

```
import pandas as pd
```

DataSet for Experiment No.01

Read Data Set

```
df= pd.read_csv('iris.csv')
```

```
pd.read_csv('iris.csv')
```

|     | sepal_length | sepal_width | petal_length | petal_width | species   |
|-----|--------------|-------------|--------------|-------------|-----------|
| 0   | 5.1          | 3.5         | 1.4          | 0.2         | setosa    |
| 1   | 4.9          | 3.0         | 1.4          | 0.2         | setosa    |
| 2   | 4.7          | 3.2         | 1.3          | 0.2         | setosa    |
| 3   | 4.6          | 3.1         | 1.5          | 0.2         | setosa    |
| 4   | 5.0          | 3.6         | 1.4          | 0.2         | setosa    |
| ... | ...          | ...         | ...          | ...         | ...       |
| 145 | 6.7          | 3.0         | 5.2          | 2.3         | virginica |
| 146 | 6.3          | 2.5         | 5.0          | 1.9         | virginica |
| 147 | 6.5          | 3.0         | 5.2          | 2.0         | virginica |
| 148 | 6.2          | 3.4         | 5.4          | 2.3         | virginica |
| 149 | 5.9          | 3.0         | 5.1          | 1.8         | virginica |

[150 rows x 5 columns]

```
df.describe()
```

|       | sepal_length | sepal_width | petal_length | petal_width |
|-------|--------------|-------------|--------------|-------------|
| count | 150.000000   | 150.000000  | 150.000000   | 150.000000  |
| mean  | 5.843333     | 3.054000    | 3.758667     | 1.198667    |
| 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    |

Preprocessing

```
df.shape
```

```
(150, 5)
```

```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
```

```
RangeIndex: 150 entries, 0 to 149
```

```
Data columns (total 5 columns):
```

| # | Column       | Non-Null Count | Dtype   |
|---|--------------|----------------|---------|
| 0 | sepal_length | 150 non-null   | float64 |

```
1  sepal_width  150 non-null  float64
2  petal_length 150 non-null  float64
3  petal_width  150 non-null  float64
4  species      150 non-null  object
```

```
dtypes: float64(4), object(1)
```

```
memory usage: 6.0+ KB
```

```
df.info
```

```
<bound method DataFrame.info of      sepal_length  sepal_width
petal_length  petal_width  species
0            5.1         3.5         1.4         0.2      setosa
1            4.9         3.0         1.4         0.2      setosa
2            4.7         3.2         1.3         0.2      setosa
3            4.6         3.1         1.5         0.2      setosa
4            5.0         3.6         1.4         0.2      setosa
..          ...         ...         ...         ...         ...
145          6.7         3.0         5.2         2.3  virginica
146          6.3         2.5         5.0         1.9  virginica
147          6.5         3.0         5.2         2.0  virginica
148          6.2         3.4         5.4         2.3  virginica
149          5.9         3.0         5.1         1.8  virginica
```

```
[150 rows x 5 columns]>
```

```
df.dtypes
```

```
sepal_length    float64
sepal_width     float64
petal_length     float64
petal_width     float64
species         object
dtype: object
```

```
df.ndim
```

```
2
```

```
df.head
```

```
<bound method NDFrame.head of      sepal_length  sepal_width
petal_length  petal_width  species
0            5.1         3.5         1.4         0.2      setosa
1            4.9         3.0         1.4         0.2      setosa
2            4.7         3.2         1.3         0.2      setosa
3            4.6         3.1         1.5         0.2      setosa
4            5.0         3.6         1.4         0.2      setosa
..          ...         ...         ...         ...         ...
145          6.7         3.0         5.2         2.3  virginica
146          6.3         2.5         5.0         1.9  virginica
147          6.5         3.0         5.2         2.0  virginica
```

|     |     |     |     |     |           |
|-----|-----|-----|-----|-----|-----------|
| 148 | 6.2 | 3.4 | 5.4 | 2.3 | virginica |
| 149 | 5.9 | 3.0 | 5.1 | 1.8 | virginica |

