# LAB4 tutorial for Machine Learning Linear Regression

The document description are designed by Jla Yanhong in 2022. Sept. 22th

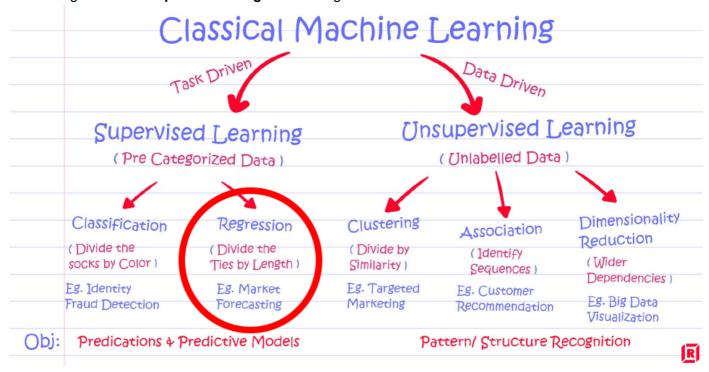
### **Objective**

- · Master the linear regression algorithm.
- · Understanding Gradient Descent
- · Polynomial Regression
- · Learn how to evaluate regression models
- · Complete the LAB assignment and submit it to BB.

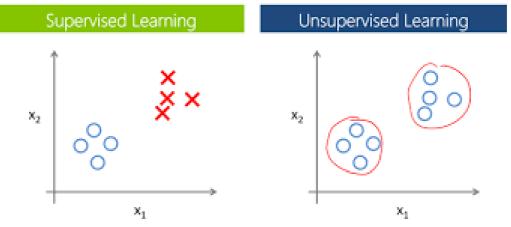
### 1 Linear Regression

### 1.1 Conceptual Overview

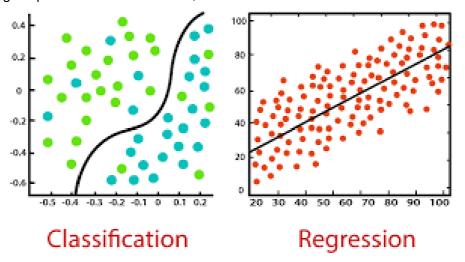
Linear Regression is "supervised" "regression" algorithm.



Supervised meaning we use labeled data to train the model.



Regression meaning we predict a numerical value, instead of a "class".



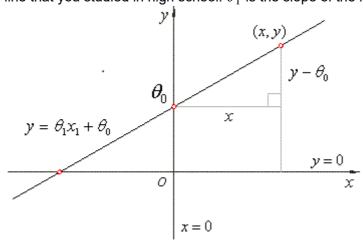
### 1.2 Linear Regression

In statistics, linear regression is a linear approach to modelling the relationship between a dependent variable and one or more independent variables.

We will define a linear relationship between these two variables as follows:

$$y = \theta_0 + \theta_1 x_1$$

This is the equation for a line that you studied in high school.  $\theta_1$  is the slope of the line and  $\theta_0$  is the y intercept.



When there are multiple independent variables, the linear relationship becomes as follows:

$$y = \theta_0 + \theta_1 x_1 + \theta_2 x_2 + \ldots + \theta_n x_n$$

**Simple Linear Regression** 

$$y = \theta_0 + \theta_1 x_1$$

**Multiple Linear Regression** 

$$y = \theta_0 + \theta_1 x_1 + \theta_2 x_2 + \dots + \theta_n x_n$$

Let  $\vec{x}$  be the independent variable and **y** be the dependent variable. The general form of the model's prediction:

$$\hat{y} = h_{\theta}(x) = \theta_0 + \theta_1 x_1 + \theta_2 x_2 + \dots + \theta_n x_n = \theta^T \cdot x$$

 $\theta$  is the model's parameter vector, containing the bias term  $\theta_0$  and the feature weights  $\theta_1$  to  $\theta_n$ . (n = number of features).

 $\theta^T \cdot x$  is the dot product of the vectors  $\theta$  and x, which is, of course, equal to  $\theta_0 + \theta_1 x_1 + \theta_2 x_2 + \ldots + \theta_n x_n$ .

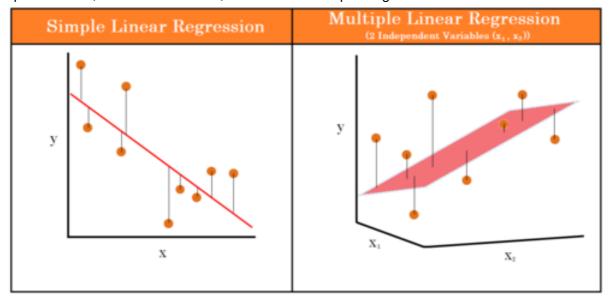
There is an error between the real y and the predicted  $\hat{y}$ :

$$y = \hat{y} + \varepsilon = h_{\theta}(x) + \varepsilon = \theta^{T} \cdot x + \varepsilon$$

Our challenge today is to determine the value of  $\theta$ .

For a single variable, such that the line corresponding to those values is the best fitting line .

For multiple variables, such as 2 variables, the surface corresponding to these values is the surface of best fit.



### 1.3 Cost Function

Cost Function evaluates the model's predictions and tells us how accurate are the model's predictions. The lower the value of the cost function, better accurate the predictions of the model. They are many cost functions to choose, but we will use the Mean Squared Error (MSE) cost function.

The MSE function calculates the average of the squared difference between the prediction and the actual value (y).

$$J(\theta) = MSE = \frac{1}{2m} \sum_{j=1}^{m} (h_{\theta}(x^{j}) - y^{j})^{2} = \frac{1}{2m} \sum_{j=1}^{m} (\theta_{0} + \theta_{1}x_{1}^{j} + \theta_{2}x_{2}^{j} + \dots + \theta_{n}x_{n}^{j} - y^{j})^{2}$$

 $x_i^j$ : The  $i_{th}$  feature of the  $j^{th}$  sample

 $y_i^j$  : the  $i_{th}$  label of the  $j^{th}$  sample

 $(x^j, y^j)$ : the  $j^{th}$  sample

m: total number of instances in your dataset

### 1.4 Least-squares estimation

Now that we have determined the cost function, the only thing left to do is minimize it. This is done by finding the partial derivative of  $J(\Theta)$ , equating it to 0 and then finding an expression for  $\Theta$ .

The loss function can be written as:

$$J(\theta) = \frac{1}{2m} \sum_{j=1}^{m} (h_{\theta}(x^{j}) - y^{j})^{2}$$
$$= \frac{1}{2m} \sum_{j=1}^{m} (\theta^{T} \cdot x^{j} - y^{j})^{2}$$
$$= \frac{1}{2m} (X\theta - y)^{T} (X\theta - y)$$

As the loss is convex the optimum solution lies at gradient zero. The gradient of the loss function is :

$$\frac{\partial J(\theta)}{\partial \theta} = \frac{\partial \frac{1}{2m} (X\theta - y)^T (X\theta - y)}{\partial \theta}$$

$$= \frac{\partial \frac{1}{2m} (\theta^T X^T - y^T) (X\theta - y)}{\partial \theta}$$

$$= \frac{\partial \frac{1}{2m} (\theta^T X^T X \theta - \theta^T X^T y - y^T X \theta + y^T y)}{\partial \theta}$$

$$= \frac{1}{2m} (2X^T X \theta - X^T y - (y^T X)^T)$$

$$= \frac{1}{m} X^T X \theta - X^T y$$

When X is a matrix of full rank, setting the gradient to zero produces the optimum parameter:

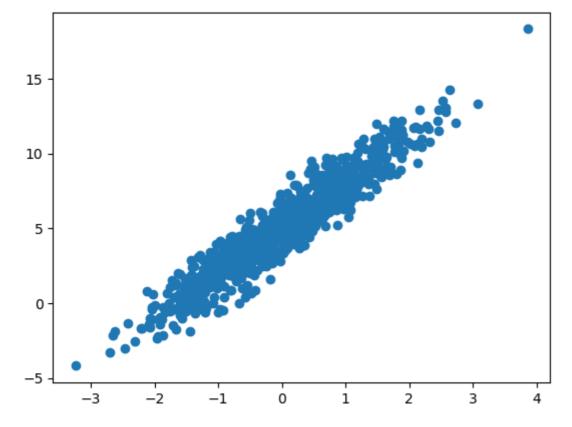
$$\theta^* = (X^T X)^{-1} X^T y$$

\end{aligned}

### **Hands-on Coding**

### In [105]:

```
1
   import numpy as np
   from sklearn.model_selection import train_test_split
   from sklearn.metrics import r2_score
 4
 5
    import matplotlib.pyplot as plt
 6
   np. random. seed (42)
 7
    X = \text{np. random. randn} (1000, 1)
 8
   \# y = 5 + 3 * X + Gaussian noise -> because in real-world it is very unlikely to get data that
9
   y = 5 + 3 * X + np. random. random (1000, 1)
10
   plt.scatter(X, y)
11
   plt.show()
12
13
```



### In [106]:

```
X b = np.c [np.ones((1000, 1)), X] # Adding the bias term which is equal to 1
 2
   # Dividing the data into train and test sets
 4
   X_train, X_test, y_train, y_test = train_test_split(X_b, y, test_size=0.2, random_state=42)
    theta_optimize = np. linalg. inv(X_train. T. dot(X_train)). dot(X_train. T). dot(y_train)
 6
 7
 8
 9
   #obtained feature weights w0 = 5.04, w1 = 2.94
   print(theta optimize)
10
11
12
   # Predicting new data with the obtained feature weights
   y_pred = X_test.dot(theta_optimize)
13
   r2_score(y_test, y_pred)
15
```

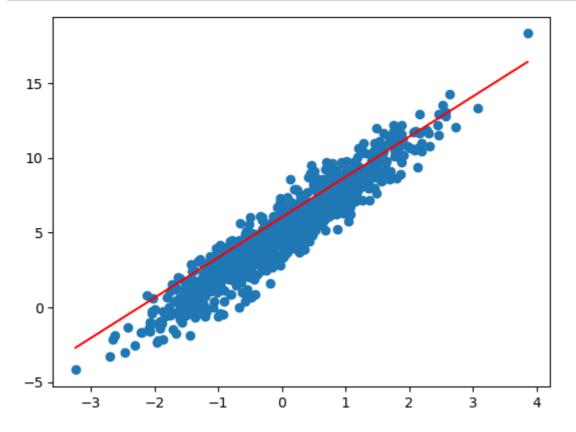
[[5. 08703256] [2. 93815133]]

### Out[106]:

0.8961012486926588

### In [107]:

```
plt.scatter(X, y)
plt.plot([min(X), max(X)], [min(y_pred), max(y_pred)], color='red') # regression line
plt.show()
```



You have seen it has predicted the feature weights very close to the actual values ( y = 5 + 3 \* X + Gaussiannoise), but due to the noise in the data it is unable to predict the exact values, but the predictions were close enough.

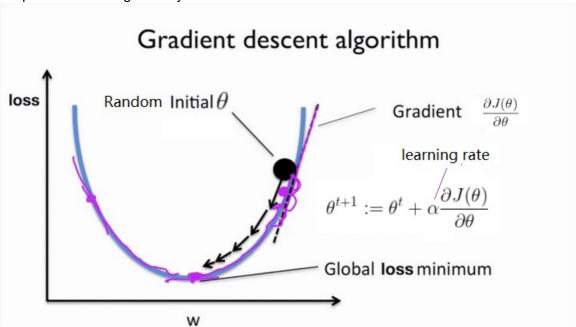
### **Disadvantages**

- It is computationally expensive if you have a large number of features.
- If there are any redundant features in your data, they the matrix inversion in the normal equation is not possible. In that case, the inverse can be replaced by the pseudo inverse.

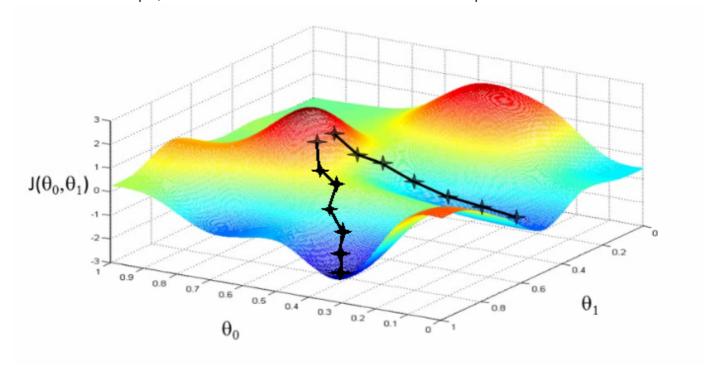
### 1.5 Gradient Descent

In this section you can learn how the gradient descent algorithm works and implement it from scratch in python.

Gradient Descent minimizes the cost function by iteratively moving in the direction of steepest descent, updating the parameters along the way.



In a real world example, it is similar to find out a best direction to take a step downhill.



