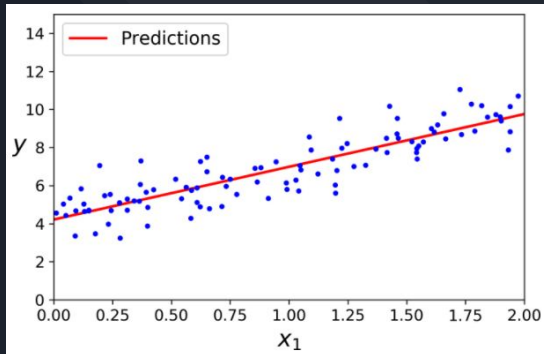


# Machine Learning: Supervised Learning - Linear Regression using Normal Equation



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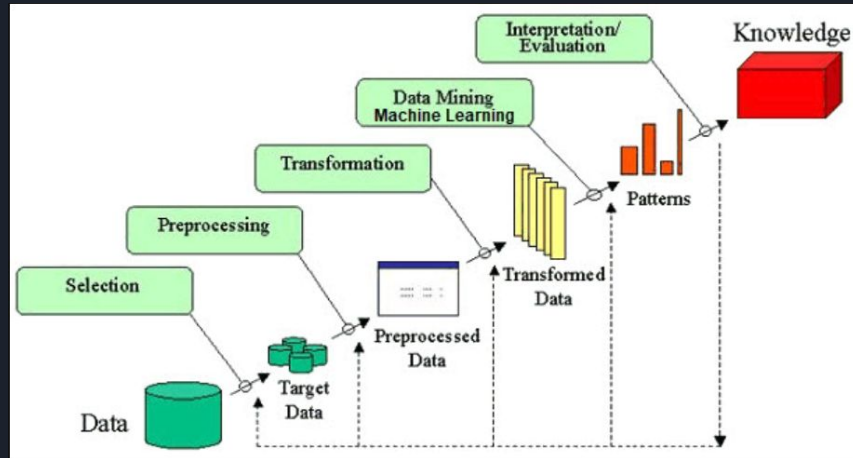


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# Linear Regression using Normal Equation

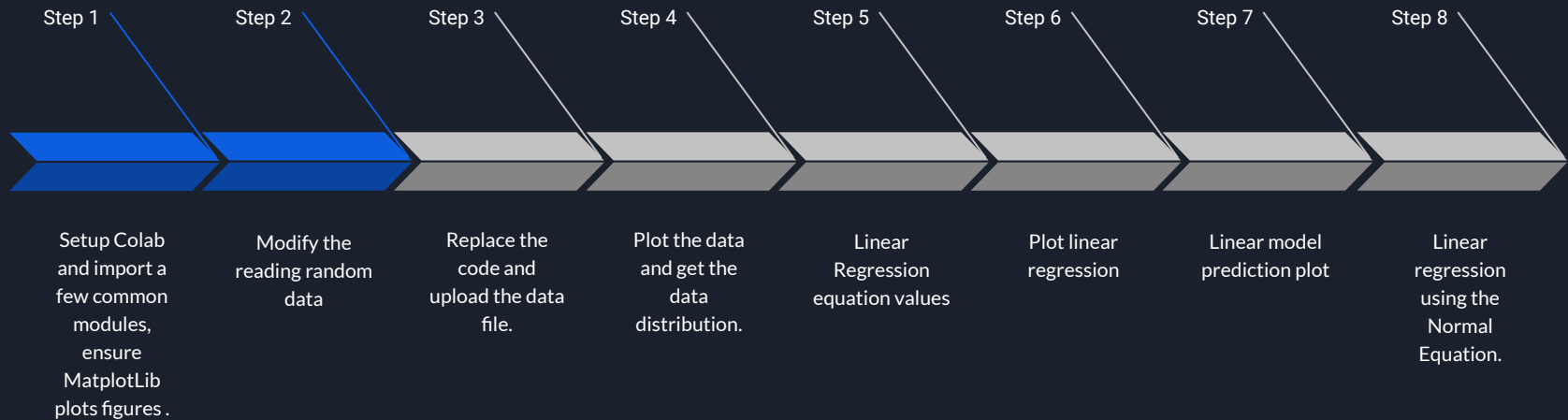
- Linear regression is one of the most important and popular predictive techniques in data analysis. Normal Equation is an analytical approach to Linear Regression with a Least Square Cost Function.
- A regression is a statistical analysis assessing the association between two variables. It is used to find the relationship between two variables.
- In this project we are performing everything except preprocessing data.



