

Source Code with Explanation-

```
#Importing the pandas for data processing and numpy for numerical
computing import numpy as np import pandas as pd
# Importing the Boston Housing dataset from the sklearn
from sklearn.datasets import load_boston boston =
load_boston() #Converting the data into pandas
dataframe data = pd.DataFrame(boston.data) #First look
at the data data.head()
```

	0	1	2	3	4	5	6	7	8	9	10	11	12
0	0.00632	18.0	2.31	0.0	0.538	6.575	65.2	4.0900	1.0	296.0	15.3	396.90	4.98
1	0.02731	0.0	7.07	0.0	0.469	6.421	78.9	4.9671	2.0	242.0	17.8	396.90	9.14
2	0.02729	0.0	7.07	0.0	0.469	7.185	61.1	4.9671	2.0	242.0	17.8	392.83	4.03
3	0.03237	0.0	2.18	0.0	0.458	6.998	45.8	6.0622	3.0	222.0	18.7	394.63	2.94
4	0.06905	0.0	2.18	0.0	0.458	7.147	54.2	6.0622	3.0	222.0	18.7	396.90	5.33

```
#Adding the feature names to the dataframe
data.columns = boston.feature_names #Adding the
target variable to the dataset data['PRICE'] =
boston.target #Looking at the data with names and
target variable data.head(n=10)
```

	CRIM	ZN	INDUS	CHAS	NOX	RM	AGE	DIS	RAD	TAX	PTRATIO	B	LSTAT	PRICE
0	0.00632	18.0	2.31	0.0	0.538	6.575	65.2	4.0900	1.0	296.0	15.3	396.90	4.98	24.0
1	0.02731	0.0	7.07	0.0	0.469	6.421	78.9	4.9671	2.0	242.0	17.8	396.90	9.14	21.6
2	0.02729	0.0	7.07	0.0	0.469	7.185	61.1	4.9671	2.0	242.0	17.8	392.83	4.03	34.7
3	0.03237	0.0	2.18	0.0	0.458	6.998	45.8	6.0622	3.0	222.0	18.7	394.63	2.94	33.4
4	0.06905	0.0	2.18	0.0	0.458	7.147	54.2	6.0622	3.0	222.0	18.7	396.90	5.33	36.2
5	0.02985	0.0	2.18	0.0	0.458	6.430	58.7	6.0622	3.0	222.0	18.7	394.12	5.21	28.7
6	0.08829	12.5	7.87	0.0	0.524	6.012	66.6	5.5605	5.0	311.0	15.2	395.60	12.43	22.9
7	0.14455	12.5	7.87	0.0	0.524	6.172	96.1	5.9505	5.0	311.0	15.2	396.90	19.15	27.1
8	0.21124	12.5	7.87	0.0	0.524	5.631	100.0	6.0821	5.0	311.0	15.2	386.63	29.93	16.5
9	0.17004	12.5	7.87	0.0	0.524	6.004	85.9	6.5921	5.0	311.0	15.2	386.71	17.10	18.9

```
#Shape of the data print(data.shape)
#Checking the null values in the dataset
data.isnull().sum() CRIM          0 ZN
0 INDUS          0 CHAS          0 NOX          0
RM          0 AGE          0 DIS          0
RAD          0 TAX          0 PTRATIO          0 B
0 LSTAT          0 PRICE          0 dtype: int64
#Checking the statistics of the data
data.describe()

# This is sometimes very useful, for example if you look at the CRIM the max is
88.97 and 75% of the value is below 3.677083 and
# mean is 3.613524 so it means the max values is actually an outlier or there are
outliers present in the column
```