We take a step towards the direction to get down. From the each step, you look out the direction again to get down faster and downhill quickly.

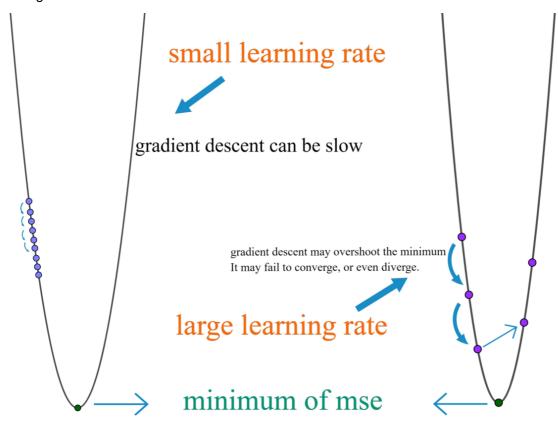
To find the best minimum, repeat steps to apply various values for  $\theta$ . In other words, repeat steps until convergence.

repeat until convergence {

$$\theta^{t+1} := \theta^t - \alpha \frac{1}{m} \frac{\partial J(\theta)}{\partial \theta}$$

The choice of correct learning rate is very important as it ensures that Gradient Descent converges in a reasonable time.

- If we choose α to be very small, Gradient Descent will take small steps to reach local minima and will take a longer time to reach minima.
- If we choose **α to be very large**, Gradient Descent can overshoot the minimum. It may fail to converge or even diverge.



**Hands-on Coding** 

implement Linear Regression from scratch

### In [108]:

```
import numpy as np
 2
   from sklearn.metrics import r2_score
 4
   np. random. seed (42)
 6
   learning_rate = 0.1
 7
   iterations = 50
   m = 100 # total number of samples
   theta = np. random. randn(2, 1) # random initialization
   for iteration in range (iterations):
10
        gradient = 2/m * X_train. T. dot(X_train. dot(theta) - y_train)
11
12
        theta = theta - learning_rate * gradient
13
14 # Output
15
   print(theta)
16
17
18 | # array([[5.08703256],
19 #
             [2. 93815133]])
20
21 # Predicting new values with gradient descent
22 | y_pred = X_test. dot(theta)
   r2_score(y_test, y_pred)
```

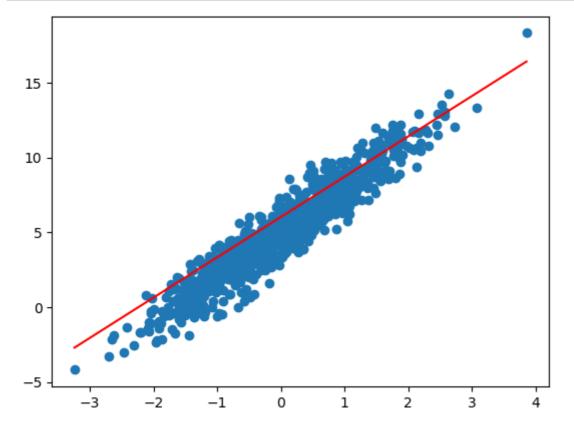
[[5. 08703256] [2. 93815133]]

### Out[108]:

0.8961012486937926

```
In [109]:
```

```
plt.scatter(X, y)
plt.plot([min(X), max(X)], [min(y_pred), max(y_pred)], color='red') # regression line
plt.show()
```



Implementing Linear Regression in Scikit-Learn

### In [110]:

```
from sklearn.linear model import LinearRegression
 2
   X_train = X_train[:, 1] # The sklearn model will automatically add the bias term so we do not ha
 4
    X \text{ test} = X \text{ test}[:, 1]
 5
 6
 7
    linear_regression = LinearRegression()
   linear_regression.fit(X_train.reshape(-1, 1), y_train)
 8
 9
    print(linear_regression.intercept_)
10
    # Output
    #array([5.08703256])
11
12
13
    print(linear_regression.coef_)
14
   # Ouput
15
   #array([[2.93815133]])
16
    # Predicting new values and calculating the r2 score
17
    linear regression.score(X test.reshape(-1, 1), y test)
18
19
20
    # output
21
    #0.8961012486926588
```

```
[5. 08703256]
[[2. 93815133]]
```

### Out[110]:

0.8961012486926588

As you can see the values we got from the normal equation, gradient descent, sklearn are nearly the same.

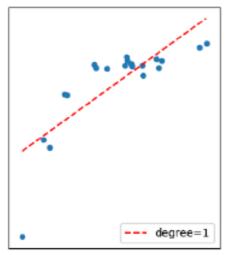
#### **Disadvantages**

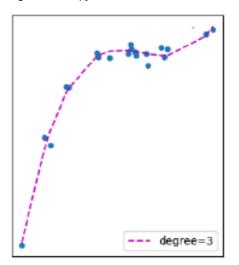
- There are possibilities for the gradient descent to stuck in a local minimum if you use another cost function that is not of a convex shape.
- You should find the appropriate value for the learning rate.

Well if you have read this far and everything makes sense pat yourself on the back!. You have learned all the underlying concepts of linear regression. A

## 1.6 <u>Polynomial Regression</u> (https://en.wikipedia.org/wiki/Polynomial regression)

We use polynomial regression when the relationship between the independent and dependent variables is nonlinear.





Simple Linear Regression

Polynomial Regression

This is accomplished by "exponentiating" our variable by taking it to powers greater than 1.

Simple Linear Regression

$$y = \theta_0 + \theta_1 x_1$$

**Multiple Linear Regression** 

$$y = \theta_0 + \theta_1 x_1 + \theta_2 x_2 + ... + \theta_n x_n$$

Ploynomial Linear Regression

$$y = \theta_0 + \theta_1 x_1 + \theta_2 x_1^2 + ... + \theta_n x_1^n$$

### **Hands-on Coding**

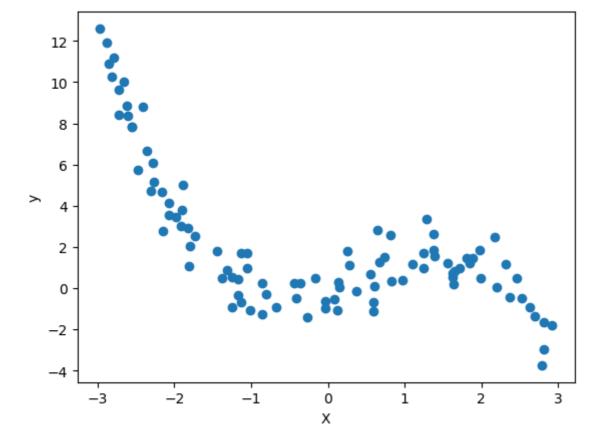
#### In [111]:

```
#step 1: Import the required libraries
import numpy as np
from scipy import stats
```

