# Lab 2 (02-01-2024)

This lab experiments help you master how to perform basic data analysis and preliminary data visualization.

```
In [ ]: Registration_Number = "22011103010"
Name = "Deepthi"

# Python Program to Get IP Address
import socket
hostname = socket.gethostname()
IPAddr = socket.gethostbyname(hostname)

print("My name is " + Name + " and my roll no : " + Registration_Number)
print("Computer IP Address is: " + IPAddr)
```

My name is Deepthi and my roll no : 22011103010 Computer IP Address is: 10.123.163.220

## **Experiment 1**

Load iris dataset using scikit learn.

```
In [ ]: # Experiment 1
    from sklearn import datasets
    import matplotlib.pyplot as plt
    import numpy as np

    iris = datasets.load_iris()
    X = np.array(iris["data"])
    Y = np.array(iris["target"])
    category_names = iris["target_names"]
    feature_names = iris["feature_names"]
In [ ]: # Description of the IRIS Dataset
    print(iris["DESCR"])
```

.. \_iris\_dataset:

Iris plants dataset

\_\_\_\_\_\_

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

:Number of Instances: 150 (50 in each of three classes)
:Number of Attributes: 4 numeric, predictive attributes and the class
:Attribute Information:

- sepal length in cm
- sepal width in cm
- petal length in cm
- petal width in cm
- class:
  - Iris-Setosa
  - Iris-Versicolour
  - Iris-Virginica

### :Summary Statistics:

=========	====	====	======	=====	=======	=======
	Min	Max	Mean	SD	Class Cor	relation
=========	====	====	======	=====	=======	=======
sepal length:	4.3	7.9	5.84	0.83	0.7826	
sepal width:	2.0	4.4	3.05	0.43	-0.4194	
petal length:	1.0	6.9	3.76	1.76	0.9490	(high!)
petal width:	0.1	2.5	1.20	0.76	0.9565	(high!)
=========	====	====	======	=====	=======	=======

:Missing Attribute Values: None

:Class Distribution: 33.3% for each of 3 classes.

:Creator: R.A. Fisher

:Donor: Michael Marshall (MARSHALL%PLU@io.arc.nasa.gov)

:Date: July, 1988

The famous Iris database, first used by Sir R.A. Fisher. The dataset is taken from Fisher's paper. Note that it's the same as in R, but not as in the UCI Machine Learning Repository, which has two wrong data points.

This is perhaps the best known database to be found in the pattern recognition literature. Fisher's paper is a classic in the field and is referenced frequently to this day. (See Duda & Hart, for example.) The data set contains 3 classes of 50 instances each, where each class refers to a type of iris plant. One class is linearly separable from the other 2; the latter are NOT linearly separable from each other.

|details-start|
\*\*References\*\*
|details-split|

- Fisher, R.A. "The use of multiple measurements in taxonomic problems" Annual Eugenics, 7, Part II, 179-188 (1936); also in "Contributions to Mathematical Statistics" (John Wiley, NY, 1950).
- Duda, R.O., & Hart, P.E. (1973) Pattern Classification and Scene Analysis. (Q327.D83) John Wiley & Sons. ISBN 0-471-22361-1. See page 218.
- Dasarathy, B.V. (1980) "Nosing Around the Neighborhood: A New System Structure and Classification Rule for Recognition in Partially Exposed Environments". IEEE Transactions on Pattern Analysis and Machine Intelligence, Vol. PAMI-2, No. 1, 67-71.

