

Assignment No. 1

Aim : Data Wrangling I Perform the following operations using Python on any open source dataset (e.g., data.csv) 1. Import all the required Python Libraries. 2. Locate an open source data from the web (e.g., <https://www.kaggle.com>). Provide a clear description of the data and its source (i.e., URL of the web site). 3. Load the Dataset into pandas dataframe. 4. Data Preprocessing: check for missing values in the data using pandas `isnull()`, `describe()` function to get some initial statistics. Provide variable descriptions. Types of variables etc. Check the dimensions of the data frame. 5. Data Formatting and Data Normalization: Summarize the types of variables by checking the data types (i.e., character, numeric, integer, factor, and logical) of the variables in the data set. If variables are not in the correct data type, apply proper type conversions. 6. Turn categorical variables into quantitative variables in Python. In addition to the codes and outputs, explain every operation that you do in the above steps and explain everything that you do to import/read/scrape the data set.

Code :

```
[1]: import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
from pandas import DataFrame, Series
```

```
[6]: import seaborn as ans
```

```
[12]: data = ans.load_dataset("iris")
```

```
[14]: print(data)
```

	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]

```
[18]: print(data)
```

	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]

```
[19]: data.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
```

```
[21]: data.head()
```

```
[21]:
```

	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

```
[22]: data.tail()
```

```
[22]:      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
```

```
[23]: data.describe()
```

```
[23]:      sepal_length  sepal_width  petal_length  petal_width
count      150.000000      150.000000      150.000000      150.000000
mean         5.843333         3.057333         3.758000         1.199333
std          0.828066         0.435866         1.765298         0.762238
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
```

```
[24]: top_left_corner_df = data.iloc[:4, :4]
```

```
[25]: print(top_left_corner_df)
```

```
      sepal_length  sepal_width  petal_length  petal_width
0          5.1          3.5          1.4          0.2
1          4.9          3.0          1.4          0.2
2          4.7          3.2          1.3          0.2
3          4.6          3.1          1.5          0.2
```

```
[27]: data.to_csv()
```

```
[27]: ',sepal_length,sepal_width,petal_length,petal_width,species\r\n0,5.1,3.5,1.4,0.2
,setosa\r\n1,4.9,3.0,1.4,0.2,setosa\r\n2,4.7,3.2,1.3,0.2,setosa\r\n3,4.6,3.1,1.5
,0.2,setosa\r\n4,5.0,3.6,1.4,0.2,setosa\r\n5,5.4,3.9,1.7,0.4,setosa\r\n6,4.6,3.4
,1.4,0.3,setosa\r\n7,5.0,3.4,1.5,0.2,setosa\r\n8,4.4,2.9,1.4,0.2,setosa\r\n9,4.9
,3.1,1.5,0.1,setosa\r\n10,5.4,3.7,1.5,0.2,setosa\r\n11,4.8,3.4,1.6,0.2,setosa\r\
n12,4.8,3.0,1.4,0.1,setosa\r\n13,4.3,3.0,1.1,0.1,setosa\r\n14,5.8,4.0,1.2,0.2,se
tosa\r\n15,5.7,4.4,1.5,0.4,setosa\r\n16,5.4,3.9,1.3,0.4,setosa\r\n17,5.1,3.5,1.4
,0.3,setosa\r\n18,5.7,3.8,1.7,0.3,setosa\r\n19,5.1,3.8,1.5,0.3,setosa\r\n20,5.4,
3.4,1.7,0.2,setosa\r\n21,5.1,3.7,1.5,0.4,setosa\r\n22,4.6,3.6,1.0,0.2,setosa\r\n
23,5.1,3.3,1.7,0.5,setosa\r\n24,4.8,3.4,1.9,0.2,setosa\r\n25,5.0,3.0,1.6,0.2,set
osa\r\n26,5.0,3.4,1.6,0.4,setosa\r\n27,5.2,3.5,1.5,0.2,setosa\r\n28,5.2,3.4,1.4,
0.2,setosa\r\n29,4.7,3.2,1.6,0.2,setosa\r\n30,4.8,3.1,1.6,0.2,setosa\r\n31,5.4,3
.4,1.5,0.4,setosa\r\n32,5.2,4.1,1.5,0.1,setosa\r\n33,5.5,4.2,1.4,0.2,setosa\r\n3
4,4.9,3.1,1.5,0.2,setosa\r\n35,5.0,3.2,1.2,0.2,setosa\r\n36,5.5,3.5,1.3,0.2,seto
sa\r\n37,4.9,3.6,1.4,0.1,setosa\r\n38,4.4,3.0,1.3,0.2,setosa\r\n39,5.1,3.4,1.5,0
.2,setosa\r\n40,5.0,3.5,1.3,0.3,setosa\r\n41,4.5,2.3,1.3,0.3,setosa\r\n42,4.4,3.
```