- 4 import matplotlib.pyplot as plt
- 5 from sklearn.preprocessing import PolynomialFeatures
- 6 from sklearn.linear model import LinearRegression
- 7 from sklearn.metrics import r2 score, mean squared error
- 8 | from sklearn.model\_selection import train\_test\_split

### In [112]:

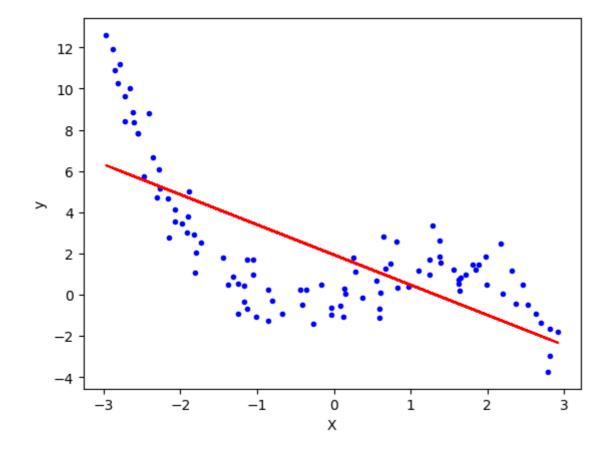
```
#step2: Load the data set. Here, We're generating data
 2
    sample_cnt= 100
    #X = np.linspace(start = -8, stop = 8, num = sample_cnt)
    X = \text{np. random. uniform}(-3, 3, \text{size}=100)
 4
 6
    # curve using polynomial
    \theta 0, \theta 1, \theta 2, \theta 3 = 0.2, 1, 0.5, -0.4
    y = \theta 0 + \theta 1*X + \theta 2*(X**2) + \theta 3*(X**3) + np. random. normal(0, 1, size = sample_cnt)
 8
9
10
11
12
    plt.scatter(X, y)
13
    plt.xlabel('X')
14
    plt.ylabel('y')
15
    plt.savefig('regu-0.png', dpi=200)
16
    plt.show()
17
```



### In [113]:

```
#split data
 2
   X_train, X_test, y_train, y_test = train_test_split(X, y, train_size=0.8, random_state=1)
 4
   # step 3: using a linear regression model to predict
   lin reg = LinearRegression()
   lin_reg.fit(X_train.reshape(-1,1), y_train.reshape(-1,1))
 6
 8
   y_pred = lin_reg. predict(X_train. reshape(-1, 1))
 9
10
   a = 1in reg. coef [0][0]
   b = lin_reg.intercept_[0]
11
12
   plt.plot(X, y, 'b.')
13
   # plt.plot([min(X_train), max(X_train)], [min(y_pred), max(y_pred)], color='red') # regression
14
   plt.plot(X_train, y_pred, 'r-')
15
   plt.xlabel('X')
16
17
   plt.ylabel('y')
   plt.savefig('regu-1.png', dpi=200)
   print("The equation: y = {} x+{} ". format(a, b))
19
   #Calculate the error and evaluate the model
   print(lin_reg.intercept_, lin_reg.coef_)
21
   print(r2 score(y pred, y train))
23
   plt.show()
24
```

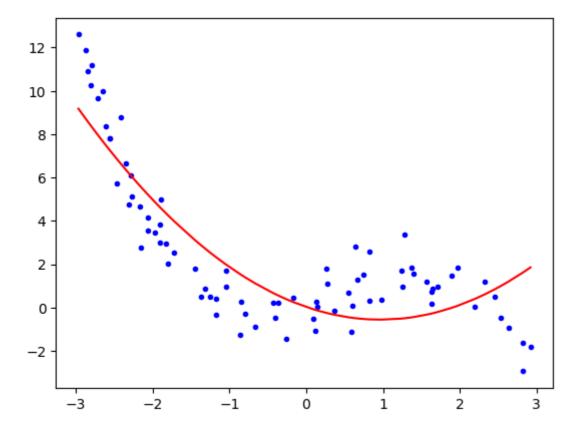
The equation: y = -1.4623971689071653x+1.9344273366540403 [1.93442734] [[-1.46239717]] 0.0640863835493316



### In [114]:

```
# Use a polynomial with degree of 2
    ploy = PolynomialFeatures(degree=2, include_bias=False)
 2
 3
    X_train2 = ploy.fit_transform(X_train.reshape(-1,1))
 4
 5
   lin_reg = LinearRegression()
    lin_reg.fit(X_train2, y_train.reshape(-1,1))
    print(lin_reg.intercept_, lin_reg.coef_) # [ 2.60996757] [[-0.12759678  0.9144504 ]]
 7
    a = 1in reg. coef [0][0]
   b = lin_{reg.coef_[0][1]}
    c = lin reg. intercept [0]
10
11
12
13
14
    y_pred = np. dot(X_train2, lin_reg. coef_.T) + lin_reg. intercept_
15
   plt.plot(X_train, y_train, 'b.')
16
    plt.plot(np.sort(X_train), y_pred[np.argsort(X_train)], color='r')
17
18
19
    plt. savefig('regu-2.png', dpi=200)
20
   plt.show()
    print ("The equation: y = \{:.3f\}x^2 + \{:.3f\}x + \{:.3f\}". format (a, b, c))
21
    print(r2_score(y_pred, y_train))
22
   print(r2 score(np.dot(ploy.transform(X test.reshape(-1,1)), lin reg.coef .T) + lin reg.intercer
```

### [0.0253991] [[-1.21577111 0.62996561]]



The equation:  $y = -1.216x^2+0.630x+0.025$ 

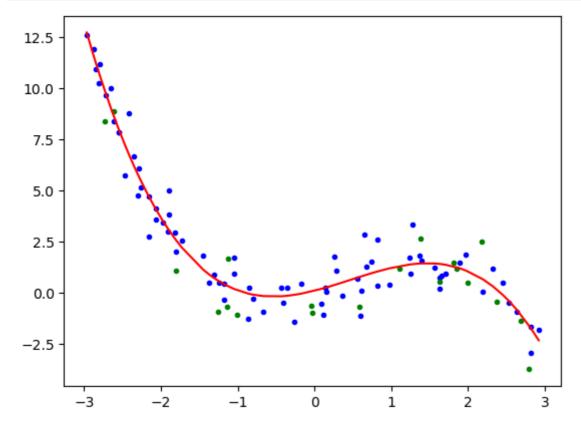
0. 63958477057975590. 1295465340821086

As the data extending to polynomial features, the value would be extremely large or small because of the power operation. That will influence the use of gradient descent which runs in background when we call fit(). So a normalization or standardization is necessary. See StandardScaler in preprocessing.