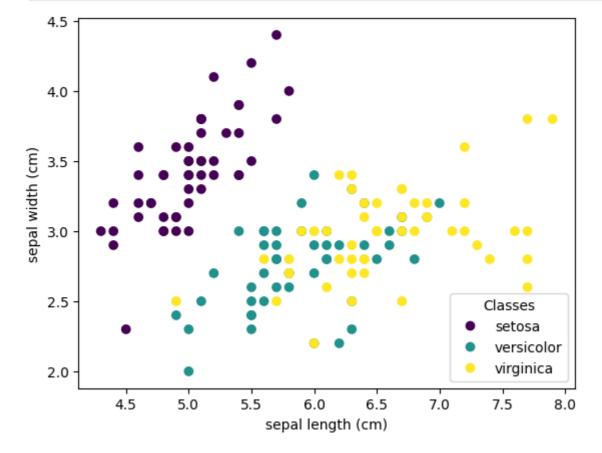
- Gates, G.W. (1972) "The Reduced Nearest Neighbor Rule". IEEE Transactions on Information Theory, May 1972, 431-433.
- See also: 1988 MLC Proceedings, 54-64. Cheeseman et al"s AUTOCLASS II conceptual clustering system finds 3 classes in the data.
- Many, many more ...

|details-end|

```
In [ ]: import pandas as pd
    csv_url = 'https://archive.ics.uci.edu/ml/machine-learning-databases/iris/iris.d
    df = pd.read_csv(csv_url, header = None)
    desc = df.describe()
    print(desc)
```

```
0
                             1
                                           2
       150.000000
                    150.000000
                                 150.000000
                                             150.000000
count
         5.843333
                      3.054000
                                   3.758667
                                                1.198667
mean
std
         0.828066
                      0.433594
                                   1.764420
                                                0.763161
min
         4.300000
                      2.000000
                                   1.000000
                                                0.100000
25%
         5.100000
                      2.800000
                                   1.600000
                                                0.300000
50%
         5.800000
                      3.000000
                                   4.350000
                                                1.300000
75%
         6.400000
                      3.300000
                                   5.100000
                                                1.800000
max
         7.900000
                      4.400000
                                   6.900000
                                                2.500000
```

```
In []: # Below code produces a scatter plot
   _, ax = plt.subplots()
   scatter = ax.scatter(X[:, 0], X[:, 1], c=Y)
   ax.set(xlabel=feature_names[0], ylabel=feature_names[1])
   _ = ax.legend(
        scatter.legend_elements()[0], category_names, loc="lower right", title="Clas")
```