2,1.3,0.2,setosa\r\n43,5.0,3.5,1.6,0.6,setosa\r\n44,5.1,3.8,1.9,0.4,setosa\r\n45,4.8,3.0,1.4,0.3,setosa\r\n46,5.1,3.8,1.6,0.2,setosa\r\n47,4.6,3.2,1.4,0.2,setosa\r\n48,5.3,3.7,1.5,0.2,setosa\r\n49,5.0,3.3,1.4,0.2,setosa\r\n50,7.0,3.2,4.7,1.4,versicolor\r\n51,6.4,3.2,4.5,1.5,versicolor\r\n52,6.9,3.1,4.9,1.5,versicolor\r\n53,5.5,2.3,4.0,1.3,versicolor\r\n54,6.5,2.8,4.6,1.5,versicolor\r\n55,5.7,2.8,4.5,1.3,versicolor\r\n56,6.3,3.3,4.7,1.6,versicolor\r\n57,4.9,2.4,3.3,1.0,versicolor\r\n58,6.6,2.9,4.6,1.3,versicolor\r\n59,5.2,2.7,3.9,1.4,versicolor\r\n60,5.0,2.0,3.5,1.0,versicolor\r\n61,5.9,3.0,4.2,1.5,versicolor\r\n62,6.0,2.2,4.0,1.0,versicolor\r\n63,6.1,2.9,4.7,1.4,versicolor\r\n64,5.6,2.9,3.6,1.3,versicolor\r\n65,6.7,3.1,4.4,1.4,versicolor\r\n66,5.6,3.0,4.5,1.5,versicolor\r\n67,5.8,2.7,4.1,1.0,versicolor\r\n68,6.2,2.2,4.5,1.5,versicolor\r\n69,5.6,2.5,3.9,1.1,versicolor\r\n70,5.9,3.2,4.8,1.8,versicolor\r\n71,6.1,2.8,4.0,1.3,versicolor\r\n72,6.3,2.5,4.9,1.5,versicolor\r\n73,6.1,2.8,4.7,1.2,versicolor\r\n74,6.4,2.9,4.3,1.3,versicolor\r\n75,6.6,3.0,4.4,1.4,versicolor\r\n76,6.8,2.8,4.8,1.4,versicolor\r\n77,6.7,3.0,5.0,1.7,versicolor\r\n78,6.0,2.9,4.5,1.5,versicolor\r\n79,5.7,2.6,3.5,1.0,versicolor\r\n80,5.5,2.4,3.8,1.1,versicolor\r\n81,5.5,2.4,3.7,1.0,versicolor\r\n82,5.8,2.7,3.9,1.2,versicolor\r\n83,6.0,2.7,5.1,1.6,versicolor\r\n84,5.4,3.0,4.5,1.5,versicolor\r\n85,6.0,3.4,4.5,1.6,versicolor\r\n86,6.7,3.1,4.7,1.5,versicolor\r\n87,6.3,2.3,4.4,1.3,versicolor\r\n88,5.6,3.0,4.1,1.3,versicolor\r\n89,5.5,2.5,4.0,1.3,versicolor\r\n90,5.5,2.6,4.4,1.2,versicolor\r\n91,6.1,3.0,4.6,1.4,versicolor\r\n92,5.8,2.6,4.0,1.2,versicolor\r\n93,5.0,2.3,3.3,1.0,versicolor\r\n94,5.6,2.7,4.2,1.3,versicolor\r\n95,5.7,3.0,4.2,1.2,versicolor\r\n96,5.7,2.9,4.2,1.3,versicolor\r\n97,6.2,2.9,4.3,1.3,versicolor\r\n98,5.1,2.5,3.0,1.1,versicolor\r\n99,5.7,2.8,4.1,1.3,versicolor\r\n100,6.3,3.3,6.0,2.5,versicolor\r\n101,5.8,2.7,5.1,1.9,versicolor\r\n102,7.1,3.0,5.9,2.1,versicolor\r\n103,6.3,2.9,5.6,1.8,versicolor\r\n104,6.5,3.0,5.8,2.2,versicolor\r\n105,7.6,3.0,6.6,2.1,versicolor\r\n106,4.9,2.5,4.5,1.7,versicolor\r\n107,7.3,2.9,6.3,1.8,versicolor\r\n108,6.7,2.5,5.8,1.8,versicolor\r\n109,7.2,3.6,6.1,2.5,versicolor\r\n110,6.5,3.2,5.1,2.0,versicolor\r\n111,6.4,2.7,5.3,1.9,versicolor\r\n112,6.8,3.0,5.5,2.1,versicolor\r\n113,5.7,2.5,5.0,2.0,versicolor\r\n114,5.8,2.8,5.1,2.4,versicolor\r\n115,6.4,3.2,5.3,2.3,versicolor\r\n116,6.5,3.0,5.5,1.8,versicolor\r\n117,7.7,3.8,6.7,2.2,versicolor\r\n118,7.7,2.6,6.9,2.3,versicolor\r\n119,6.0,2.2,5.0,1.5,versicolor\r\n120,6.9,3.2,5.7,2.3,versicolor\r\n121,5.6,2.8,4.9,2.0,versicolor\r\n122,7.7,2.8,6.7,2.0,versicolor\r\n123,6.3,2.7,4.9,1.8,versicolor\r\n124,6.7,3.3,5.7,2.1,versicolor\r\n125,7.2,3.2,6.0,1.8,versicolor\r\n126,6.2,2.8,4.8,1.8,versicolor\r\n127,6.1,3.0,4.9,1.8,versicolor\r\n128,6.4,2.8,5.6,2.1,versicolor\r\n129,7.2,3.0,5.8,1.6,versicolor\r\n130,7.4,2.8,6.1,1.9,versicolor\r\n131,7.9,3.8,6.4,2.0,versicolor\r\n132,6.4,2.8,5.6,2.2,versicolor\r\n133,6.3,2.8,5.1,1.5,versicolor\r\n134,6.1,2.6,5.6,1.4,versicolor\r\n135,7.7,3.0,6.1,2.3,versicolor\r\n136,6.3,3.4,5.6,2.4,versicolor\r\n137,6.4,3.1,5.5,1.8,versicolor\r\n138,6.0,3.0,4.8,1.8,versicolor\r\n139,6.9,3.1,5.4,2.1,versicolor\r\n140,6.7,3.1,5.6,2.4,versicolor\r\n141,6.9,3.1,5.1,2.3,versicolor\r\n142,5.8,2.7,5.1,1.9,versicolor\r\n143,6.8,3.2,5.9,2.3,versicolor\r\n144,6.7,3.3,5.7,2.5,versicolor\r\n145,6.7,3.0,5.2,2.3,versicolor\r\n146,6.3,2.5,5.0,1.9,versicolor\r\n147,6.5,3.0,5.2,2.0,versicolor\r\n148,6.2,3.4,5.4,2.3,versicolor\r\n149,5.9,3.0,5.1,1.8,versicolor\r\n'

