- 1. Follow the procedure mentioned in <u>Chapter 4 Training Linear Models</u> to make it work on Colab.
- 2. Save the abalone train.cvs to a local drive
 - Note: the abalone_train.cvs has this format

- 3. Change the process mentioned in <u>Step 1</u> by <u>reading CVS test data from a local drive</u>: <u>abalone_train.cvs</u>
 - Process
 - a. You can modify the code in Linear regression using the Normal Equation. Instead of reading random data

You need to read data from a local drive and transform the data to fit the <u>Python</u> code.

```
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)
```

Answer:

Go to colab

→ Setup

First, let's import a few common modules, ensure MatplotLib plots figures inline and prepare ϵ Python 3.5 or later is installed (although Python 2.x may work, it is deprecated so we strongly Scikit-Learn \geq 0.20.

```
# 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', labelsize=14)
    mpl.rc('xtick', labelsize=12)
    mpl.rc('ytick', labelsize=12)
    # Where to save the figures
    PROJECT_ROOT_DIR = "."
```

▼ Linear regression using the Normal Equation

```
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
          0.585
   1
           0.655
X1 = abalone["Length"]
# array([0.435, 0.585, ...., 0.45])
X2 = np.array(X1)
# X is
   array([[0.435],
        [0.585],
           [0.655],
```

Upload file

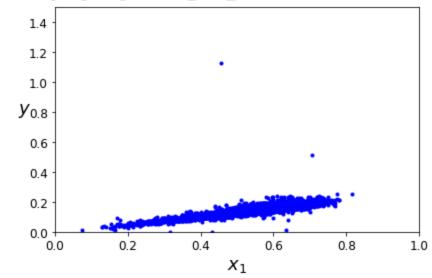
```
# X2 is
   # array([0.435, 0.585, ...., 0.45])
   X2 = np.array(X1)
   # X is
    # array([[0.435],
           [0.585],
   #
             [0.655],
   #
             ...,
   #
             [0.53],
             [0.395],
             [0.45]])
   X = X2.reshape(-1, 1)
   y1 = abalone["Height"]
   y2 = np.array(y1)
   y = y2.reshape(-1, 1)
Choose Files abalone_train.csv
```

• abalone_train.csv(application/vnd.ms-excel) - 149229 bytes, last modified: 6/2/2021 - 100% done Saving abalone_train.csv to abalone_train.csv

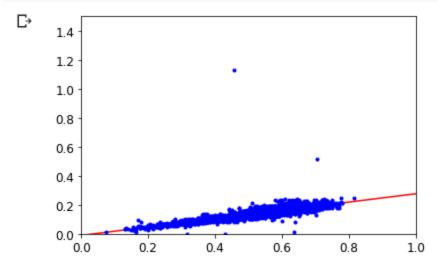
The result

```
plt.plot(X, y, "b.")
plt.xlabel("$x_1$", fontsize=18)
plt.ylabel("$y$", rotation=0, fontsize=18)
plt.axis([0, 1, 0, 1.5])
save_fig("generated_data_plot")
plt.show()
```

Saving figure generated_data_plot

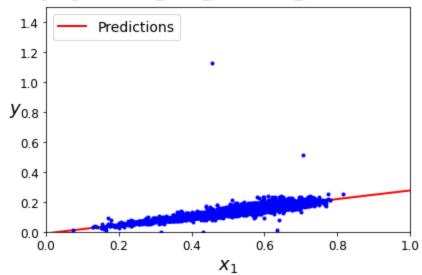


```
plt.plot(X_new, y_predict, "r-")
plt.plot(X, y, "b.")
plt.axis([0, 1, 0, 1.5])
plt.show()
```



```
[14] 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, 1, 0, 1.5])
    save_fig("linear_model_predictions_plot")
    plt.show()
```

Saving figure linear_model_predictions_plot



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

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: