**Supposed that, X\_j\_val** = X\_train\_feature[217:434,:]. So that **X\_j\_train** will merge two matrix with get all columns data and specific rows: **X\_train\_feature[0:217,:]** and **X\_train\_feature[434:1085,:]**

.....

- Assign data to **y\_j\_train** is the same with **X\_j\_train** without choosing column (':') because **y\_j\_train** just has 1 column and n row. So we need to get all data in specific rows

```
y_j_train = np.concatenate([y_train_feature[0:numberSet*j],
y_train_feature[numberSet*(j+1):len(y_train_feature)]])
```

- Call the method **fit(X\_j\_train, y\_j\_train)** to calculate w values and predict the result by method **predict(X\_j\_val)** that stored in **y\_j\_val\_pre**.
- Calculate the RMSE of each loop by **rmse(y\_j\_val, y\_j\_val\_pre)** function and add it to **rmse\_val**
- Divided RMSE value by 5 after finished and return it as average RMSE of this model

### 2.2.10.first\_Model\_Featurel(x)

**Input:** Original matrix after shuffle

**Output:** The feature matrix corresponding the model defined

**Approach and Implement:** The first model I implemented is

Life Expectancy= Polio\*w1 + Diphtheria\*w2 + Income composition of resources\*w3 + Schooling\*w4

I choose this model because this is 4 best features (min RMSE) in 10 features. So I make an addition between them to create an RMSE model.

I use concatenate method I merge corresponding column (column 2,3,8,9 with begin index=0) with each other to create the matrix feature X and return it



### 2.2.11. *second\_Model\_Featurel(x)*

**Input:** Original matrix after shuffle

**Output:** The feature matrix corresponding the model defined

**Approach and Implement:** The second model I implemented is

$$\text{Life Expectancy} = (\text{Schooling}^3) * w1 + (\text{Income composition of resources}^2) * w2 + \text{Diphtheria} * w3$$

I choose this model because this is 3 best features (min RMSE) in 10 features. After a number of times tries, I find this model has the good RMSE with cubic function

I use concatenate method I merge corressponding column ((column 3)^1, (column 8)^2, (column 9)^3) with begin index=0) with each other to create the matrix feature X and return

### 2.2.12. *third\_Model\_Featurel(x)*

**Input:** Original matrix after shuffle

**Output:** The feature matrix corresponding the model defined

**Approach and Implement:** The third model I implemented is

$$\text{Life Expectancy} = (\text{Schooling} * \text{Income composition of resources}) * w1 + (\text{BMI} + \text{Polio} + \text{Diphtheria}) * w2$$

I choose this model because this is 5 best features (min RMSE) in 10 features. After a number of times tries, I find this model has the good RMSE with combining multiplication and summation

I use concatenate method I merge corressponding column ((column 8\*column 9) and (column 1 + column 2 + column 3)) with begin index=0) with each other to create the matrix feature X and return

## 2.3. Requirements

### 2.3.1. *Model using all 10 features the topic provides*

**Approach:**

- Calculating the w value by training data with 10 features ( in file train.csv )
- Using w value to calculate the predict result
- Determining the RMSE between the predict result and the result in file test.csv

**Implement:**



- I call the method `fit(X_train, y_train)` in class `OLSLinearRegression` to calculate the `w` values and save the entire object to `lr` with features in file `train.csv`
- The object `lr` calls the method `predict(X_test)` to calculate the predict the result with 10 features in file `test.csv` and `w` value in object `lr` that calculated in the previous step
- Having the predict result calculated by `w`, I calculate the RMSE between the predict result and the result in file `test.csv`

### 2.3.2. Models using all 1 feature the topic provides, find the best model

#### Approach:

- Calculating the `w` value by training data with only 1 feature ( in file `train.csv` )
- Using `w` value to calculate the predict result
- Determining the RMSE between the predict result and the result
- All of above are implemented by **5-fold cross validation** ( Divide dataset into 5 groups , 1 group for testing, 4 groups remaining for training data, and calculate the RMSE, do it 5 times until each group becomes validating group and divided RMSE by 5 to get the average RMSE after executing )
- Do it with 10 features in `train.csv` ( 10 times, each time works with 1 feature)

#### Implement:

- Initializing `X_train_clone`, `y_train_clone` and `numberSet` by `Init_Data()` function ( I have explained clearly in the previous topic [2.2.7](#) )
- Calculate the RMSE value in a model by calling the `cross_validation(X_train_feature, y_train_feature, numberSet)` ( I have explained clearly in the previous topic [2.2.9](#) )
- To calculate 10 RMSE values in 10 features in `train.csv`, I initialize an array to store 10 RMSE values and call the `cross_validation(X_train_feature, y_train_feature, numberSet)` in loop with index `i` in range (0,10)
- In loop, `X_train_feature = X_train_clone[:,i:i+1]` and `y_train_feature = y_train_clone`.