# Project Description

- Follow the procedure mentioned in Training Linear Models and make work in Colab.
- Save the [abalone\\_train.csv](#) to a local drive and upload in step 3.

```
names=["Length", "Diameter", "Height", "Whole weight", "Shucked weight", "Viscera weight", "Shell weight", "Age"]
```





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# Implementation

## Step 1:

Setup Colab and import a few common modules, ensure Matplotlib plots figures inline and prepare a function to save the figures.

```
# Python ≥3.5 is required
import sys
assert sys.version_info >= (3, 5)

# Scikit-Learn ≥0.20 is required
import sklearn
assert sklearn.__version__ >= "0.20"

# Common imports
import numpy as np
import os

# to make this notebook's output stable across runs
np.random.seed(42)

# To plot pretty figures
%matplotlib inline
import matplotlib as mpl
import matplotlib.pyplot as plt
mpl.rc('axes', labelsiz=14)
mpl.rc('xtick', labelsiz=12)
mpl.rc('ytick', labelsiz=12)

# Where to save the figures
PROJECT_ROOT_DIR = "."
CHAPTER_ID = "training_linear_models"
IMAGES_PATH = os.path.join(PROJECT_ROOT_DIR, "images", CHAPTER_ID)
os.makedirs(IMAGES_PATH, exist_ok=True)

def save_fig(fig_id, tight_layout=True, fig_extension="png", resolution=300):
    path = os.path.join(IMAGES_PATH, fig_id + "." + fig_extension)
    print("Saving figure", fig_id)
    if tight_layout:
        plt.tight_layout()
    plt.savefig(path, format=fig_extension, dpi=resolution)
```

# Implementation

## Step 2:

Modify the reading random data

```
import numpy as np

X = 2 * np.random.rand(100, 1)
y = 4 + 3 * X + np.random.randn(100, 1)
```

## Step 3:

Replace the code and upload the data file.

```
import numpy as np
import pandas as pd

# X = 2 * np.random.rand(100, 1)
# y = 4 + 3 * X + np.random.randn(100, 1)
from google.colab import files
uploaded = files.upload()

import io
abalone = pd.read_csv(
    io.BytesIO(uploaded['abalone_train.csv']),
    names=["Length", "Diameter", "Height", "Whole weight", "Shucked weight",
           "Viscera weight", "Shell weight", "Age"])

# X1 is
#      0      0.435
#      1      0.585
#      2      0.655
#      ....
X1 = abalone["Length"]

# X2 is
#      array([0.435, 0.585, ..., 0.45])
X2 = np.array(X1)

# X is
#      array([[0.435],
#             [0.585],
#             [0.655],
#             ...,
#             [0.53 ],
#             [0.395],
#             [0.45 ]])
X = X2.reshape(-1, 1)

y1 = abalone["Height"]
y2 = np.array(y1)
y = y2.reshape(-1, 1)
```

Choose Files abalone\_train.csv  
• abalone\_train.csv(text/csv) - 145915 bytes, last modified: 5/26/2021 - 100% done  
Saving abalone\_train.csv to abalone\_train.csv

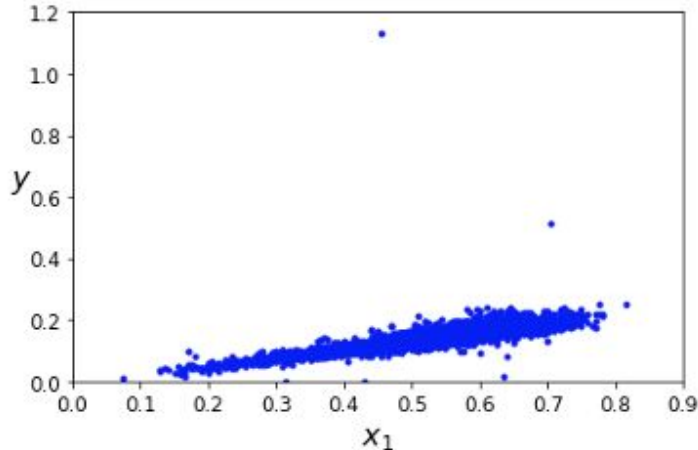
# Implementation

## Step 4:

Plot the data and get the data distribution.

```
plt.plot(X, y, "b.")  
plt.xlabel("$x_1$", fontsize=18)  
plt.ylabel("$y$", rotation=0, fontsize=18)  
plt.axis([0, 0.9, 0, 1.2])  
save_fig("generated_data_plot")  
plt.show()
```

Saving figure generated\_data\_plot



## Step 5:

Linear Regression equation values.

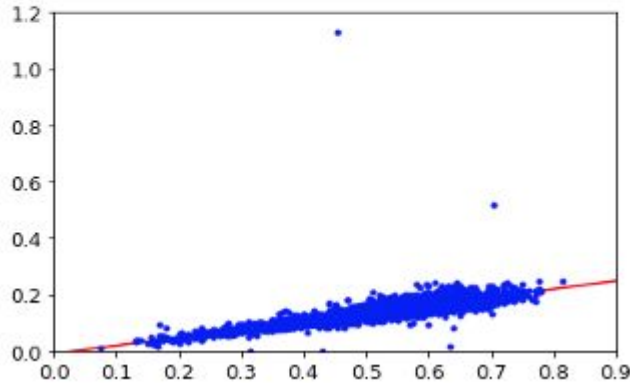
```
[26] X_b = np.c_[np.ones((3320, 1)), X] # add x0 = 1 to each instance  
theta_best = np.linalg.inv(X_b.T.dot(X_b)).dot(X_b.T).dot(y)  
  
[27] theta_best  
  
array([[ -0.0108267 ],  
       [ 0.28716253]])  
  
[28] X_new = np.array([[0], [2]])  
X_new_b = np.c_[np.ones((2, 1)), X_new] # add x0 = 1 to each instance  
y_predict = X_new_b.dot(theta_best)  
  
array([[ -0.0108267 ],  
       [ 0.56349837]])
```



# Implementation

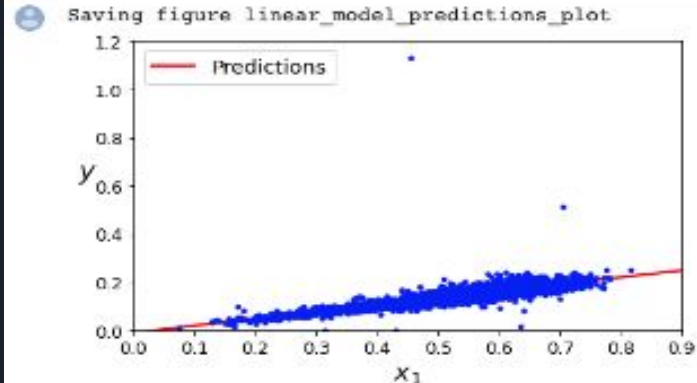
Step 6:  
Plot linear regression.

```
plt.plot(X_new, y_predict, "r-")  
plt.plot(X, y, "b.")  
plt.axis([0, 0.9, 0, 1.2])  
plt.show()
```



Step 7:  
Linear model prediction plot.

```
plt.plot(X_new, y_predict, "r-", linewidth=2, label="Predictions")  
plt.plot(X, y, "b.")  
plt.xlabel("$x_1$", fontsize=18)  
plt.ylabel("$y$", rotation=0, fontsize=18)  
plt.legend(loc="upper left", fontsize=14)  
plt.axis([0, 0.9, 0, 1.2])  
save_fig("linear_model_predictions_plot")  
plt.show()
```



# Implementation

## Step 8:

Linear regression using the Normal Equation.

```
from sklearn.linear_model import LinearRegression

lin_reg = LinearRegression()
lin_reg.fit(X, y)
lin_reg.intercept_, lin_reg.coef_

(array([-0.0108267]), array([[0.28716253]]))
```

```
[33] lin_reg.predict(X_new)

array([[ -0.0108267 ],
       [ 0.56349837]])
```

The `LinearRegression` class is based on the `scipy.linalg.lstsq()` function (the name stands for "least squares"), which you could call directly:

```
[34] theta_best_svd, residuals, rank, s = np.linalg.lstsq(X_b, y, rcond=1e-6)
theta_best_svd

array([[ -0.0108267 ],
       [ 0.28716253]])
```

This function computes  $\mathbf{X}^+ \mathbf{y}$ , where  $\mathbf{X}^+$  is the *pseudoinverse* of  $\mathbf{X}$  (specifically the Moore-Penrose inverse). You can use `np.linalg.pinv()` to compute the pseudoinverse directly:

```
np.linalg.pinv(X_b).dot(y)

array([[ -0.0108267 ],
       [ 0.28716253]])
```



## Bibliography/References

- <https://www.geeksforgeeks.org/ml-normal-equation-in-linear-regression/>
- <https://towardsdatascience.com/performing-linear-regression-using-the-normal-equation-6372ed3c57>
- [https://colab.research.google.com/github/ageron/handson-ml2/blob/master/04\\_training\\_linear\\_models.ipynb#scrollTo=QnCG6urINAUM](https://colab.research.google.com/github/ageron/handson-ml2/blob/master/04_training_linear_models.ipynb#scrollTo=QnCG6urINAUM)

ThankYou