Pipeline can help us assemble several preprocessing functions and the learning process together.

### In [115]:

```
# Use a polynomial with degree of 3
   from sklearn.pipeline import Pipeline
 2
    from sklearn.preprocessing import StandardScaler
 4
    from sklearn.preprocessing import PolynomialFeatures
 5
    poly_reg = Pipeline([
 6
 7
        ('poly', PolynomialFeatures(degree=3)),
        ('std_scaler', StandardScaler()),
 8
 9
        ('lin_reg', LinearRegression())
    ])
10
11
    poly reg. fit (X train. reshape (-1, 1), y train. reshape (-1, 1))
12
13
14
    y_pred = poly_reg. predict(X_train. reshape(-1, 1))
15
16
17
18
    plt.plot(X_train, y_train, 'b.')
19
    plt.plot(X_test, y_test, 'g.')
20
    plt.plot(np.sort(X_train), y_pred[np.argsort(X_train)], color='r')
21
22
    plt. savefig('regu-3. png', dpi=200)
23
    plt.show()
24
    params =poly_reg.named_steps['lin_reg']
25
    print ("The equation: ", poly_reg. named_steps['lin_reg']. coef_, poly_reg. named_steps['lin_reg']. int
    print ("The equation: y = \{:.3f\}x^3 + \{:.3f\}x^2 + \{:.3f\}x + \{:.3f\}". format (params. coef_[0][0], params. c
26
27
    print(r2_score(y_pred, y_train))
    print(r2_score(poly_reg.predict(X_test.reshape(-1,1)), y_test))
28
29
    #print(mean squared error(y pred, y))
```

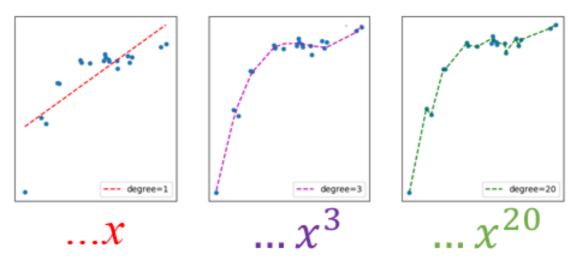


0.9413635998834047

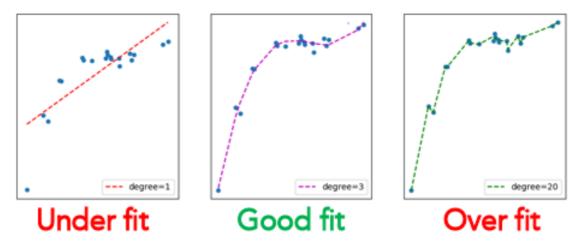
0.8303023309883463



In practice, we need to select the "degree" of the polynomial.



Our selection should seek to fit the current data well, and generalize to new data.



This challenge is known as the "bias variance tradeoff".

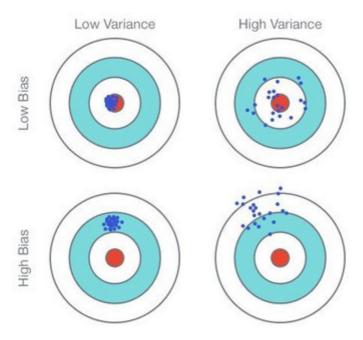
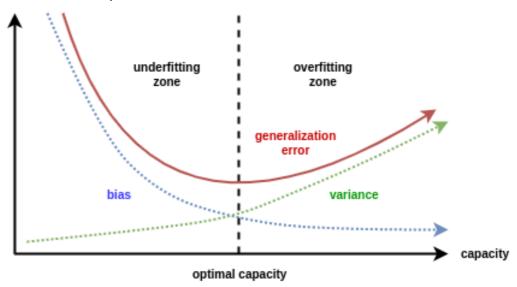


Fig. 1: Graphical Illustration of bias-<u>variance trade</u>-off , Source: Scott Fortmann-Roe., Understanding Bias-Variance Trade-off

Bias usually caused by underfitting, Variance caused by overfitting. What can we do to solve this problem?

• 1. Choose a better polynomial degree. As we increase the degree, the bias decreases, but the variance increases. We want to stop where these two factors are minimized.



Note: polynomial regression is very sensitive to outliers, and we must take care when selecting the degree to avoid overfitting.

2. Regularization .Regularization in scikit learn is <u>RidgeRegression</u>
 (<a href="https://en.wikipedia.org/wiki/Ridge\_regression#:~:text=Ridge%20regression%20is%20a%20method%20">https://en.wikipedia.org/wiki/Ridge\_regression#:~:text=Ridge%20regression%20is%20a%20method%20"
 ,which is in <u>linear\_model (https://scikit-learn.org/stable/modules/generated/sklearn.linear\_model.Ridge.html)</u>. Use it if you need regularization in your model.

Ridge regression is also known as L2 regularization. It is a regularized version of linear regression to find a better fitting line. It adds I2 penalty terms in the cost function and thereby reducing coefficients lower towards zero and minimizing their impact on the training data. It is useful to avoid over-fitting of

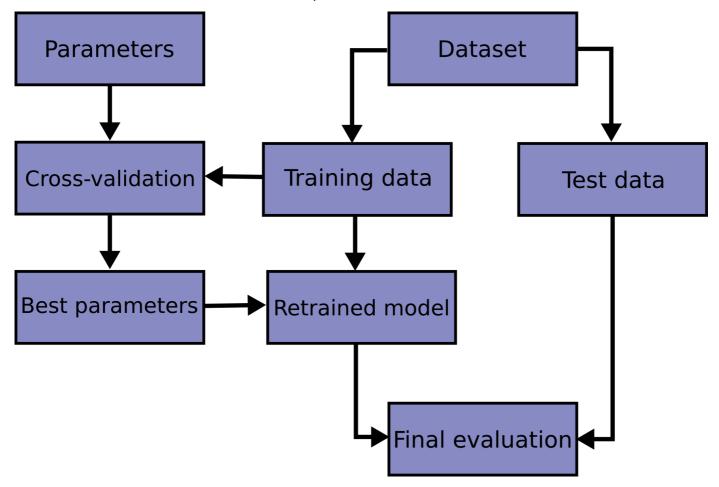
the data in a model. The alpha parameter of a model is used to calculate cost function to manage biasvariance tradeoff and reduce an error in prediction.