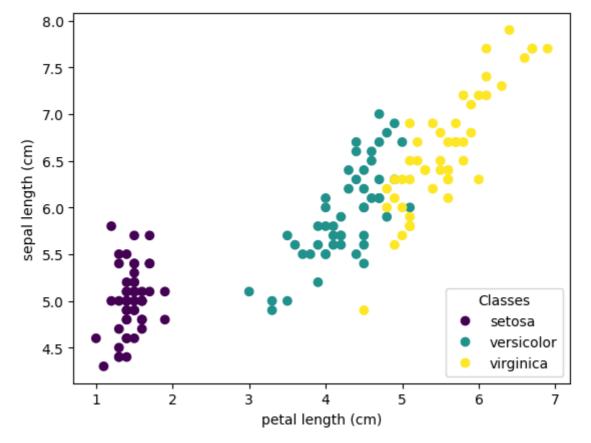


## What is the inference from above plot?

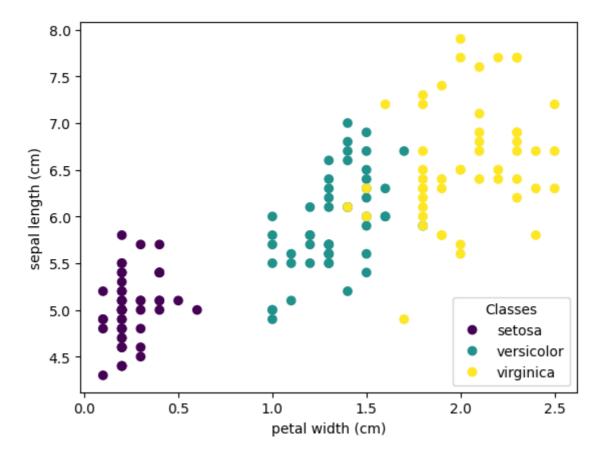
Type your answer here:

```
In []: # Repeat the above scatter plot for all pairs of features
# 1) sepal length in cm vs petal length in cm
# 2) sepal length in cm vs petal width in cm
# 3) petal length in cm vs petal width in cm

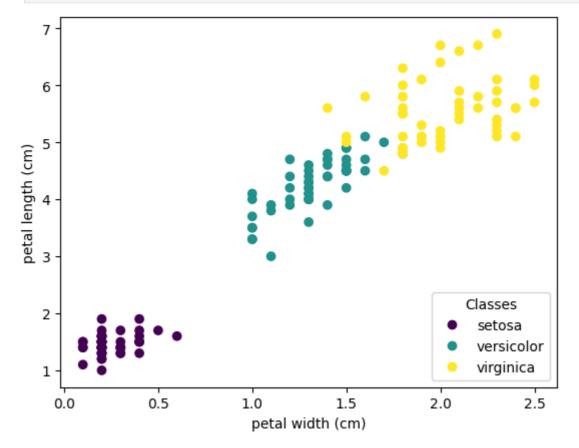
_, ax = plt.subplots()
scatter = ax.scatter(X[:, 2], X[:, 0], c=Y)
ax.set(xlabel=feature_names[2], ylabel=feature_names[0])
_ = ax.legend(
    scatter.legend_elements()[0], category_names, loc="lower right", title="Clas")
```



```
In [ ]: _, ax = plt.subplots()
scatter = ax.scatter(X[:, 3], X[:, 0], c=Y)
ax.set(xlabel=feature_names[3], ylabel=feature_names[0])
_ = ax.legend(
    scatter.legend_elements()[0], category_names, loc="lower right", title="Clas")
```



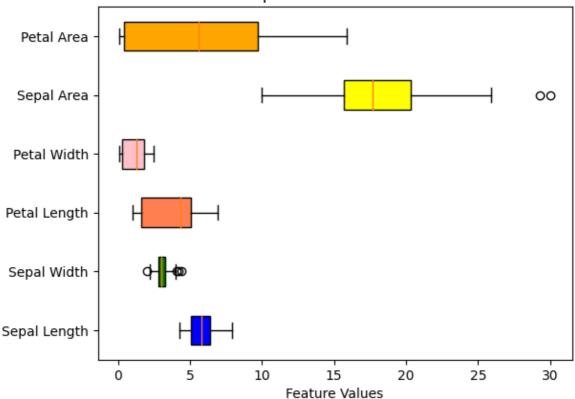
```
In [ ]: __, ax = plt.subplots()
    scatter = ax.scatter(X[:, 3], X[:, 2], c=Y)
    ax.set(xlabel=feature_names[3], ylabel=feature_names[2])
    _ = ax.legend(
        scatter.legend_elements()[0], category_names, loc="lower right", title="Clas")
```



```
In []: # Create two new features
    # sepal area = sepal length x sepal width,
    # petal area = petal length x petal width
    # Create new features
    sepal_area = X[:, 0] * X[:, 1]
    petal_area = X[:, 2] * X[:, 3]
    X = np.column_stack((X, sepal_area, petal_area))
```

```
In []: # Use boxplot for each feature plot in the same plot different
import matplotlib.pyplot as plt
import numpy as np
fig, ax = plt.subplots()
boxplots = ax.boxplot(X, vert=False, patch_artist=True)
ax.set_yticklabels(['Sepal Length', 'Sepal Width', 'Petal Length', 'Petal Width'
ax.set_xlabel('Feature Values')
ax.set_title('Boxplots for Each Feature')
colors = ['blue', 'green', 'coral', 'pink', 'yellow', 'orange']
for box, color in zip(boxplots['boxes'], colors):
    box.set_facecolor(color)
```

### Boxplots for Each Feature



```
In [ ]: # Find the average of each feature (including petal area, sepal area) for each c
    # and compare with the average of the complete dataset
    import numpy as np

data_with_labels = np.column_stack((X, Y))

unique_classes = np.unique(Y)

class_averages = {}
for class_label in unique_classes:
    class_data = data_with_labels[data_with_labels[:, -1] == class_label]
```