#### For example

- With index `i=0`: `X_train_feature = X_train_clone[:,0:1]` means getting all data at column 0 as the first feature in `train.csv`
- With index `i=1`: `X_train_feature = X_train_clone[:,1:2]` means getting all data at column 1 as the second feature in `train.csv`
- .....
- Finally, after doing loop to calculate the RMSE of 10 features, I get an array storing 10 RMSE values (`rmse_arr`). I use `argmin()` function to get the index in the array that has minimum values ( It means the  $i^{\text{th}}$  feature has smallest RMSE ) and save to `bestIndex` because the model has smallest RMSE is the best model

- The best feature will be display by `X_train[:,bestIndex]` and save to `X_best_feature` (Get the column with bestIndex)
- I call the method `fit(X_train, y_train)` in class `OLSLinearRegression` with `X_train = X_best_feature` ( The best feature ) to calculate the `w` values and save the entire object to `lr` with features in file `train.csv`
- The object `lr` calls the method `predict(X_test)` to calculate the predict the result with the feature having the corresponding `bestIndex` in file `test.csv` and `w` value in object `lr` that calculated in the previous step
- Having the predict result calculated by `w`, I calculate the RMSE between the predict result and the result in file `test.csv`

### 2.3.3. Freedom models, find the best model

#### Approach:

- Create 3 models ( `X_train_feature` ) by 3 function (`first_Mode_Featurel`, `second_Mode_Featurel`, `third_Mode_Featurel`) that I have explained in 2.2.10, 2.2.11, 2.2.12
- Calculate 3 RMSE of 3 models
- Find the best model by minimum RMSE
- Train the best model in `train.csv` to get `w` values in class `OLSLinearRegression`
- Calculate the predict result by using `w` value with coressponding model in `test.csv`
- Calculate the RMSE between result in `test.csv` and the predict result

#### Implement:

- Create 3 models ( `X_train_feature` ) and save in list `X_train_feature_array` by 3 function(`first_Mode_Featurel(x)`,`second_Mode_Featurel(x)`,`third_Mode_Featurel(x)`)) that I have explained in 2.2.10, 2.2.11, 2.2.12 with `x=X_train_clone`
- Create 3 models ( `X_test_feature` ) and save in list `X_test_feature_array` by 3 function(`first_Mode_Featurel(x)`,`second_Mode_Featurel(x)`,`third_Mode_Featurel(x)`)) that I have explained in 2.2.10, 2.2.11, 2.2.12 with `x=X_test`. This step is used for when I get the best model in train.csv, I can get the best model in test.csv with same index and structure
- Calculate the RMSE value in each model by calling the `cross_validation(X_train_feature, y_train_feature, numberSet)` ( I have explained clearly in the previous topic 2.2.9) in list `RMSE_array`
- Finally, after calculating the RMSE of 3 models, I get an array storing 3 RMSE values (`RMSE_arr`). I use `argmin()` function to get the index in the array that has minimum

values ( It means the  $i^{\text{th}}$  feature has smallest RMSE ) and save to **bestIndex** because the model has smallest RMSE is the best model

- The best feature will be display by `X_train_feature_array[bestIndex]` and save to `X_best_feature` (Get the model with bestIndex)
- I call the method `fit(X_train, y_train)` in class `OLSLinearRegression` with `X_train = X_best_feature` ( The best feature ) to calculate the **w** values and save the entire object to **lr** with features in file **train.csv**
- The object **lr** calls the method `predict(X_test)` to calculate the predict the result with the feature having the corresponding **bestIndex** in file **test.csv** and **w** value in object **lr** that calculated in the previous step
- Having the predict result calculated by **w**, I calculate the RMSE between the predict result and the result in file **test.csv**

### 3. RESULTS & EVALUTIONS

#### 3.1.Model using all 10 features the topic provides

W values

```
w values-----
w1: 0.015101362735318279
w2: 0.09021998065775627
w3: 0.04292181752549435
w4: 0.13928911689488216
w5: -0.5673328270884068
w6: -0.0001007651148748953
w7: 0.7407134377587112
w8: 0.19093579767396474
w9: 24.505973591149445
w10: 2.393516607832779
```

RMSE on **test.csv**

```
RMSE: 7.064046430584705
```

Regression Formula

**Life Expectancy** =  $0.015101 \cdot \text{Adult Mortality} + 0.090220 \cdot \text{BMI} + 0.042922 \cdot \text{Polio} + 0.139289 \cdot \text{Diphtheria} - 0.567333 \cdot \text{HIV/AIDS} - 0.000101 \cdot \text{GDP} + 0.740713 \cdot \text{Thinness age 10-19} + 0.190936 \cdot \text{Thinness age 5-9} + 24.505974 \cdot \text{Income composition of resources} + 2.393517 \cdot \text{Schooling}$

## Evaluation

- W values have inverse ratio with feature value
- The RMSE value on **test.csv** with 10 features is small
- If the features are modified, it means W values are changed that leads to the shift in RMSE value

### 3.2.Models using all 1 feature the topic provides, find the best model

RMSE 10 features on train.csv

```
RMSE feature 1: 46.29318840485384
RMSE feature 2: 27.89566460172918
RMSE feature 3: 18.03199333638845
RMSE feature 4: 15.999435116642521
RMSE feature 5: 67.1076781800127
RMSE feature 6: 60.208226896875715
RMSE feature 7: 51.75287025474305
RMSE feature 8: 51.665179224465724
RMSE feature 9: 13.291015950250417
RMSE feature 10: 11.786003414181945
```