	CRIM	ZN	INDUS	CHAS	NOX	RM	AGE	DIS	RAD	TAX	PTRATIO
count	506.000000	506.000000	506.000000	506.000000	506.000000	506.000000	506.000000	506.000000	506.000000	506.000000	506.000000
mean	3.613524	11.363636	11.136779	0.069170	0.554695	6.284634	68.574901	3.795043	9.549407	408.237154	18.455534
std	8.601545	23.322453	6.860353	0.253994	0.115878	0.702617	28.148861	2.105710	8.707259	168.537116	2.164946
min	0.006320	0.000000	0.460000	0.000000	0.385000	3.561000	2.900000	1.129600	1.000000	187.000000	12.600000
25%	0.082045	0.000000	5.190000	0.000000	0.449000	5.885500	45.025000	2.100175	4.000000	279.000000	17.400000
50%	0.256510	0.000000	9.690000	0.000000	0.538000	6.208500	77.500000	3.207450	5.000000	330.000000	19.050000
75%	3.677083	12.500000	18.100000	0.000000	0.624000	6.623500	94.075000	5.188425	24.000000	666.000000	20.200000
max	88.976200	100.000000	27.740000	1.000000	0.871000	8.780000	100.000000	12.126500	24.000000	711.000000	22.000000

```
data.info() <class
'pandas.core.frame.DataFrame'>
RangeIndex: 506 entries, 0 to 505
Data columns (total 14 columns):
#   Column      Non-Null Count  Dtype -
--  -
0    CRIM        506 non-null    float64
1    ZN          506 non-null    float64
2    INDUS       506 non-null    float64
3    CHAS        506 non-null    float64
```

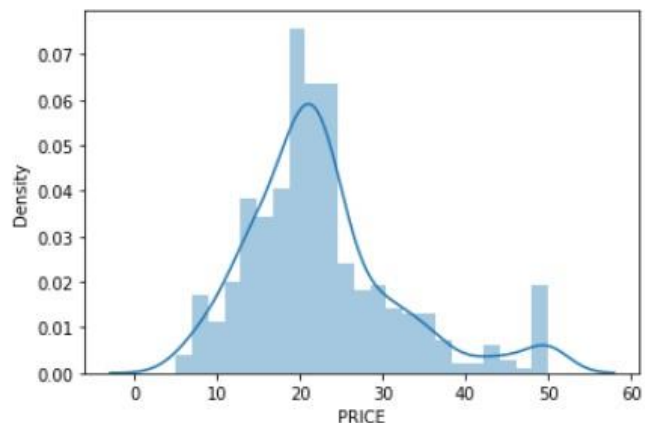
```
4   NOX      506 non-null   float64
5   RM       506 non-null   float64
6   AGE      506 non-null   float64
7   DIS      506 non-null   float64
8   RAD      506 non-null   float64
9   TAX      506 non-null   float64
10  PTRATIO  506 non-null   float64
11  B        506 non-null   float64
12  LSTAT    506 non-null   float64
13  PRICE    506 non-null   float64
```

```
dtypes: float64(14) memory usage: 55.5
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```
#checking the distribution of the target variable import seaborn as sns
sns.distplot(data.PRICE) #The distribution seems normal, has not be the data
normal we would have perform log transformation or took to square root of the
data to make the data normal. # Normal distribution is need for the machine
learning for better predictiblity of the model
```

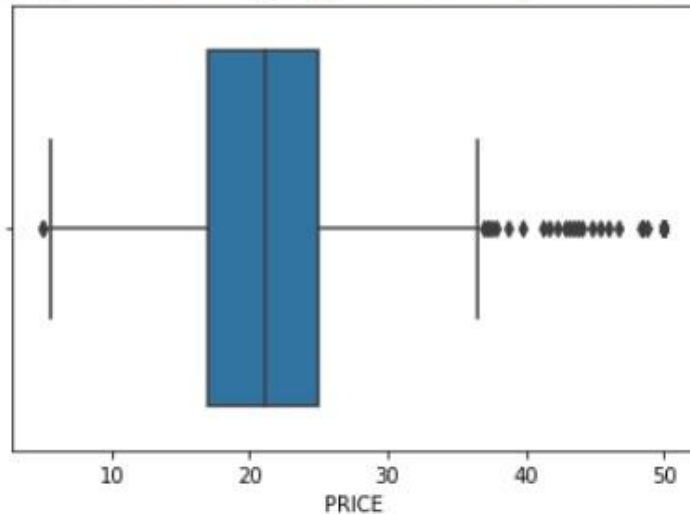
```
<matplotlib.axes._subplots.AxesSubplot at 0x7f44d082c670>
```



```
#Distribution using box plot
```

```
sns.boxplot(data.PRICE)
```