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```
Labex2
            class_averages[class_label] = np.mean(class_data[:, :-1], axis=0)
        overall_average = np.mean(X, axis=0)
        print("Class-wise averages:")
        for class_label, avg_values in class_averages.items():
            print(f"Class {int(class_label)}: {avg_values}")
        print("\nOverall average:")
        print(overall_average)
       Class-wise averages:
       Class 0: [ 5.006 3.428
                                 1.462 0.246 17.2578 0.3656]
       Class 1: [ 5.936 2.77
                                         1.326 16.5262 5.7204]
                                 4.26
       Class 2: [ 6.588 2.974
                                 5.552 2.026 19.6846 11.2962]
       Overall average:
       [ 5.84333333  3.05733333  3.758
                                            1.19933333 17.82286667 5.79406667]
In [ ]: !pip install statsmodels
       Requirement already satisfied: statsmodels in c:\users\deepthi\appdata\local\prog
       rams\python\python38\lib\site-packages (0.14.1)
       Requirement already satisfied: numpy<2,>=1.18 in c:\users\deepthi\appdata\local\p
       rograms\python\python38\lib\site-packages (from statsmodels) (1.24.3)
       Requirement already satisfied: scipy!=1.9.2,>=1.4 in c:\users\deepthi\appdata\loc
       al\programs\python\python38\lib\site-packages (from statsmodels) (1.10.1)
       Requirement already satisfied: pandas!=2.1.0,>=1.0 in c:\users\deepthi\appdata\lo
```

cal\programs\python\python38\lib\site-packages (from statsmodels) (2.0.3) Requirement already satisfied: patsy>=0.5.4 in c:\users\deepthi\appdata\local\pro grams\python\python38\lib\site-packages (from statsmodels) (0.5.6) Requirement already satisfied: packaging>=21.3 in c:\users\deepthi\appdata\local \programs\python\python38\lib\site-packages (from statsmodels) (23.1) Requirement already satisfied: python-dateutil>=2.8.2 in c:\users\deepthi\appdata \local\programs\python\python38\lib\site-packages (from pandas!=2.1.0,>=1.0->stat smodels) (2.8.2)

Requirement already satisfied: pytz>=2020.1 in c:\users\deepthi\appdata\local\pro grams\python\python38\lib\site-packages (from pandas!=2.1.0,>=1.0->statsmodels) (2023.3)

Requirement already satisfied: tzdata>=2022.1 in c:\users\deepthi\appdata\local\p rograms\python\python38\lib\site-packages (from pandas!=2.1.0,>=1.0->statsmodels) (2023.3)

Requirement already satisfied: six in c:\users\deepthi\appdata\local\programs\pyt hon\python38\lib\site-packages (from patsy>=0.5.4->statsmodels) (1.16.0)

```
In [ ]: # Check what is QQ plot and plot pair of features
        import matplotlib.pyplot as plt
        import numpy as np
        import statsmodels.api as sm
        num_features = X.shape[1]
        fig, axes = plt.subplots(nrows=num_features, ncols=num_features, figsize=(12, 12
        fig.subplots_adjust(hspace=0.5, wspace=0.5)
        for i in range(num features):
            for j in range(num_features):
                if i != j:
                     # Scatter plot for pairs of features
                    axes[i, j].scatter(X[:, i], X[:, j])
```

```
axes[i, j].set_xlabel(f'Feature {i}')
                                                                           axes[i, j].set_ylabel(f'Feature {j}')
                                                                           # QQ plot for each feature against a normal distribution
                                                                           sm.qqplot(X[:, i], line='q', ax=axes[i, j], fit=True)
                                                                           axes[i, j].set_title(f'QQ Plot - Feature {i} vs Feature {j}')
         plt.subplots_adjust(left=0.1,
                                                                                                                     bottom=0.1,
                                                                                                                     right=0.9,
                                                                                                                     top=0.9,
                                                                                                                     wspace=0.4,
                                                                                                                     hspace=0.4)
         plt.show()
                                                                  QQ Plot - Feature 0 vq@&dobreFeature 0 vq@&dobreFeature 0 vq@&dobreFeature 0 vq@&dobreFeature 0 vq.
            0.75
             0.50
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QQ Plot - Feature 1 vs Feature Detical QuaQQeBloth # Frature QuaqQ
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```