[150 rows x 5 columns]>

df.head()

|   | sepal_length | sepal_width | petal_length | petal_width | species |
|---|--------------|-------------|--------------|-------------|---------|
| 0 | 5.1          | 3.5         | 1.4          | 0.2         | setosa  |
| 1 | 4.9          | 3.0         | 1.4          | 0.2         | setosa  |
| 2 | 4.7          | 3.2         | 1.3          | 0.2         | setosa  |
| 3 | 4.6          | 3.1         | 1.5          | 0.2         | setosa  |
| 4 | 5.0          | 3.6         | 1.4          | 0.2         | setosa  |

df.tail

```
<bound method NDFrame.tail of
petal_length  petal_width  species
0             5.1         3.5      1.4         0.2      setosa
1             4.9         3.0      1.4         0.2      setosa
2             4.7         3.2      1.3         0.2      setosa
3             4.6         3.1      1.5         0.2      setosa
4             5.0         3.6      1.4         0.2      setosa
..            ...         ...      ...         ...      ...
145           6.7         3.0      5.2         2.3     virginica
146           6.3         2.5      5.0         1.9     virginica
147           6.5         3.0      5.2         2.0     virginica
148           6.2         3.4      5.4         2.3     virginica
149           5.9         3.0      5.1         1.8     virginica
```

[150 rows x 5 columns]>

df.tail()

|     | sepal_length | sepal_width | petal_length | petal_width | species   |
|-----|--------------|-------------|--------------|-------------|-----------|
| 145 | 6.7          | 3.0         | 5.2          | 2.3         | virginica |
| 146 | 6.3          | 2.5         | 5.0          | 1.9         | virginica |
| 147 | 6.5          | 3.0         | 5.2          | 2.0         | virginica |
| 148 | 6.2          | 3.4         | 5.4          | 2.3         | virginica |
| 149 | 5.9          | 3.0         | 5.1          | 1.8         | virginica |

df.notnull()

|     | sepal_length | sepal_width | petal_length | petal_width | species |
|-----|--------------|-------------|--------------|-------------|---------|
| 0   | True         | True        | True         | True        | True    |
| 1   | True         | True        | True         | True        | True    |
| 2   | True         | True        | True         | True        | True    |
| 3   | True         | True        | True         | True        | True    |
| 4   | True         | True        | True         | True        | True    |
| ..  | ...          | ...         | ...          | ...         | ...     |
| 145 | True         | True        | True         | True        | True    |

|     |      |      |      |      |      |
|-----|------|------|------|------|------|
| 146 | True | True | True | True | True |
| 147 | True | True | True | True | True |
| 148 | True | True | True | True | True |
| 149 | True | True | True | True | True |

[150 rows x 5 columns]

df.isnull()

|     | sepal_length | sepal_width | petal_length | petal_width | species |
|-----|--------------|-------------|--------------|-------------|---------|
| 0   | False        | False       | False        | False       | False   |
| 1   | False        | False       | False        | False       | False   |
| 2   | False        | False       | False        | False       | False   |
| 3   | False        | False       | False        | False       | False   |
| 4   | False        | False       | False        | False       | False   |
| ..  | ...          | ...         | ...          | ...         | ...     |
| 145 | False        | False       | False        | False       | False   |
| 146 | False        | False       | False        | False       | False   |
| 147 | False        | False       | False        | False       | False   |
| 148 | False        | False       | False        | False       | False   |
| 149 | False        | False       | False        | False       | False   |

[150 rows x 5 columns]

df.isnull

| <bound method DataFrame.isnull of |             |         | sepal_length | sepal_width |           |
|-----------------------------------|-------------|---------|--------------|-------------|-----------|
| petal_length                      | petal_width | species |              |             |           |
| 0                                 | 5.1         | 3.5     | 1.4          | 0.2         | setosa    |
| 1                                 | 4.9         | 3.0     | 1.4          | 0.2         | setosa    |
| 2                                 | 4.7         | 3.2     | 1.3          | 0.2         | setosa    |
| 3                                 | 4.6         | 3.1     | 1.5          | 0.2         | setosa    |
| 4                                 | 5.0         | 3.6     | 1.4          | 0.2         | setosa    |
| ..                                | ...         | ...     | ...          | ...         | ...       |
| 145                               | 6.7         | 3.0     | 5.2          | 2.3         | virginica |
| 146                               | 6.3         | 2.5     | 5.0          | 1.9         | virginica |
| 147                               | 6.5         | 3.0     | 5.2          | 2.0         | virginica |
| 148                               | 6.2         | 3.4     | 5.4          | 2.3         | virginica |
| 149                               | 5.9         | 3.0     | 5.1          | 1.8         | virginica |