[28]: ash = data.copy()

```
[29]: print(ash)
```

	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]

```
[ ]:
```

```
[31]: data.count()
```

```
[31]: sepal_length    150
      sepal_width    150
      petal_length   150
      petal_width    150
      species        150
      dtype: int64
```

```
[34]: data.cummax()
```

```
[34]:
```

	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	setosa
1	5.1	3.5	1.4	0.2	setosa
2	5.1	3.5	1.4	0.2	setosa
3	5.1	3.5	1.5	0.2	setosa
4	5.1	3.6	1.5	0.2	setosa
..
145	7.9	4.4	6.9	2.5	virginica
146	7.9	4.4	6.9	2.5	virginica
147	7.9	4.4	6.9	2.5	virginica
148	7.9	4.4	6.9	2.5	virginica
149	7.9	4.4	6.9	2.5	virginica

[150 rows x 5 columns]

```
[35]: data.cummin()
```

```
[35]:      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           3.0           1.3           0.2  setosa
      3           4.6           3.0           1.3           0.2  setosa
      4           4.6           3.0           1.3           0.2  setosa
      ..          ...          ...          ...          ...          ...
     145          4.3           2.0           1.0           0.1  setosa
     146          4.3           2.0           1.0           0.1  setosa
     147          4.3           2.0           1.0           0.1  setosa
     148          4.3           2.0           1.0           0.1  setosa
     149          4.3           2.0           1.0           0.1  setosa
```