# **Experiment 2**

Theoretical Ouantiles

0.0

Sample Quantiles

Load Melbourne housing dataset supplied alongwith. The filename is "real\_estate.csv" file. You may use either pandas or numpy.

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0

Theoretical Quantiles Theoretical Quantiles

10

Theoretical Quantiles

10

0.25

Sample Quantiles

10

Theoretical Ouantiles

```
In [ ]: import pandas as pd
    file_path = "real_estate.csv"
    df = pd.read_csv(file_path)
    print(df.head())
```

	Address	Rooms	Price	Bedroom2	Bathroom	Landsize	YearBuilt	\
0	85 Turner St	2	1480000	2	1	202	NaN	
1	25 Bloomburg St	2	1035000	2	1	156	1900.0	
2	5 Charles St	3	1465000	3	2	134	1900.0	
3	40 Federation La	3	850000	3	2	94	NaN	
4	55a Park St	4	1600000	3	1	120	2014.0	

		Regionname	Suburb	Type
0	Northern	Metropolitan	Abbotsford	h
1	Northern	Metropolitan	Abbotsford	h
2	Northern	Metropolitan	Abbotsford	h
3	Northern	Metropolitan	Abbotsford	h
4	Northern	Metropolitan	Abbotsford	h

## **Unique Identifier**

• Address (replace the address with a unique number)

### Features of the dataset

- Rooms
- Price
- Number of Bedrooms
- Number of Bathrooms
- Land size
- Year built
- Region name
- Sub urban name

## Target variable (output variable t)

- Type of House
  - h house, cottage, villa, semi, terrace
  - u unit, duplex
  - t townhouse

#### In the above dataset,

- Convert the "Year built" into a categorical data
  - the houses built before 1800 is given 8
  - the houses built from 1800 to 1850 is given 7
  - the houses build from 1850 to 1900 is given 6
  - the houses built between 1900 to 1920 is given as 5
  - the houses built between 1920 to 1950 is given as 4
  - the houses built between 1950 to 1980 is given as 3
  - the houses built between 1980 to 2000 is given as 2
  - the houses built between 2000 to 2023 is given as 1
  - if there is no data on house built consider it as 0
- Use normalization techniques used in previous lab for "price" and "lab-size"?
- Choose any one "Region name" for that answer following questions

- Which is the dominant house-type?
- what is the average cost of different types of house?
- Use scatter plot for the following pair of features
  - o price vs number of bedrooms
  - o price vs number of bathrooms
  - land size vs price