<matplotlib.axes._subplots.AxesSubplot at 0x7f44d077ed60>



#Checking the correlation of the independent feature with the dependent feature
 # Correlation is a statistical technique that can show whether and how strongly pairs of variables are related. An intelligent correlation analysis can lead to a greater understanding of your data #checking Correlation of the data correlation

```
= data.corr() correlation.loc['PRICE']
```

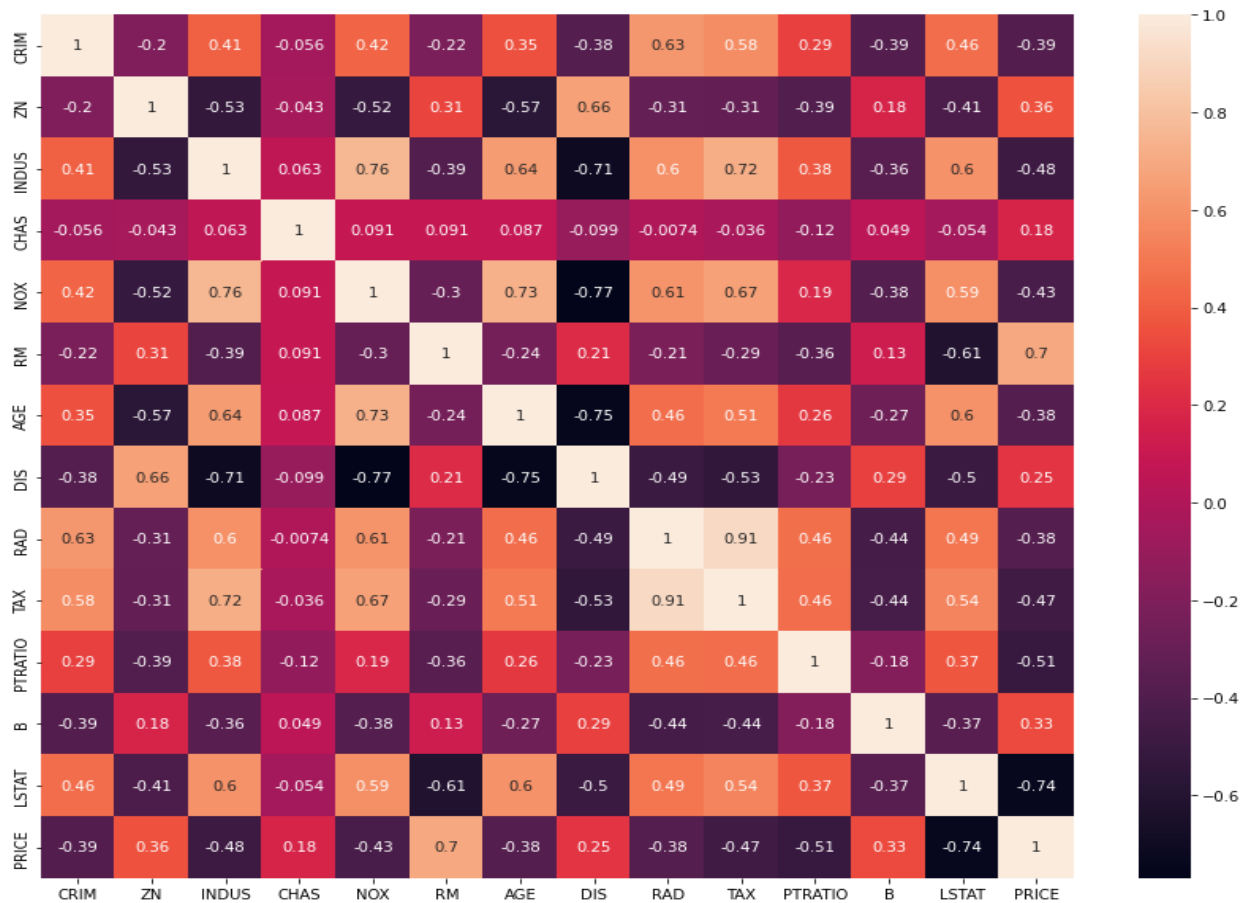
```
CRIM      -0.388305
ZN        0.360445
INDUS     -0.483725
CHAS      0.175260
NOX       -0.427321
RM        0.695360
AGE       -0.376955
DIS        0.249929
RAD       -0.381626
TAX       -0.468536
PTRATIO   -0.507787
B         0.333461
LSTAT     -0.737663
PRICE     1.000000
```