[150 rows x 5 columns]

```
[36]: data.dropna()
```

```
[36]:      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]

```
[37]: data.any()
```

```
[37]: sepal_length    True
      sepal_width    True
      petal_length    True
      petal_width    True
      species        True
      dtype: bool
```

```
[39]: data.get(40)
```

```
[40]: mr = data.get(40)
```

```
[41]: print(mr)
```

None

```
[4]: import seaborn as sea
```

```
[5]: data = sea.get_dataset_names()
```

```
[6]: print(data)
```

```
['anagrams', 'anscombe', 'attention', 'brain_networks', 'car_crashes',  
'diamonds', 'dots', 'dowjones', 'exercise', 'flights', 'fmri', 'geyser', 'glue',  
'healthexp', 'iris', 'mpg', 'penguins', 'planets', 'seaice', 'taxi', 'tips',  
'titanic', 'anagrams', 'anagrams', 'anscombe', 'anscombe', 'attention',  
'attention', 'brain_networks', 'brain_networks', 'car_crashes', 'car_crashes',  
'diamonds', 'diamonds', 'dots', 'dots', 'dowjones', 'dowjones', 'exercise',  
'exercise', 'flights', 'flights', 'fmri', 'fmri', 'geyser', 'geyser', 'glue',  
'glue', 'healthexp', 'healthexp', 'iris', 'iris', 'mpg', 'mpg', 'penguins',  
'penguins', 'planets', 'planets', 'seaice', 'seaice', 'taxi', 'taxi', 'tips',  
'tips', 'titanic', 'titanic', 'anagrams', 'anscombe', 'attention',  
'brain_networks', 'car_crashes', 'diamonds', 'dots', 'dowjones', 'exercise',  
'flights', 'fmri', 'geyser', 'glue', 'healthexp', 'iris', 'mpg', 'penguins',  
'planets', 'seaice', 'taxi', 'tips', 'titanic']
```

```
[7]: data = sea.load_dataset("iris")
```

```
[8]: data.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
```

```
[9]: data.describe()
```

```
[9]:
```

	sepal_length	sepal_width	petal_length	petal_width
count	150.000000	150.000000	150.000000	150.000000
mean	5.843333	3.057333	3.758000	1.199333
std	0.828066	0.435866	1.765298	0.762238
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

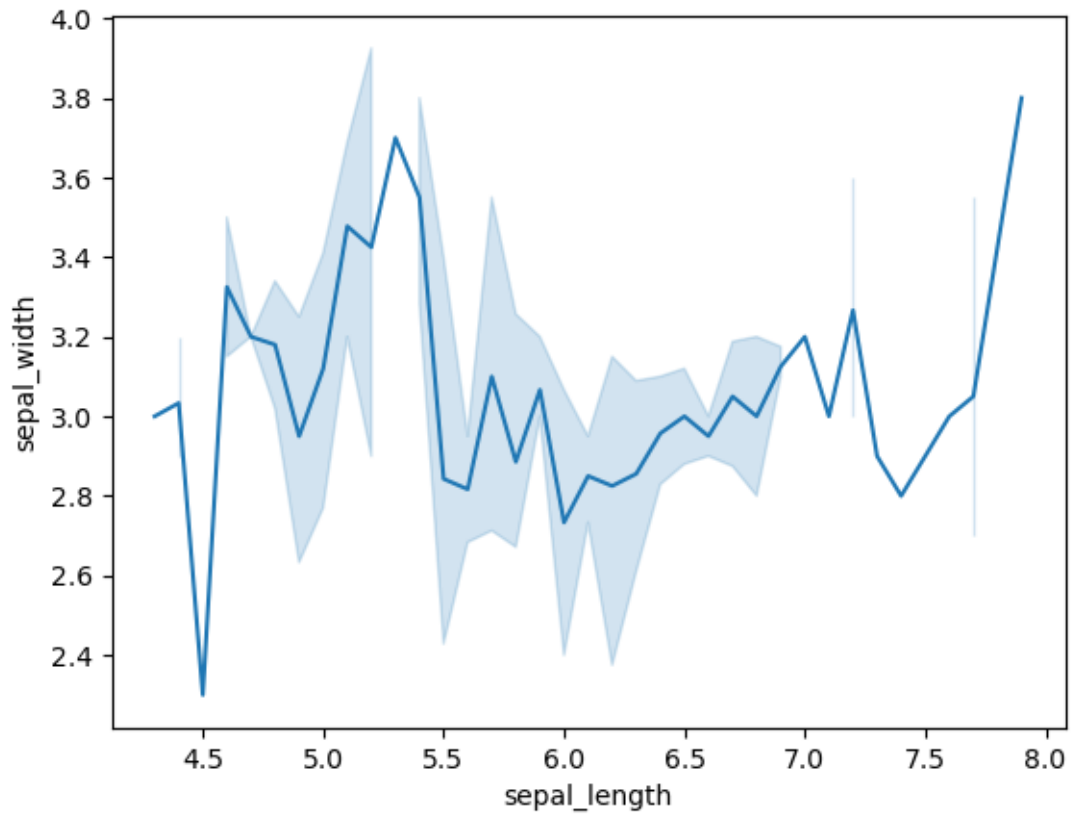
```
[10]: data.head()
```