[150 rows x 5 columns]>

df.sum()

|              |   |
|--------------|---|
| sepal_length | 876.5                                       |
| sepal_width  | 458.1                                       |
| petal_length | 563.8                                       |
| petal_width  | 179.8                                       |
| species      | setosasetosasetosasetosasetosasetosaseto... |
| dtype:       | object                                      |

```
df.max()
```

```
sepal_length    7.9
sepal_width     4.4
petal_length    6.9
petal_width     2.5
species         virginica
dtype: object
```

```
df.min()
```

```
sepal_length    4.3
sepal_width     2.0
petal_length    1.0
petal_width     0.1
species         setosa
dtype: object
```

```
df.min()
```

```
sepal_length    4.3
sepal_width     2.0
petal_length    1.0
petal_width     0.1
species         setosa
dtype: object
```

```
df.count()
```

```
sepal_length    150
sepal_width     150
petal_length    150
petal_width     150
species         150
dtype: int64
```

```
df.sort_index()
```

|     | sepal_length | sepal_width | petal_length | petal_width | species   |
|-----|--------------|-------------|--------------|-------------|-----------|
| 0   | 5.1          | 3.5         | 1.4          | 0.2         | setosa    |
| 1   | 4.9          | 3.0         | 1.4          | 0.2         | setosa    |
| 2   | 4.7          | 3.2         | 1.3          | 0.2         | setosa    |
| 3   | 4.6          | 3.1         | 1.5          | 0.2         | setosa    |
| 4   | 5.0          | 3.6         | 1.4          | 0.2         | setosa    |
| ... | ...          | ...         | ...          | ...         | ...       |
| 145 | 6.7          | 3.0         | 5.2          | 2.3         | virginica |
| 146 | 6.3          | 2.5         | 5.0          | 1.9         | virginica |
| 147 | 6.5          | 3.0         | 5.2          | 2.0         | virginica |
| 148 | 6.2          | 3.4         | 5.4          | 2.3         | virginica |
| 149 | 5.9          | 3.0         | 5.1          | 1.8         | virginica |

```
[150 rows x 5 columns]
```

```
df.sort_values
```

```
<bound method DataFrame.sort_values of      sepal_length  sepal_width  
petal_length  petal_width  species  
0           5.1         3.5         1.4         0.2      setosa  
1           4.9         3.0         1.4         0.2      setosa  
2           4.7         3.2         1.3         0.2      setosa  
3           4.6         3.1         1.5         0.2      setosa  
4           5.0         3.6         1.4         0.2      setosa  
..          ...         ...         ...         ...         ...  
145          6.7         3.0         5.2         2.3  virginica  
146          6.3         2.5         5.0         1.9  virginica  
147          6.5         3.0         5.2         2.0  virginica  
148          6.2         3.4         5.4         2.3  virginica  
149          5.9         3.0         5.1         1.8  virginica
```

```
[150 rows x 5 columns]>
```

```
df['sepal_length'].sort_values()
```

```
13      4.3  
42      4.4  
38      4.4  
8        4.4  
41      4.5
```

```
...  
122     7.7  
118     7.7  
117     7.7  
135     7.7  
131     7.9
```

```
Name: sepal_length, Length: 150, dtype: float64
```

```
df['sepal_length'].sort_index()
```

```
0      5.1  
1      4.9  
2      4.7  
3      4.6  
4      5.0
```

```
...  
145     6.7  
146     6.3  
147     6.5  
148     6.2  
149     5.9
```

```
Name: sepal_length, Length: 150, dtype: float64
```

```
df['sepal_length'].sort_index()
0      5.1
1      4.9
2      4.7
3      4.6
4      5.0
...
145     6.7
146     6.3
147     6.5
148     6.2
149     5.9
Name: sepal_length, Length: 150, dtype: float64

df['sepal_length'].mean()
5.843333333333334

df['sepal_length'].describe()
count      150.000000
mean        5.843333
std         0.828066
min         4.300000
25%         5.100000
50%         5.800000
75%         6.400000
max         7.900000
Name: sepal_length, dtype: float64

df['sepal_length'].sum()
876.5

df['sepal_length'].mean()
5.843333333333334

df['sepal_length'].max()
7.9

df['sepal_length'].min()
4.3

df['sepal_length'].count()
150

df['species'].value_counts()
```

```

species
setosa      50
versicolor  50
virginica   50
Name: count, dtype: int64

df['sepal_length'].value_counts()

sepal_length
5.0      10
5.1       9
6.3       9
5.7       8
6.7       8
5.8       7
5.5       7
6.4       7
4.9       6
5.4       6
6.1       6
6.0       6
5.6       6
4.8       5
6.5       5
6.2       4
7.7       4
6.9       4
4.6       4
5.2       4
5.9       3
4.4       3
7.2       3
6.8       3
6.6       2
4.7       2
7.6       1
7.4       1
7.3       1
7.0       1
7.1       1
5.3       1
4.3       1
4.5       1
7.9       1
Name: count, dtype: int64