Name: PRICE, dtype: float64 # plotting the heatmap

```
import matplotlib.pyplot as plt fig, axes =
plt.subplots(figsize=(15,12))
```

```
sns.heatmap(correlation, square = True, annot = True)
```

By looking at the correlation plot LSTAT is negatively correlated with -0.75 and RM is positively correlated to the price and PTRATIO is correlated negatively with -0.51



Checking the scatter plot with the most correlated features

```
plt.figure(figsize = (20,5)) features =
['LSTAT', 'RM', 'PTRATIO'] for i, col in enumerate(features):
    plt.subplot(1, len(features) , i+1)
    x = data[col] y = data.PRICE
    plt.scatter(x, y, marker='o')
    plt.title("Variation in House prices")
    plt.xlabel(col)
    plt.ylabel('"House prices in $1000"')
```



```
# Splitting the dependent feature and independent feature
#X = data[['LSTAT','RM','PTRATIO']] X = data.iloc[:, :-1]
y= data.PRICE

# In order to provide a standardized input to our neural network, we need the
perform the normalization of our dataset. # This can be seen as an step to
reduce the differences in scale that may arise from the existent features. # We
perform this normalization by subtracting the mean from our data and dividing
it by the standard deviation. # One more time, this normalization should only
be performed by using the mean and standard deviation from the training set, #
in order to avoid any information leak from the test set.

mean = X_train.mean(axis=0)
std = X_train.std(axis=0)

X_train = (X_train - mean) / std
X_test = (X_test - mean) / std
#Linear Regression
from sklearn.linear_model import LinearRegression
regressor = LinearRegression()
#Fitting the model
regressor.fit(X_train,y_train)
# Model Evaluation

#Prediction on the test dataset y_pred =
regressor.predict(X_test) # Predicting RMSE the Test
set results from sklearn.metrics import
mean_squared_error rmse =
(np.sqrt(mean_squared_error(y_test, y_pred)))
print(rmse)

from sklearn.metrics import r2_score
r2 = r2_score(y_test, y_pred)
print(r2) # Neural Networks #Scaling
the dataset
from sklearn.preprocessing import StandardScaler
sc = StandardScaler() X_train =
sc.fit_transform(X_train)
X_test = sc.transform(X_test) # Due to the small amount of presented data in this
dataset, we must be careful to not create an overly complex model, # which could
lead to overfitting our data. For this, we are going to adopt an architecture
```


based on two Dense layers, # the first with 128 and the second with 64 neurons, both using a ReLU activation function. # A dense layer with a linear activation will be used as output layer. # In order to allow us to know if our model is properly learning, we will use a mean squared error loss function and to report the performance of it we will adopt the mean average error metric. # By using the summary method from Keras, we can see that we have a total of 10,113 parameters, which is acceptable for us.

```
#Creating the neural network model
import keras
from keras.layers import Dense, Activation, Dropout
from keras.models import Sequential

model = Sequential()
model.add(Dense(128, activation = 'relu', input_dim = 13))
model.add(Dense(64, activation = 'relu')) model.add(Dense(32, activation = 'relu'))
model.add(Dense(16, activation = 'relu')) model.add(Dense(1))
#model.compile(optimizer='adam', loss='mse', metrics=['mae'])
model.compile(optimizer = 'adam', loss = 'mean_squared_error', metrics=['mae'])
!pip install ann_visualizer
!pip install graphviz
from ann_visualizer.visualize import ann_viz;
#Build your model here
ann_viz(model, title="DEMO ANN");
history = model.fit(X_train, y_train, epochs=100, validation_split=0.05)
# By plotting both loss and mean average error, we can see that our model was
capable of learning patterns in our data without overfitting taking place (as
shown by the validation set curves) from plotly.subplots import make_subplots
import plotly.graph_objects as go

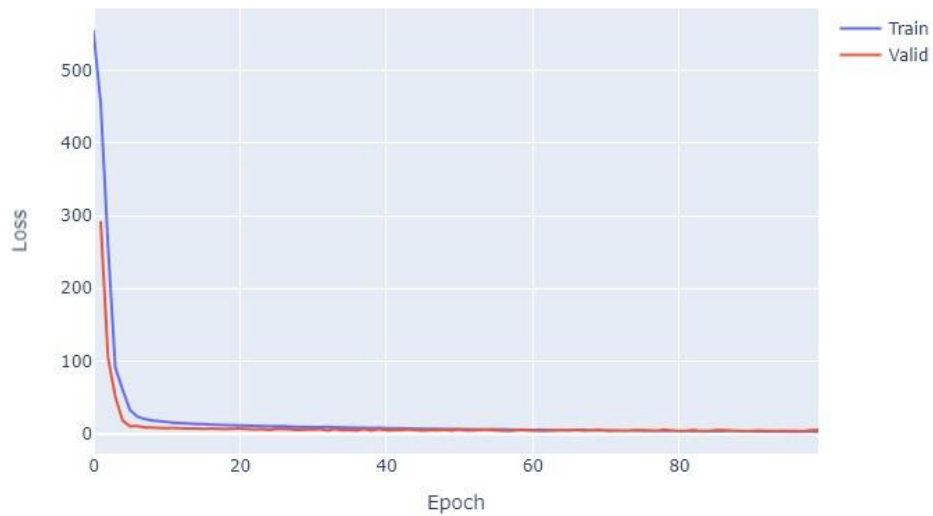
fig = go.Figure()
fig.add_trace(go.Scattergl(y=history.history['loss'],
                           name='Train'))

fig.add_trace(go.Scattergl(y=history.history['val_loss'],
                           name='Valid'))

fig.update_layout(height=500, width=700,
```