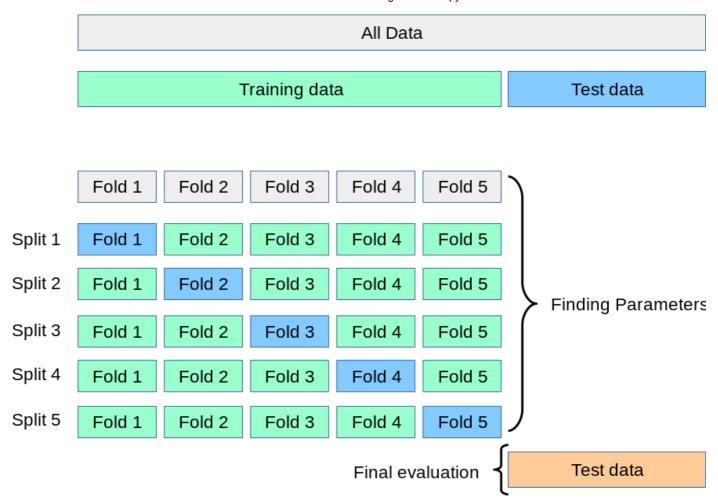
$$J(\theta) = \frac{1}{2m} \sum_{j=1}^{m} (h_{\theta}(x^{j}) - y^{j})^{2} + \alpha \sum_{i=1}^{n} \theta_{i}$$
$$= \frac{1}{2m} (X\theta - y)^{T} (X\theta - y) + \alpha \theta^{T} \theta$$

3. Use training set, validation set and test set or <u>cross validation (https://scikit-learn.org/stable/modules/cross\_validation.html)</u> to acquire a better model.

# 2 <u>Cross-validation (https://scikit-learn.org/stable/modules/cross\_validation.html)</u>

We often use cross validation to select parameters or to select models.



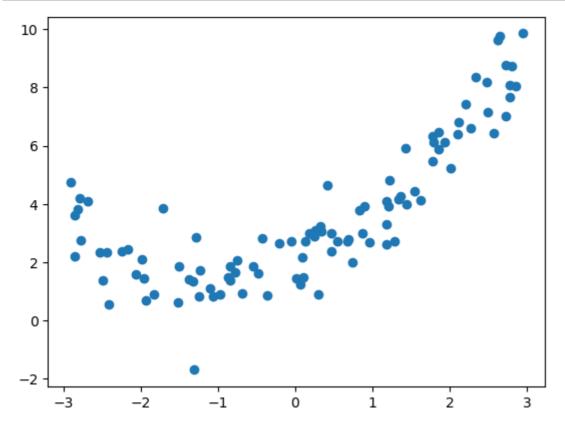


### For more detail:

- <a href="https://scikit-learn.org/stable/modules/cross\_validation.html">https://scikit-learn.org/stable/modules/cross\_validation.html</a> (https://scikit-learn.org/stable/modules/cross\_validation.html)
- <a href="https://en.wikipedia.org/wiki/Cross-validation\_(statistics)">https://en.wikipedia.org/wiki/Cross-validation\_(statistics)</a> (https://en.wikipedia.org/wiki/Cross-validation\_(statistics))
- <a href="https://scikit-learn.org/stable/modules/generated/sklearn.model\_selection.GridSearchCV.html">https://scikit-learn.org/stable/modules/generated/sklearn.model\_selection.GridSearchCV.html</a> (<a href="https://scikit-learn.org/stable/modules/generated/sklearn.model\_selection.GridSearchCV.html">https://scikit-learn.org/stable/modules/generated/sklearn.model\_selection.GridSearchCV.html</a> (<a href="https://scikit-learn.org/stable/modules/generated/sklearn.model\_selection.GridSearchCV.html">https://scikit-learn.org/stable/modules/generated/sklearn.model\_selection.GridSearchCV.html</a> (<a href="https://scikit-learn.org/stable/modules/generated/sklearn.model\_selection.GridSearchCV.html">https://scikit-learn.org/stable/modules/generated/sklearn.model\_selection.GridSearchCV.html</a>)

### In [117]:

```
import numpy as np
    import matplotlib.pyplot as plt
 2
   import pandas as pd
 4 from sklearn. model selection import train test split
    from sklearn.linear model import Lasso, Ridge
    from sklearn.model_selection import GridSearchCV
    from sklearn. pipeline import Pipeline
    from sklearn.preprocessing import StandardScaler
    from sklearn.preprocessing import PolynomialFeatures
10
    from sklearn.linear model import LinearRegression
11
    sample cnt = 100
    X = \text{np. random. uniform}(-3, 3, \text{size}=100). \text{ reshape}(-1, 1)
12
13
14
15
    # curve using polynomial
    \theta 0, \quad \theta 1, \quad \theta 2, \quad \theta 3 = 2, \quad 1, \quad 0.5, \quad 0
16
    y = \theta 0 + \theta 1*X + \theta 2*(X**2) + \theta 3*(X**3)
17
   y += np. random. normal(0, 1, size = sample cnt). reshape(-1, 1)
    X_train, X_test, y_train, y_test = train_test_split(X, y, random_state=1)
19
20
21
    plt.scatter(X, y)
22
    plt.show()
```



### In [118]:

```
1
 2
 3
    model = Pipeline([
 4
         ('poly', PolynomialFeatures()),
 5
         ('std_scaler', StandardScaler()),
 6
         ('ridge', Ridge())
 7
    7)
    params = { 'poly_degree': [1, 2, 3, 4],
 8
 9
    'ridge alpha':list(x / 10 for x in range(0, 101)),
10
               'ridge__max_iter':[5,10,100,1000]}
11
12
    grid search = GridSearchCV (model, param grid=params, cv=5)
13
    grid search. fit (X train, y train)
14
15
    print('The highest score of cross validation: {:.3f}\n'.format(grid_search.best_score_))
    print('The best parameters: {}\n'.format( grid search.best params ))
17
    print ('The highest score: {:..3f}\n'..format(grid_search.score(X_test, y_test)))
18
    # y hat = grid search.predict(np.array(X test))
19
20 | \# t = np. arange(len(X test))
   # plt.plot(t, y_test, 'r-', linewidth=2, label='Test')
# plt.plot(t, y_hat, 'g-', linewidth=2, label='Predict')
23 | # plt.legend(loc='upper right')
24 # plt. grid()
25
   # plt. show()
```

```
The highest score of cross validation:0.837

The best parameters: {'poly_degree': 2, 'ridge_alpha': 2.5, 'ridge_max_iter': 5}
```

### 3 Evaluation for Regression Models

Metrics commonly used to evaluate regression models are:

• Mean Absolute Error

The highest score: 0.897

- Mean Squared Error (MSE)
- Root Mean Squared Error (RMSE)
- R-Squared

### 1) Mean Absolute Error

$$MAE = rac{1}{N} \sum_{i=1}^{N} |y_i - \hat{ ext{y}}_i|$$

```
def MAE(y, y_pre):
    return np.mean(np.abs(y - y pre))
```

### **Advantages**

 MAE is not sensitive to outliers. Use MAE when you do not want outliers to play a big role in error calculated.