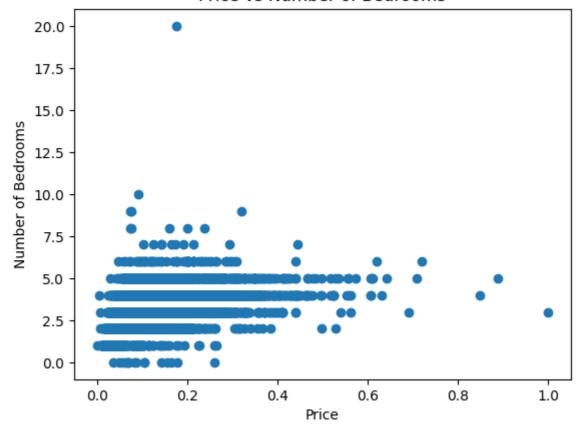
```
In [ ]: def categorize_year_built(year):
            if year < 1800:
                return 8
            elif 1800 <= year <= 1850:
                return 7
            elif 1850 <= year <= 1900:
                return 6
            elif 1900 <= year <= 1920:
                return 5
            elif 1920 <= year <= 1950:
                return 4
            elif 1950 <= year <= 1980:
                return 3
            elif 1980 <= year <= 2000:
                return 2
            elif 2000 <= year <= 2023:
                return 1
            else:
                return 0
        df['Year_built_categorical'] = df['YearBuilt'].apply(categorize_year_built)
In [ ]: from sklearn.preprocessing import MinMaxScaler
        scaler = MinMaxScaler()
        df[['Price', 'Landsize']] = scaler.fit_transform(df[['Price', 'Landsize']])
In [ ]: region df = df[df['Regionname'] == 'Region 1']
In [ ]: dominant_house_type_numerical = df['Type'].mode().iloc[0]
        print(f"Dominant House Type (Numerical): {dominant house type numerical}")
       Dominant House Type (Numerical): h
In [ ]: average_cost_by_type = df.groupby('Type')['Price'].mean()
        print(f"Average Cost of Different House Types in Region_1:\n{average_cost_by_typ
       Average Cost of Different House Types in Region 1:
       Type
           0.129856
            0.095203
            0.058343
       Name: Price, dtype: float64
In [ ]: import matplotlib.pyplot as plt
        plt.scatter(df['Price'], df['Bedroom2'])
        plt.xlabel('Price')
        plt.ylabel('Number of Bedrooms')
        plt.title('Price vs Number of Bedrooms')
```

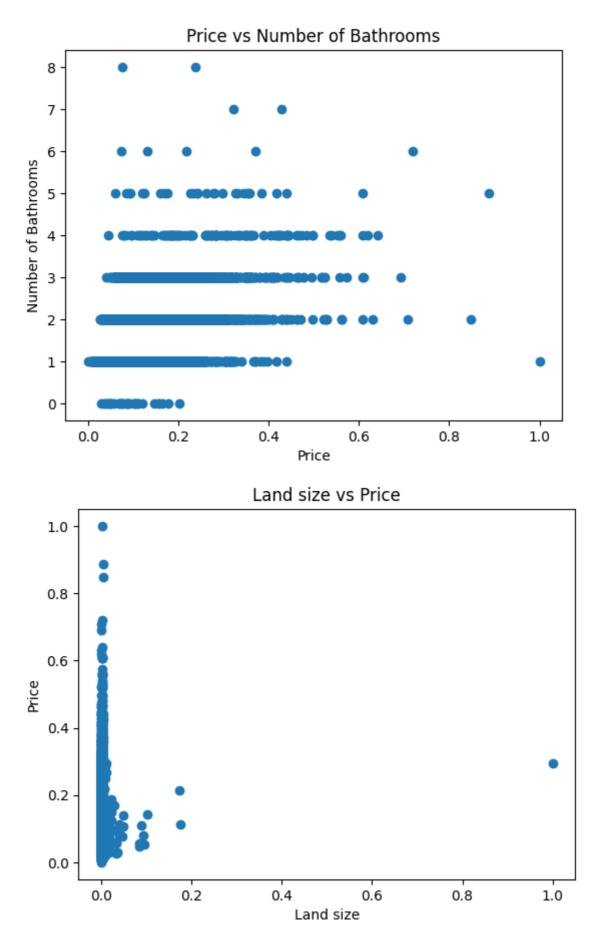
```
plt.show()

plt.scatter(df['Price'], df['Bathroom'])
plt.xlabel('Price')
plt.ylabel('Number of Bathrooms')
plt.title('Price vs Number of Bathrooms')
plt.show()

plt.scatter(df['Landsize'], df['Price'])
plt.xlabel('Land size')
plt.ylabel('Price')
plt.title('Land size vs Price')
plt.show()
```

## Price vs Number of Bedrooms





# **Covariance matrix**

Create a numpy array with 3 features land size, price, number of bedrooms. the size of the array will be 3xN. N is the number of samples.