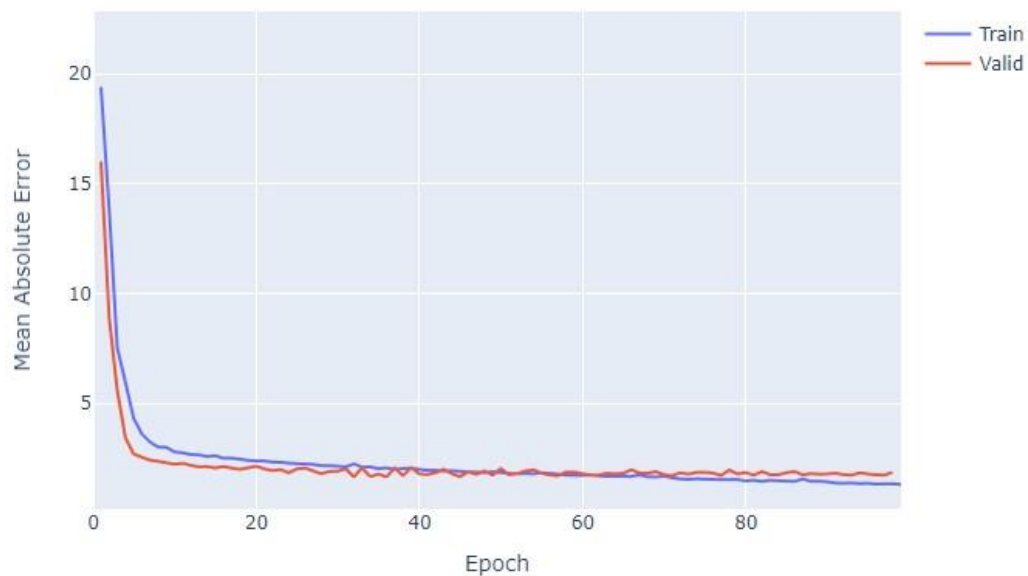
```
axis_title='Epoch',  
axis_title='Loss')
```

```
fig.show()
```



```
fig = go.Figure()  
fig.add_trace(go.Scattergl(y=history.history['mae'],  
                           name='Train'))  
  
fig.add_trace(go.Scattergl(y=history.history['val_mae'],  
                           name='Valid'))
```

```
fig.update_layout(height=500, width=700,  
                  axis_title='Epoch',  
                  axis_title='Mean Absolute Error')  
fig.show()
```




```

#Evaluation of the model
y_pred = model.predict(X_test)
mse_nn, mae_nn = model.evaluate(X_test, y_test)

print('Mean squared error on test data: ', mse_nn)
print('Mean absolute error on test data: ', mae_nn)
4/4 [=====] - 0s 4ms/step - loss: 10.5717 - mae: 2.2670
Mean squared error on test data: 10.571733474731445
Mean absolute error on test data: 2.2669904232025146
#Comparison with traditional approaches #First let's try with a
simple algorithm, the Linear Regression: from sklearn.metrics
import mean_absolute_error

lr_model = LinearRegression()
lr_model.fit(X_train, y_train)
y_pred_lr = lr_model.predict(X_test)
mse_lr = mean_squared_error(y_test,
y_pred_lr) mae_lr =
mean_absolute_error(y_test,
y_pred_lr)

print('Mean squared error on test data: ', mse_lr)
print('Mean absolute error on test data: ', mae_lr)
from sklearn.metrics import r2_score r2 =
r2_score(y_test, y_pred) print(r2)
0.8812832788381159
# Predicting RMSE the Test set results from
sklearn.metrics import mean_squared_error rmse =
(np.sqrt(mean_squared_error(y_test, y_pred)))
print(rmse) 3.320768607496587
# Make predictions on new data import sklearn new_data =
sklearn.preprocessing.StandardScaler().fit_transform([[0.1, 10.0,
5.0, 0, 0.4, 6.0, 50, 6.0, 1, 400, 20, 300, 10]])
prediction = model.predict(new_data)
print("Predicted house price:", prediction) 1/1
[=====] - 0s 70ms/step
Predicted house price:
[[11.104753]]

#new_data      =

sklearn.preprocessing.StandardScaler().fit_transform([[0.1,      10.0,

```

`5.0, 0, 0.4, 6.0, 50, 6.0, 1, 400, 20, 300, 10]]))` is a line of code that standardizes the input features of a new data point.

In this specific case, we have a new data point represented as a list of 13 numeric values (`[0.1, 10.0, 5.0, 0, 0.4, 6.0, 50, 6.0, 1, 400, 20, 300, 10]`) that represents the values for the 13 features of the Boston House Price dataset.

The `StandardScaler()` function from the `sklearn.preprocessing` module is used to standardize the data. Standardization scales each feature to have zero mean and unit variance, which is a common preprocessing step in machine learning to ensure that all features contribute equally to the model.

The `fit_transform()` method is used to fit the scaler to the data and apply the standardization transformation. The result is a new data point with standardized feature values.