### **Disadvantages**

MAE is not differentiable globally. This is not convenient when we use it as a loss function, due to the
gradient optimization method.

### 2) Mean Squared Error (MSE)

$$MSE = rac{1}{N} \sum_{i=1}^{N} (y_i - \hat{\mathrm{y}}_i)^2$$

```
def MSE(y, y_pre):
    return np.mean((y - y pre) ** 2)
```

### **Advantages**

- Graph of MSE is differentiable which means it can be easily used as a loss function.
- MSE can be decomposed into variance and bias squared. This helps us understand the effect of variance or bias in data to the overall error.

$$MSE(\hat{\mathbf{y}}) = Var(\hat{\mathbf{y}}) + (Bias(\hat{\mathbf{y}}))^2$$

### \*\* Disadvantages\*\*

- The value calculated MSE has a different unit than the target variable since it is squared. (Ex. meter →
  meter²)
- If there exists outliers in the data, then they are going to result in a larger error. Therefore, MSE is not robust to outliers (this can also be an advantage if you are looking to penalize outliers).

### 3) Root Mean Squared Error (RMSE)

$$RMSE = \sqrt{rac{1}{N}\sum_{i=1}^{N}(y_i - \hat{ ext{y}}_i)^2}$$

```
def RMSE(y, y_pre):
    return np.sqrt(np.mean((y - y_pre) ** 2))
```

### **Advantages**

• The error calculated has the same unit as the target variables making the interpretation relatively easier.

### **Disadvantages**

• Just like MSE, RMSE is also susceptible to outliers.

### 4) R-Squared

$$R^{2} = 1 - \frac{\sum_{i} (\hat{y}^{(i)} - y^{(i)})^{2}}{\sum_{i} (\overline{y} - y^{(i)})^{2}}$$

$$R^{2} = 1 - \frac{(\sum_{i} (\hat{y}_{i} - y_{i})^{2}) / m}{(\sum_{i} (\hat{y}_{i} - y_{i})^{2}) / m}$$

$$= 1 - \frac{MSE(\hat{y}, y)}{Var(y)}$$

```
def R2(y, y_pre):
    u = np. sum((y - y_pre) ** 2)
    v = np. sum((y - np. mean(y)) ** 2)
    return 1 - (u / y)
```

Value closer to 1 is better.

### **4 LAB Assignment**

Now it's time to implement linear regression techniques in practice. In this lab, you will use linear regression to fit a house price model. You will use some real-world data as the test set to evaluate your model.

### **4.1 Before Assignment**

### 4.1.1 Load dataset & Import the required libraries

**Datasets**: scikit-learn provides a number of datasets which can be directly loaded by using a function. First we load a dataset as an example.

```
In [119]:
```

```
import warnings
from sklearn import datasets
boston = datasets.load_boston()
print(boston.DESCR)
```

.. boston dataset:

Boston house prices dataset

\*\*Data Set Characteristics:\*\*

:Number of Instances: 506

:Number of Attributes: 13 numeric/categorical predictive. Median Value (attribut e 14) is usually the target.

:Attribute Information (in order):

- CRIM per capita crime rate by town
- ZN proportion of residential land zoned for lots over 25,000 sq.ft.
- INDUS proportion of non-retail business acres per town
- CHAS Charles River dummy variable (= 1 if tract bounds river; 0 otherw

ise)

- NOX nitric oxides concentration (parts per 10 million)
- RM average number of rooms per dwelling
- AGE proportion of owner-occupied units built prior to 1940
   DIS weighted distances to five Boston employment centres
- RAD index of accessibility to radial highways
- TAX full-value property-tax rate per \$10,000 - PTRATIO pupil-teacher ratio by town
- B 1000 (Bk 0.63) 2 where Bk is the proportion of black people by t

own

- LSTAT % lower status of the population
- MEDV Median value of owner-occupied homes in \$1000's

:Missing Attribute Values: None

:Creator: Harrison, D. and Rubinfeld, D.L.

This is a copy of UCI ML housing dataset.

https://archive.ics.uci.edu/ml/machine-learning-databases/housing/ (https://archive.ics.uci.edu/ml/machine-learning-databases/housing/)

This dataset was taken from the StatLib library which is maintained at Carnegie Mell on University.

The Boston house-price data of Harrison, D. and Rubinfeld, D.L. 'Hedonic prices and the demand for clean air', J. Environ. Economics & Management, vol.5, 81-102, 1978. Used in Belsley, Kuh & Welsch, 'Regression diagnostics ...', Wiley, 1980. N.B. Various transformations are used in the table on pages 244-261 of the latter.

The Boston house-price data has been used in many machine learning papers that address regression problems.

.. topic:: References

- Belsley, Kuh & Welsch, 'Regression diagnostics: Identifying Influential Data and Sources of Collinearity', Wiley, 1980. 244-261.
- Quinlan, R. (1993). Combining Instance-Based and Model-Based Learning. In Procee dings on the Tenth International Conference of Machine Learning, 236-243, University of Massachusetts, Amherst. Morgan Kaufmann.

d:\miniconda3\envs\cs405\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWa rning: Function load\_boston is deprecated; `load\_boston` is deprecated in 1.0 and wi ll be removed in 1.2.

The Boston housing prices dataset has an ethical problem. You can refer to the documentation of this function for further details.

The scikit-learn maintainers therefore strongly discourage the use of this dataset unless the purpose of the code is to study and educate about ethical issues in data science and machine learning.