An example of how to find the covariance matrix is given below:

## Find the covariance matrix for entire real\_estate database.

```
In [ ]: numeric_columns = df.select_dtypes(include='number')
    covariance_matrix = numeric_columns.cov()

# Display the covariance matrix
    print("Covariance Matrix:")
    print(covariance_matrix)
```

#### Covariance Matrix:

```
        Rooms
        Price
        Bedroom2
        Bathroom
        Landsize
        \

        Rooms
        0.913454
        0.034038
        0.871655
        0.391990
        0.000226

        Price
        0.034038
        0.005143
        0.032968
        0.023167
        0.000025

        Bedroom2
        0.871655
        0.032968
        0.933003
        0.390651
        0.000228

        Bathroom
        0.391990
        0.023167
        0.390651
        0.478465
        0.000237

        Landsize
        0.000226
        0.000025
        0.000228
        0.000237
        0.000085

        YearBuilt
        -2.388385
        -0.901477
        -1.940503
        4.010633
        0.003116

        Year_built_categorical
        0.046326
        0.018412
        0.034478
        -0.033262
        -0.000359
```

	YearBuilt	Year_built_categorical
Rooms	-2.388385	0.046326
Price	-0.901477	0.018412
Bedroom2	-1.940503	0.034478
Bathroom	4.010633	-0.033262
Landsize	0.003116	-0.000359
YearBuilt	1389.333350	-53.548282
Year built categorical	-53.548282	3.552879

# Basic linear algebra

- Find the eigen values of the covariance matrix of the entire dataset (use eigen value decomposition)
- Find the singular values of the covariance matrix of the entire dataset

```
In [ ]: import numpy as np
        # Assuming df is your DataFrame containing the real estate data
        numeric_columns = df.select_dtypes(include='number')
        covariance_matrix = numeric_columns.cov()
        # Eigenvalues and eigenvectors using eigenvalue decomposition
        eigenvalues, eigenvectors = np.linalg.eig(covariance_matrix)
        # Singular value decomposition
        singular_values = np.linalg.svd(covariance_matrix, compute_uv=False)
        # Display the results
        print("Eigenvalues:")
        print(eigenvalues)
        print("\nEigenvectors:")
        print(eigenvectors)
        print("\nSingular Values:")
        print(singular_values)
       Eigenvalues:
       [1.39141839e+03 1.99687287e+00 1.50112883e+00 2.45974845e-01
        5.12847740e-02 2.64571042e-03 8.47212265e-05]
       Eigenvectors:
       [ 1.71771308e-03 -6.60327910e-01 7.11216096e-03 -2.22758131e-01
          7.16990742e-01 -1.47174138e-02 7.96124391e-05]
        [ 6.47937132e-04 -2.60734982e-02 9.15977346e-03 4.18905181e-02
          9.40952555e-03 9.98688637e-01 -3.67477677e-03]
        [ 1.39574984e-03 -6.66911347e-01 3.66813180e-03 -2.64766293e-01
         -6.96500259e-01 2.21862359e-04 -4.00855013e-05]
        [-2.88126935e-03 -3.42834521e-01 -1.12247950e-01 9.31118367e-01
         -2.62952393e-02 -4.67290624e-02 -3.58093614e-04]
        [-2.24801292e-06 -1.97465426e-04 1.42749802e-04 5.10656160e-04
         -6.06848941e-05 3.65226576e-03 9.99993169e-01]
        [-9.99249687e-01 1.13404405e-04 -3.79617135e-02 -7.54516477e-03
          5.57665490e-04 1.31067923e-03 2.29498786e-06]
        [ 3.85544128e-02 3.13196201e-02 -9.92880328e-01 -1.07164385e-01
          5.60097314e-03 1.43420206e-02 1.50688889e-04]]
       Singular Values:
       [1.39141839e+03 1.99687287e+00 1.50112883e+00 2.45974845e-01
        5.12847740e-02 2.64571042e-03 8.47212265e-05]
```