Best feature

```
Best feature is feature: 10
Feature 10 [[ 9.9]
 [ 9.8]
 [ 9.5]
 ...
 [10. ]
 [ 9.8]
 [ 9.8]]
```

W values

```
W: 5.5573993976919205
```

RMSE on **test.csv**

```
RMSE: 10.26095039165537
```

Regression Formula

**Life Expectancy** = 5.5573994\*Schooling

## Evaluation

No.	Feature	RMSE
1	Adult Mortality	46.245725793822416
2	BMI	27.955334582292416
3	Polio	18.02915163159739
4	Diphtheria	15.845876548938545
5	HIV/AIDS	67.09917048487907
6	GDP	60.20975942082864
7	Thinness age 10-19	51.855291331394504
8	Thinness age 5-9	51.75814155839241
9	Income composition of resources	13.27861308365102
10	Schooling	11.771538278894493
RMSE of Model Schooling on <b>test.csv</b>		<b>10.26095039165537</b>

- W values have inverse ratio with feature value
- The RMSE of **Schooling** feature is smallest (11.77) that means **Schooling** is the best feature to create a model with 1 feature
- The RMSE of **HIV/AIDS** feature is biggest (67.1) that means **HIV/AIDS** is the worst feature that should not be used to create a model
- There is a large difference between RMSE value of each feature. Some features have small RMSE, but the others are bigger than a lot (**Polio** and **GDP**)
- There are some features that have RMSE value is a little bit the same with the other (**Schooling** and **Income composition of resources**, **Polio** and **Diphtheria**)
- The RMSE value on **test.csv** with 1 feature is greater than it with 10 features (10.26 > 7.06). So that the model with 1 feature is worse than the other with 10 features.
- If the features are modified, it means W values are changed that leads to the shift in RMSE value

### 3.3.Freedom models, find the best model

#### Model 1

$$\text{Life Expectancy} = (\text{Schooling}^3) * w_1 + (\text{Income composition of resources}^2) * w_2 + \text{Diphtheria} * w_3$$

## Model 2

$$\text{Life Expectancy} = \text{Polio} * w1 + \text{Diphtheria} * w2 + \text{Income composition of resources} * w3 + \text{Schooling} * w4$$

## Model 3

$$\text{Life Expectancy} = (\text{Schooling} * \text{Income composition of resources}) * w1 + (\text{BMI} + \text{Polio} + \text{Diphtheria}) * w2$$

RMSE 3 models on **train.csv**

```
RMSE model 1: 9.55407523675269
RMSE model 2: 13.200600645811434
RMSE model 3: 12.951990251649391
```

Best model

```
Best model is model: 1
Best model: [[99.    91.    0.822 17.2 ]
 [ 7.    69.    0.613 12.9 ]
 [42.    42.    0.507  9.7 ]
 ...
 [98.    98.    0.569  9.8 ]
 [78.    77.    0.668 11.  ]
 [91.    91.    0.662 12.8 ]]
```

W values

```
W: [[0.14820296]
 [0.27274154]
 [5.55143115]
 [2.31214422]]
```

RMSE on **test.csv**

```
RMSE: 9.692944694229913
```

Regression Formula

$$\text{Life Expectancy} = 0.14820296 * \text{Polio} + 0.27274154 * \text{Diphtheria} + 5.55143115 * \text{Income composition of resources} + 2.31214422 * \text{Schooling}$$



## Evaluation

No.	Feature	RMSE
1	Model 1	9.55407523675269
RMSE of Model 1 on <b>test.csv</b>		<b>9.692944694229913</b>
2	Model 2	13.200600645811434
3	Model 3	12.951990251649391

- W values have inverse ratio with feature value
- The RMSE of **Model 1** is smallest (9.55) that means **Model 1** is the best model
- The RMSE of **Model 2** is biggest (13.2) that means **Model 2** is the worst model
- Difference between RMSE value of each model is not too high (9.55, 13.2, 12.95).
- The RMSE value on **test.csv** with **Model 1** is greater than it with 10 features at requirement 1 ( $9.69 > 7.06$ ) and smaller than 1 feature at requirement 2 ( $9.69 < 10.26$ ). So that the **Model 1** is worse than the model with 10 features and better than the model with 1 feature.
- The feature with small RMSE values creates small RMSE model
- If the features are modified, it means W values are changed that leads to the shift in RMSE value

## 4. REFERENCES

- (\*) Shuffle Data: <https://www.delftstack.com/howto/numpy/python-numpy-shuffle-two-arrays/>
- (\*\*) Convert Pandas DataFrame to Numpy Array (using the function `DataFrame.to_numpy()`):  
<https://pythonexamples.org/convert-pandas-dataframe-to-numpy-array/#:~:text=To%20convert%20Pandas%20DataFrame%20to,returned%20ndarray%20is%202%2Ddimensional.>
- Merge 2 numpy array with `np.concatenate`:  
<https://numpy.org/doc/stable/reference/generated/numpy.concatenate.html>