+ Code — + Text

## **OUTLINE**

- Simulation for Parabola of Loss Function
- RMSE
- · Confidence Intervals

Before starting, lets load the same Cars24 dataset we can work with

	selling_price	year	km_driven	mileage	engine	max_power	make	model	transmission_type	seats_coupe	seats_family
0	1.20	2012.0	120000	19.70	796.0	46.30	Maruti	Alto Std	1	0	1
1	5.50	2016.0	20000	18.90	1197.0	82.00	Hyundai	Grand i10 Asta	1	0	1
2	2.15	2010.0	60000	17.00	1197.0	80.00	Hyundai	i20 Asta	1	0	1
3	2.26	2012.0	37000	20.92	998.0	67.10	Maruti	Alto K10 2010- 2014 VXI	1	0	1
								Ecosport 2015-			

```
# define X and y
X = df["max_power"].values
Y = df["selling price"].values
```

```
# standardize the data
u = X.mean()
std = X.std()
X = (X-u)/std
```

## ▼ Cost Function Visualisation

How does the "cost function" surface look for univariate linear regression?

To understand this, lets take different possible values of W0 and W1

- For a set of W0 and W1 values, we can find MSE np.sum((Y-Y\_hat)\*\*2)/Y.shape[0]
- · We can plot MSE for possible values of
  - W0=[-100,100],

- W1=[-100,100]
- with step-size=1

How can we generate different combinations of W0 and W1?

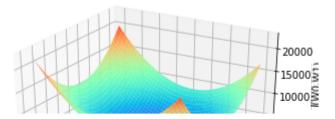
- We will use np.meshgrid(W0,W1)
- It will return two matrices
  - 1. Contains X-coordinate (W0)
  - 2. Contains Y-coordinate (W1)

You can add a print to statement to undersrtand the output of meshgrid

```
W0 = np.arange(-100,100,1)
W1 = np.arange(-100,100,1)

W0,W1 = np.meshgrid(W0,W1)
J = np.zeros(W0.shape)
for i in range(J.shape[0]):
    for j in range(J.shape[1]):
        Y_hat = W1[i,j]*X + W0[i,j]
        J[i,j] = np.sum((Y-Y_hat)**2)/Y.shape[0]

fig = plt.figure()
axes = fig.gca(projection='3d')
axes.plot_surface(W0,W1,J, cmap="rainbow")
axes.set_xlabel("W0")
axes.set_zlabel("W1")
axes.set_zlabel("J(W0,W1)")
plt.show()
```



## ▼ Root Mean Square Error (RMSE)

• 
$$RMSE = \sqrt{\sum_{i=1}^{i=n} (y_i - y_i^-) \div n}$$

- ullet  $y_i$  is the actual value
- $ullet \ y_i^-$  is the predicted value
- n is the number of samples
- ANALOGOUS TO -
  - $\circ$  Euclidean Distance:  $dist(x,y) = \sqrt{\sum_{i=1}^n (x_i y_i)^2}$
  - $\circ~$  RMSE can be thought of as some kind of (normalized) distance between the vector of  $y_i^-$  and the vector of  $y_i$ .
- ▼ But how "confident" are we about the estimated line?

We calculate bootstraped estimates to get confidence intervals for our parameters

## **Code Recap**

```
# model = w0 + x1w1
def predict(x,weight):
    y_hat = weight[0] + weight[1]*x #hypothesis
    return y_hat
#MSE
```

```
def error(X,Y,weight):
    m = X.shape[0]
   total_error = 0.0
   for i in range(m):
       y hat = predict(X[i],weight)
       total error += (y hat - Y[i])**2
    return (total error/m)
def gradient(X,Y,weight):
    m = X.shape[0]
    # print(m)
    grad = np.zeros((2,)) #initializing gradient from 0
    # print(grad)
    for i in range(200): #iterate over all points
        x = X[i]
       y hat = predict(x,weight)
       y = Y[i]
        grad[0] += -2*(v - v hat)
        grad[1] += -2*(y - y hat)*x
        # print(x, y)
        # print(2*(y - y hat), -2*(y - y hat)*x)
        # print(grad)
    return grad/m
def gradient descent(X,Y, epochs=800,learning rate =0.1):
    weight = np.zeros((2,))
    error_list = []
    weight list = []
    for i in range(epochs):
        # Compute grad
        grad = gradient(X,Y,weight)
        # print(grad)
        e = error(X,Y,weight)
        #Update weights
```

```
weight[0] = weight[0] - learning_rate*grad[0]
weight[1] = weight[1] - learning_rate*grad[1]
# Storing the weight values during updates
weight_list.append((weight[0],weight[1]))
error_list.append(e)

return weight,error_list,weight_list

weight, error_list, weight_list = gradient_descent(X,Y,epochs=800)
```

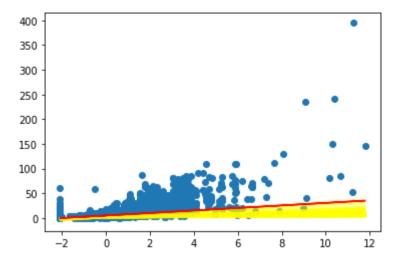
▼ Lets calculate Confidence Intervals by bootstrapping technique we learnt in P&S module

Recall that we calculated bootstraped estimates to get confidence intervals for our parameters

NOTE: Please do not run this cell below, it will take a lot of time before it shows results.

```
n_reps = 1000
bs_weights = []
for i in range(n_reps):
    inds = np.arange(len(X)) #setup an array of indices
    bs_inds = np.random.choice(inds, size=len(inds)) # bootstrap the indices
    X_bs, Y_bs = X[bs_inds], Y[bs_inds] # generate the bootstrapped sample
    bs_weight, _, _ = gradient_descent(X_bs, Y_bs, epochs=50) # perform gradient descent
    bs_weights.append(bs_weight)

plt.figure()
Y_hat = predict(X,weight)
plt.scatter(X,Y)
for i in range(n_reps):
    plt.plot(X, predict(X, bs_weights[i]), alpha=0.1, color='yellow')
plt.plot(X,Y_hat,color='red')
plt.show()
```



So, how do I get the estimates if boorstrapping technique takes so much time?

Thanks to Central Limit Theorem, we can calculate it using simple formula.

We will discuss this in one of the future classes soon.

This is a great chance to revise CLT, if in case, you have forgotten.

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