```
[10]:   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
```

```
[11]: sea.lineplot(x="sepal_length", y="sepal_width", data=data)
```

```
C:\Users\Welcome\anaconda3\Lib\site-packages\seaborn\_oldcore.py:1119:
FutureWarning: use_inf_as_na option is deprecated and will be removed in a
future version. Convert inf values to NaN before operating instead.
  with pd.option_context('mode.use_inf_as_na', True):
C:\Users\Welcome\anaconda3\Lib\site-packages\seaborn\_oldcore.py:1119:
FutureWarning: use_inf_as_na option is deprecated and will be removed in a
future version. Convert inf values to NaN before operating instead.
  with pd.option_context('mode.use_inf_as_na', True):
```

```
[11]: <Axes: xlabel='sepal_length', ylabel='sepal_width'>
```

```
[12]: data.min()
```

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

```
[13]: data.max()
```

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

```
[15]: data.mode()
```

```
[15]:   sepal_length  sepal_width  petal_length  petal_width   species
      0         5.0         3.0         1.4         0.2     setosa
      1         NaN         NaN         1.5         NaN  versicolor
      2         NaN         NaN         NaN         NaN   virginica
```

```
[7]: import seaborn as san
data = san.get_dataset_names()
data
```

```
[7]: ['anagrams',
      'anscombe',
      'attention',
      'brain_networks',
      'car_crashes',
      'diamonds',
      'dots',
      'dowjones',
      'exercise',
      'flights',
      'fmri',
      'geyser',
      'glue',
      'healthexp',
      'iris',
      'mpg',
      'penguins',
      'planets',
      'seaice',
      'taxis',
      'tips',
      'titanic',
      'anagrams',
      'anagrams',
      'anscombe',
      'anscombe',
      'attention',
      'attention',
      'brain_networks',
      'brain_networks',
      'car_crashes',
      'car_crashes',
      'diamonds',
      'diamonds',
      'dots',
      'dots',
      'dowjones',
      'dowjones',
```

'exercise',
'exercise',
'flights',
'flights',
'fmri',
'fmri',
'geyser',
'geyser',
'glue',
'glue',
'healthexp',
'healthexp',
'iris',
'iris',
'mpg',
'mpg',
'penguins',
'penguins',
'planets',
'planets',
'seaice',
'seaice',
'taxis',
'taxis',
'tips',
'tips',
'titanic',
'titanic',
'anagrams',
'anscombe',
'attention',
'brain_networks',
'car_crashes',
'diamonds',
'dots',
'dowjones',
'exercise',
'flights',
'fmri',
'geyser',
'glue',
'healthexp',
'iris',
'mpg',
'penguins',
'planets',
'seaice',

```
'taxi',
'tips',
'titanic']
```

```
[16]: df.describe(include='all')
```

```
[16]:
```

	sepal_length	sepal_width	petal_length	petal_width	species
count	150.000000	150.000000	150.000000	150.000000	150
unique	NaN	NaN	NaN	NaN	3
top	NaN	NaN	NaN	NaN	setosa
freq	NaN	NaN	NaN	NaN	50
mean	5.843333	3.057333	3.758000	1.199333	NaN
std	0.828066	0.435866	1.765298	0.762238	NaN
min	4.300000	2.000000	1.000000	0.100000	NaN
25%	5.100000	2.800000	1.600000	0.300000	NaN
50%	5.800000	3.000000	4.350000	1.300000	NaN
75%	6.400000	3.300000	5.100000	1.800000	NaN
max	7.900000	4.400000	6.900000	2.500000	NaN

```
[20]: df.sort_index(axis=1, ascending=False)
```

```
[20]:
```

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

[150 rows x 5 columns]

```
[24]: df.sort_values(by="sepal_width")
```

```
[24]:
```

	sepal_length	sepal_width	petal_length	petal_width	species
60	5.0	2.0	3.5	1.0	versicolor
62	6.0	2.2	4.0	1.0	versicolor
119	6.0	2.2	5.0	1.5	virginica
68	6.2	2.2	4.5	1.5	versicolor
41	4.5	2.3	1.3	0.3	setosa
..
16	5.4	3.9	1.3	0.4	setosa
14	5.8	4.0	1.2	0.2	setosa

32	5.2	4.1	1.5	0.1	setosa
33	5.5	4.2	1.4	0.2	setosa
15	5.7	4.4	1.5	0.4	setosa

[150 rows x 5 columns]

```
[25]: df.iloc[5]
```

```
[25]: sepal_length    5.4
      sepal_width     3.9
      petal_length    1.7
      petal_width     0.4
      species        setosa
      Name: 5, dtype: object
```

```
[26]: df[0:3]
```

```
[26]:   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
```

```
[27]: df.loc[:, ["sepal_width", "petal_length"]]
```

```
[27]:   sepal_width  petal_length
0         3.5         1.4
1         3.0         1.4
2         3.2         1.3
3         3.1         1.5
4         3.6         1.4
..      ...      ...
145        3.0         5.2
146        2.5         5.0
147        3.0         5.2
148        3.4         5.4
149        3.0         5.1
```

[150 rows x 2 columns]

```
[29]: df.iloc[:30, :]
```

```
[29]:   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
```

5	5.4	3.9	1.7	0.4	setosa
6	4.6	3.4	1.4	0.3	setosa
7	5.0	3.4	1.5	0.2	setosa
8	4.4	2.9	1.4	0.2	setosa
9	4.9	3.1	1.5	0.1	setosa
10	5.4	3.7	1.5	0.2	setosa
11	4.8	3.4	1.6	0.2	setosa
12	4.8	3.0	1.4	0.1	setosa
13	4.3	3.0	1.1	0.1	setosa
14	5.8	4.0	1.2	0.2	setosa
15	5.7	4.4	1.5	0.4	setosa
16	5.4	3.9	1.3	0.4	setosa
17	5.1	3.5	1.4	0.3	setosa
18	5.7	3.8	1.7	0.3	setosa
19	5.1	3.8	1.5	0.3	setosa
20	5.4	3.4	1.7	0.2	setosa
21	5.1	3.7	1.5	0.4	setosa
22	4.6	3.6	1.0	0.2	setosa
23	5.1	3.3	1.7	0.5	setosa
24	4.8	3.4	1.9	0.2	setosa
25	5.0	3.0	1.6	0.2	setosa
26	5.0	3.4	1.6	0.4	setosa
27	5.2	3.5	1.5	0.2	setosa
28	5.2	3.4	1.4	0.2	setosa
29	4.7	3.2	1.6	0.2	setosa

```
[30]: df.iloc[:, :17]
```

```
[30]:
```

	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]

```
[31]: df.iloc[:6, :12]
```

```
[31]:
```

	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
5	5.4	3.9	1.7	0.4	setosa

```
[32]: df.iloc[3:5, 0:2]
```

```
[32]:      sepal_length  sepal_width
3             4.6           3.1
4             5.0           3.6
```

```
[33]: df.iloc[[1, 2, 4], [0, 2]]
```

```
[33]:      sepal_length  petal_length
1             4.9           1.4
2             4.7           1.3
4             5.0           1.4
```

```
[34]: df.iloc[1:3, :]
```

```
[34]:      sepal_length  sepal_width  petal_length  petal_width  species
1             4.9           3.0           1.4           0.2    setosa
2             4.7           3.2           1.3           0.2    setosa
```

```
[35]: df.iloc[:, 1:3]
```

```
[35]:      sepal_width  petal_length
0             3.5           1.4
1             3.0           1.4
2             3.2           1.3
3             3.1           1.5
4             3.6           1.4
..          ...          ...
145           3.0           5.2
146           2.5           5.0
147           3.0           5.2
148           3.4           5.4
149           3.0           5.1
```

[150 rows x 2 columns]

```
[36]: df.iloc[1, 1]
```

```
[36]: 3.0
```

```
[38]: df['sepal_length'].iloc[5]
```

[38]: 5.4

```
[41]: cols_2_4 = df.columns[2:4]
df[cols_2_4]
```

```
[41]:      petal_length  petal_width
0           1.4           0.2
1           1.4           0.2
2           1.3           0.2
3           1.5           0.2
4           1.4           0.2
..          ...          ...
145          5.2           2.3
146          5.0           1.9
147          5.2           2.0
148          5.4           2.3
149          5.1           1.8
```

[150 rows x 2 columns]

```
[42]: df[df.columns[2:4]].iloc[5:10]
```

```
[42]:      petal_length  petal_width
5           1.7           0.4
6           1.4           0.3
7           1.5           0.2
8           1.4           0.2
9           1.5           0.1
```

```
[43]: df.isnull()
```

```
[43]:      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]

```
[44]: df.isnull().any()
```



```
[44]: sepal_length    False
      sepal_width     False
      petal_length    False
      petal_width     False
      species         False
      dtype: bool
```

```
[46]: df.isnull().sum().sum()
```

```
[46]: 0
```

```
[47]: df.isnull().sum()
```

```
[47]: sepal_length    0
      sepal_width    0
      petal_length   0
      petal_width    0
      species        0
      dtype: int64
```

```
[49]: df.isnull().sum(axis=1)
```

```
[49]: 0      0
      1      0
      2      0
      3      0
      4      0
      ..
     145     0
     146     0
     147     0
     148     0
     149     0
      Length: 150, dtype: int64
```

```
[50]: df.isna().sum()
```

```
[50]: sepal_length    0
      sepal_width    0
      petal_length   0
      petal_width    0
      species        0
      dtype: int64
```

```
[51]: df.petal_length.isnull().sum()
```

```
[51]: 0
```

```
[53]: df.groupby(['sepal_length'])['petal_width'].apply(lambda x:x.isnull().sum())
```

```
[53]: sepal_length
4.3    0
4.4    0
4.5    0
4.6    0
4.7    0
4.8    0
4.9    0
5.0    0
5.1    0
5.2    0
5.3    0
5.4    0
5.5    0
5.6    0
5.7    0
5.8    0
5.9    0
6.0    0
6.1    0
6.2    0
6.3    0
6.4    0
6.5    0
6.6    0
6.7    0
6.8    0
6.9    0
7.0    0
7.1    0
7.2    0
7.3    0
7.4    0
7.6    0
7.7    0
7.9    0
Name: petal_width, dtype: int64
```

```
[55]: df.dtypes
```

```
[55]: sepal_length    float64
sepal_width      float64
petal_length     float64
petal_width      float64
species          object
```

dtype: object

```
[61]: df['petal_length'] = df['petal_length'].astype("int")
df['petal_length']
```

```
[61]: 0      1
      1      1
      2      1
      3      1
      4      1
      ..
     145     5
     146     5
     147     5
     148     5
     149     5
      Name: petal_length, Length: 150, dtype: int32
```

```
[68]: import pandas as pd
```

```
[69]: from sklearn import preprocessing
```

```
[70]: df.head()
```

```
[70]:   sepal_length  sepal_width  petal_length  petal_width  species
0           5.1           3.5           1           0.2   setosa
1           4.9           3.0           1           0.2   setosa
2           4.7           3.2           1           0.2   setosa
3           4.6           3.1           1           0.2   setosa
4           5.0           3.6           1           0.2   setosa
```

```
[72]: min_max_scaler = preprocessing.MinMaxScaler()
print(min_max_scaler)
```

MinMaxScaler()

```
[75]: x=df.iloc[:, :4]
```

```
[76]: x_scaled = min_max_scaler.fit_transform(x)
```

```
[77]: df_normalized = pd.DataFrame(x_scaled)
```

```
[78]: df_normalized
```

```
[78]:           0           1           2           3
0    0.222222  0.625000  0.0  0.041667
1    0.166667  0.416667  0.0  0.041667
```

```

2    0.111111  0.500000  0.0  0.041667
3    0.083333  0.458333  0.0  0.041667
4    0.194444  0.666667  0.0  0.041667
..      ...      ...  ...
145  0.666667  0.416667  0.8  0.916667
146  0.555556  0.208333  0.8  0.750000
147  0.611111  0.416667  0.8  0.791667
148  0.527778  0.583333  0.8  0.916667
149  0.444444  0.416667  0.8  0.708333

```

[150 rows x 4 columns]

```
[82]: df['species'].unique()
```

```
[82]: array(['setosa', 'versicolor', 'virginica'], dtype=object)
```

```
[83]: label_encoder = preprocessing.LabelEncoder()
```

```
[84]: df['species'] = label_encoder.fit_transform(df['species'])
```

```
[85]: df['species'].unique()
```

```
[85]: array([0, 1, 2])
```

```
[86]: features_df = df.drop(columns=['species'])
```

```
[87]: enc = preprocessing.OneHotEncoder()
```

```
[93]: enc_df = pd.DataFrame(enc.fit_transform(df[['species']]))
```

```
[95]: df_encode = features_df.join(enc_df)
```

```
[96]: df_encode
```

```
[96]:
```

	sepal_length	sepal_width	petal_length	petal_width	
0	5.1	3.5	1	0.2	(0, 0)\t1.0
1	4.9	3.0	1	0.2	(0, 0)\t1.0
2	4.7	3.2	1	0.2	(0, 0)\t1.0
3	4.6	3.1	1	0.2	(0, 0)\t1.0
4	5.0	3.6	1	0.2	(0, 0)\t1.0
..
145	6.7	3.0	5	2.3	(0, 2)\t1.0
146	6.3	2.5	5	1.9	(0, 2)\t1.0
147	6.5	3.0	5	2.0	(0, 2)\t1.0
148	6.2	3.4	5	2.3	(0, 2)\t1.0
149	5.9	3.0	5	1.8	(0, 2)\t1.0

[150 rows x 5 columns]

```
[97]: df_encode.rename(columns = {0:'Iris-Setosa',1:'Iris-Versicolor',2:
    ↳ 'Iris-virginica'}, inplace = True)
```

```
[98]: df_encode
```

```
[98]:
```

	sepal_length	sepal_width	petal_length	petal_width	Iris-Setosa
0	5.1	3.5	1	0.2	(0, 0)\t1.0
1	4.9	3.0	1	0.2	(0, 0)\t1.0
2	4.7	3.2	1	0.2	(0, 0)\t1.0
3	4.6	3.1	1	0.2	(0, 0)\t1.0
4	5.0	3.6	1	0.2	(0, 0)\t1.0
..
145	6.7	3.0	5	2.3	(0, 2)\t1.0
146	6.3	2.5	5	1.9	(0, 2)\t1.0
147	6.5	3.0	5	2.0	(0, 2)\t1.0
148	6.2	3.4	5	2.3	(0, 2)\t1.0
149	5.9	3.0	5	1.8	(0, 2)\t1.0

[150 rows x 5 columns]

```
[100]: one_hot_df = pd.get_dummies(df, prefix="species",columns=['species'],
    ↳ drop_first=True)
```

```
[101]: one_hot_df
```

```
[101]:
```

	sepal_length	sepal_width	petal_length	petal_width	species_1	species_2
0	5.1	3.5	1	0.2	False	False
1	4.9	3.0	1	0.2	False	False
2	4.7	3.2	1	0.2	False	False
3	4.6	3.1	1	0.2	False	False
4	5.0	3.6	1	0.2	False	False
..
145	6.7	3.0	5	2.3	False	True
146	6.3	2.5	5	1.9	False	True
147	6.5	3.0	5	2.0	False	True
148	6.2	3.4	5	2.3	False	True
149	5.9	3.0	5	1.8	False	True

[150 rows x 6 columns]

```
[17]: class Solution:
    def solve(str, s):
        output = ""
        num=""
        for i in s:
```

```

        if i.isalpha():
            output+=i*int(num)
            num=""
        else:
            num+=i
    return output
print("Enter a string : ")
str = input()
ob = Solution()
print(ob.solve(str))

```

Enter a string :
4B3A
BBBBAAA

```

[28]: list1 = [1, 2, 3, 4, 5]
      list2 = [4, 5, 6, 7, 8]

      common = list(set(list1) & set(list2))

      print(common)

```

[4, 5]

```

[30]: L1= ['Sohan', 'Mohan', 'Rohan']
      for string in L1:
          print (string[0])

```

S
M
R

```

[31]: a = ['pandas', 'numpy', 'flask', 'python', 'python']

      s = set()

      dup = []

      for n in a:
          if n in s:
              dup.append(n)
          else:
              s.add(n)

      print(dup)

```

['python']

```
[4]: a = [1,2,5,3,4,8,9,"lis","a"]  
     length = len(a)  
     print(length)
```

9

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