In this special case, you can fetch the dataset from the original source::

```
import pandas as pd
      import numpy as np
      data url = "http://lib.stat.cmu.edu/datasets/boston"
      raw_df = pd.read_csv(data_url, sep="\s+", skiprows=22, header=None)
      data = np.hstack([raw_df.values[::2, :], raw_df.values[1::2, :2]])
      target = raw_df.values[1::2, 2]
  Alternative datasets include the California housing dataset (i.e.
  :func:~sklearn.datasets.fetch_california_housing~) and the Ames housing
  dataset. You can load the datasets as follows::
      from sklearn.datasets import fetch_california_housing
      housing = fetch california housing()
  for the California housing dataset and::
      from sklearn.datasets import fetch_openml
      housing = fetch openml (name="house prices", as frame=True)
  for the Ames housing dataset.
warnings.warn(msg, category=FutureWarning)
```

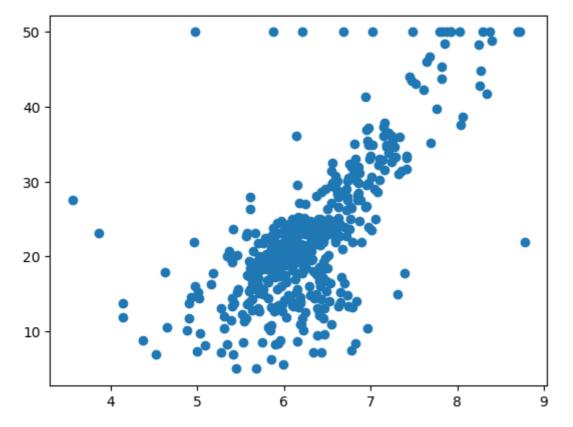
See <u>sklearn website (https://scikit-learn.org/stable/modules/classes.html#module-sklearn.datasets)</u> for details. To do this you have to import right packages and modules.

### 4.1.2 Preprocessing data

This is a small dataset containing 506 samples and 13 attributes. We need to use proper visualization methods to have an intuitive understanding. We choose the sixth attribute and draw a scattering plot to see the distribution of samples. We use *matplotlib* for data visualization.

### In [120]:

```
# Use one feature for visualization
2
   x = boston.data[:,5]
3
4
   # Get the target vector
5
   y = boston. target
6
7
   # Scattering plot of prive vs. room number
8
   from matplotlib import pyplot as plt
9
   plt.scatter(x, y)
10
   plt.show()
```



It can be seen that the samples have some exceptional distributions at the top of the plot. They may be outliers owing to some practical operation during the data input (e.g., convert any price larger than 50 into 50). However, these data are harmful to the model training, and should be removed.

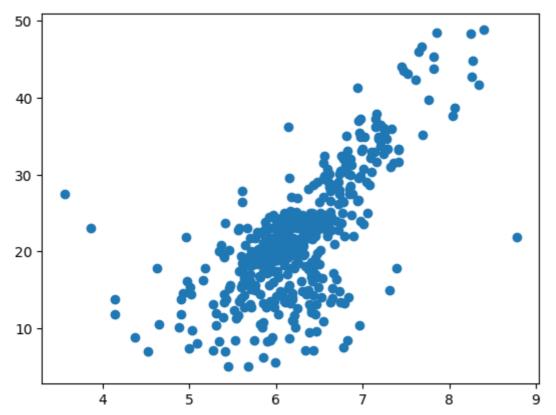
```
In [121]:
```

```
1  x = x[y<50.0]

2  y = y[y<50.0]

3  4  plt. scatter(x, y)

5  plt. show()
```



Now it can be seen that the data is nearly linear, although just in one dimension. Now we use X to denote all attributes

### In [122]:

#### Out[122]:

(490, 13)

### 4.1.3 Split data

Now we divide the whole dataset into a training set and a test set using the the scikit-learn model\_selection module.

### In [123]:

```
1 from sklearn.model_selection import train_test_split
2 X_train, X_test, y_train, y_test = train_test_split(X, y)
3 y_train.shape
```

### Out[123]:

(367,)

Usually we also use a validation set. When we use the test set for evaluation, the model will not be changed after the evaluation. However, sometime we want to optimize our model by changing its parameters according to prediction results. The solution is to split a validation set from the training set for adjusting our model. When we believe that the model is good enough, then we evaluate our model on the test set. A more rigorous and costly way is cross validation. With that method, the training set is divided into several pieces in the same size and take every piece as a validation set in turn.

#### 4.1.4 Training

### 1) Linear Regression

Now we try to implement a simple linear regression model because the dataset seems linear.

### In [124]:

```
from sklearn.linear_model import LinearRegression
lin_reg = LinearRegression()
lin_reg.fit(X_train, y_train)
```

#### Out[124]:

LinearRegression()

In a Jupyter environment, please rerun this cell to show the HTML representation or trust the notebook.

On GitHub, the HTML representation is unable to render, please try loading this page with nbviewer.org.

The model has been trained just by using a few lines of codes. Now let's make a prediction for testing

```
In [125]:

1  # Make a prediction
2  y_0_hat = lin_reg.predict(X_test[0].reshape(1,-1))
3  y_0_hat

Out[125]:
array([17.99145573])

In [126]:
1  y_test[0]

Out[126]:
14.1
```

Notice that in scikit-learn, the standard interface for machine learning is

1) instantiate a learner with super parameters or none; 2) use fit() method and feed the learner with training data; 3) use predict() for prediction.

Moreover, the data preprocessing algorithms also have the same interface, they just use transform() instead of predict().

Below are the trained parameters.

32. 00270292551771

Use the evaluation method to see if it is a good model. The score () method uses R-square.

```
In [129]:
    1 lin_reg.score(X_test, y_test)
Out[129]:
```

#### 2) Polynomial Regression

0.7595299927902152

If you have understood the concept of linear regression, you can easily implement polynomial regression.

#### 4.1.5 Evaluation model

Checking the results on test set.

### 4.2 LAB Assignment

Please use the real world dataset, **California housing price**, for model training and evaluate the model's prediction performance. You can use simple linear regression, polynomial regression or more complicated base functions such as Gaussian function or use regularization methods. Make sure at least **20% data for testing** and choose one evaluation method you think good. **Please do not just train your model and say that is good enough, you need to analyze the bias and variance**. For that end, validation or cross validation is needed. Compare the score in the training set and the validation set. If they are both good enough, then use the model on the test set.

Your test set can only be used for final evaluation!

### 4.3 Questions

1) Describe another real-world application where the regression method can be applied 2) What are the strengths of the linear/polynomial regression methods; when do they perform well? 3) What are the weaknesses of the linear/polynomial regression methods; when do they perform poorly? 4) What makes the linear regression method a good candidate for the regression problem, if you have enough knowledge about the data?

Please complete lab4 Assignment, and submit the result to bb as required (Lab02\_Assignment\_Template.ipynb)