```

#Data Fromating and Normalization

```
df['species'].replace(['setosa', 'varsicolor', 'virginica'], [1, 2, 3])
```



```

0      1
1      1
2      1
3      1
4      1
..
145    3
146    3
147    3
148    3
149    3
Name: species, Length: 150, dtype: object

```

```
df.value_counts()
```

| sepal_length | sepal_width | petal_length | petal_width | species    |   |
|--------------|-------------|--------------|-------------|------------|---|
| 4.9          | 3.1         | 1.5          | 0.1         | setosa     | 3 |
| 5.8          | 2.7         | 5.1          | 1.9         | virginica  | 2 |
|              | 4.0         | 1.2          | 0.2         | setosa     | 1 |
| 5.9          | 3.0         | 4.2          | 1.5         | versicolor | 1 |
| 6.2          | 3.4         | 5.4          | 2.3         | virginica  | 1 |
|              |             |              |             | ..         |   |
| 5.5          | 2.3         | 4.0          | 1.3         | versicolor | 1 |
|              | 2.4         | 3.7          | 1.0         | versicolor | 1 |
|              |             | 3.8          | 1.1         | versicolor | 1 |
|              | 2.5         | 4.0          | 1.3         | versicolor | 1 |
| 7.9          | 3.8         | 6.4          | 2.0         | virginica  | 1 |

```
Name: count, Length: 147, dtype: int64
```

```
df['species'].value_counts()
```

```

species
setosa      50
versicolor  50
virginica   50
Name: count, dtype: int64

```

```
df
```

|     | sepal_length | sepal_width | petal_length | petal_width | species   |
|-----|--------------|-------------|--------------|-------------|-----------|
| 0   | 5.1          | 3.5         | 1.4          | 0.2         | setosa    |
| 1   | 4.9          | 3.0         | 1.4          | 0.2         | setosa    |
| 2   | 4.7          | 3.2         | 1.3          | 0.2         | setosa    |
| 3   | 4.6          | 3.1         | 1.5          | 0.2         | setosa    |
| 4   | 5.0          | 3.6         | 1.4          | 0.2         | setosa    |
| ..  | ...          | ...         | ...          | ...         | ...       |
| 145 | 6.7          | 3.0         | 5.2          | 2.3         | virginica |
| 146 | 6.3          | 2.5         | 5.0          | 1.9         | virginica |
| 147 | 6.5          | 3.0         | 5.2          | 2.0         | virginica |
| 148 | 6.2          | 3.4         | 5.4          | 2.3         | virginica |
| 149 | 5.9          | 3.0         | 5.1          | 1.8         | virginica |

```
[150 rows x 5 columns]
df['species'].replace(['setosa','versicolor','virginica'],[1,2,3])
```

```
0      1
1      1
2      1
3      1
4      1
..
145    3
146    3
147    3
148    3
149    3
```

```
Name: species, Length: 150, dtype: object
```

```
df['species'].replace(['setosa','versicolor','virginica'],
[1,2,3],inplace=True)
```

```
df
```

|     | sepal_length | sepal_width | petal_length | petal_width | species |
|-----|--------------|-------------|--------------|-------------|---------|
| 0   | 5.1          | 3.5         | 1.4          | 0.2         | 1       |
| 1   | 4.9          | 3.0         | 1.4          | 0.2         | 1       |
| 2   | 4.7          | 3.2         | 1.3          | 0.2         | 1       |
| 3   | 4.6          | 3.1         | 1.5          | 0.2         | 1       |
| 4   | 5.0          | 3.6         | 1.4          | 0.2         | 1       |
| ..  | ...          | ...         | ...          | ...         | ...     |
| 145 | 6.7          | 3.0         | 5.2          | 2.3         | 3       |
| 146 | 6.3          | 2.5         | 5.0          | 1.9         | 3       |
| 147 | 6.5          | 3.0         | 5.2          | 2.0         | 3       |
| 148 | 6.2          | 3.4         | 5.4          | 2.3         | 3       |
| 149 | 5.9          | 3.0         | 5.1          | 1.8         | 3       |

```
[150 rows x 5 columns]
```

```
df['species'].value_counts()
```

```
species
1      50
2      50
3      50
```

```
Name: count, dtype: int64
```

```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
```

| # | Column       | Non-Null Count | Dtype   |
|---|--------------|----------------|---------|
| 0 | sepal_length | 150 non-null   | float64 |
| 1 | sepal_width  | 150 non-null   | float64 |
| 2 | petal_length | 150 non-null   | float64 |
| 3 | petal_width  | 150 non-null   | float64 |
| 4 | species      | 150 non-null   | int64   |

dtypes: float64(4), int64(1)

memory usage: 6.0 KB

df.info

```
<bound method DataFrame.info of
petal_length petal_width species sepal_length sepal_width
0          5.1          3.5          1.4          0.2          1
1          4.9          3.0          1.4          0.2          1
2          4.7          3.2          1.3          0.2          1
3          4.6          3.1          1.5          0.2          1
4          5.0          3.6          1.4          0.2          1
..          ...          ...          ...          ...          ...
145         6.7          3.0          5.2          2.3          3
146         6.3          2.5          5.0          1.9          3
147         6.5          3.0          5.2          2.0          3
148         6.2          3.4          5.4          2.3          3
149         5.9          3.0          5.1          1.8          3
```

[150 rows x 5 columns]>

df['species'].astype

```
<bound method NDFrame.astype of 0      1
1      1
2      1
3      1
4      1
..
145    3
146    3
147    3
148    3
149    3
```

Name: species, Length: 150, dtype: int64>

df.astype(float)

|   | sepal_length | sepal_width | petal_length | petal_width | species |
|---|--------------|-------------|--------------|-------------|---------|
| 0 | 5.1          | 3.5         | 1.4          | 0.2         | 1.0     |
| 1 | 4.9          | 3.0         | 1.4          | 0.2         | 1.0     |
| 2 | 4.7          | 3.2         | 1.3          | 0.2         | 1.0     |
| 3 | 4.6          | 3.1         | 1.5          | 0.2         | 1.0     |
| 4 | 5.0          | 3.6         | 1.4          | 0.2         | 1.0     |

```

..      ...      ...      ...      ...      ...
145      6.7      3.0      5.2      2.3      3.0
146      6.3      2.5      5.0      1.9      3.0
147      6.5      3.0      5.2      2.0      3.0
148      6.2      3.4      5.4      2.3      3.0
149      5.9      3.0      5.1      1.8      3.0

```

```
[150 rows x 5 columns]
```

```
df
```

```

      sepal_length  sepal_width  petal_length  petal_width  species
0              5.1           3.5           1.4           0.2         1
1              4.9           3.0           1.4           0.2         1
2              4.7           3.2           1.3           0.2         1
3              4.6           3.1           1.5           0.2         1
4              5.0           3.6           1.4           0.2         1
..      ...      ...      ...      ...      ...
145      6.7      3.0      5.2      2.3         3
146      6.3      2.5      5.0      1.9         3
147      6.5      3.0      5.2      2.0         3
148      6.2      3.4      5.4      2.3         3
149      5.9      3.0      5.1      1.8         3

```

```
[150 rows x 5 columns]
```

```
df=pd.read_csv('iris.csv')
```

```
df.describe()
```

```

      sepal_length  sepal_width  petal_length  petal_width
count    150.000000    150.000000    150.000000    150.000000
mean       5.843333     3.054000     3.758667     1.198667
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

```

```
#Categorical-> Normalization
```

```
from sklearn.preprocessing
```

```

import numpy as np
from sklearn import preprocessing

import LabelEncoder as le

```

```

-----
-----

```

ModuleNotFoundError

Traceback (most recent call

last)

Cell In[212], line 1

```
----> 1 import LabelEncoder as le
```

ModuleNotFoundError: No module named 'LabelEncoder'