

St. Francis Institute of Technology, Mumbai-400 103
Department of Information Technology

A.Y. 2021-2022
Class: TE-ITA/B, Semester: VI

Subject: **Data Science Lab**

Experiment – 1: To implement Data Preparation using Numpy and Pandas.

1. **Aim:** To implement Data Preparation using Numpy and Pandas.
2. **Objectives:** After study of this experiment, the student will be able to
 - Understand Numpy concepts
 - Understand Pandas concepts
3. **Outcomes:** After study of this experiment, the student will be able to
 - Understand data preparation, Numpy and Pandas.
4. **Prerequisite:** Fundamentals of Python Programming and Database Management System.
5. **Requirements:** Python Installation, Personal Computer, Windows operating system, Internet Connection, Microsoft Word.
6. **Pre-Experiment Exercise:**
Brief Theory:
Basic Concepts of Pandas and Numpy.
7. **Laboratory Exercise**
A. Procedure:
Software Installation:

1. Python 3.6
 - This setup requires that your machine has python 3.6 installed on it. you can refer to this url <https://www.python.org/downloads/> to download python. Once you have python downloaded and installed, you will need to setup PATH variables (if you want to run python program directly). To do that check this: <https://www.pythontutorial.net/add-python-to-path-python-is-not-recognized-as-an-internal-or-external-command/>.
 - Setting up PATH variable is optional as you can also run program without it.
2. Second and easier option is to download anaconda and use its anaconda prompt to run the commands. To install anaconda check this url <https://www.anaconda.com/download/>
3. You will also need to download and install below 2 packages after you install either python or anaconda from the steps above
 - Pandas
 - Numpy
- if you have chosen to install python 3.6 then run below commands in command prompt/terminal to install these packages

```
pip install pandas  
pip install numpy
```

- if you have chosen to install anaconda then run below commands in anaconda prompt to install these packages

```
conda install -c anaconda numpy
```

```
conda install -c anaconda pandas
```

4. Use Google Colab

Dataset Used:**Iris Dataset:**

Iris Dataset is considered as the Hello World for data science. It contains five columns namely – Petal Length, Petal Width, Sepal Length, Sepal Width, and Species Type. Iris is a flowering plant, the researchers have measured various features of the different iris flowers and recorded them digitally.

```
import pandas as pd
import numpy as np
df = pd.read_csv("https://raw.githubusercontent.com/uiuc-cse/data-fa14/gh-
pages/data/iris.csv")

df.info()

"""**Data Inspection**"""

df.head(5) # head

df.shape

df.columns

df["sepal_length"].nunique()

df["sepal_length"].unique()

# number of unique values altogether
df.columns.nunique()

# value counts
df['species'].value_counts()

"""**Dealing with NA values**"""

# show null/NA values per column
df.isnull().sum()

# show NA values as % of total observations per column
df.isnull().sum()*100/len(df)

# drop all rows containing null
df.dropna()
```

```
# drop all columns containing null
df.dropna(axis=1)

# drop columns with less than 5 NA values
df.dropna(axis=1, thresh=5)

# replace all na values with -9999
df.fillna(-9999)

# fill na values with NaN
df.fillna(np.NaN)

# fill na values with strings
df.fillna("data missing")

# fill missing values with mean column values
df.fillna(df.mean())

*****Column Operation*****

# select a column
df["sepal_length"]

# select multiple columns and create a new dataframe X
X = df[["sepal_length", "sepal_width", "species"]]
X

# select a column by column number
df.iloc[:, [1,3,4]]

# save all columns to a list
df.columns.tolist()

# sorting values by column "sepalW" in ascending order
df.sort_values(by = "sepal_width", ascending = True)

# add new calculated column
df['newcol'] = df["sepal_length"]*2
df

# create a conditional calculated column
df['newcol'] = ["short" if i<3 else "long" for i in df["sepal_width"]]
df

*****Row Operation (Sort, Filter, Slice)*****

# select rows 3 to 10
df.iloc[3:10,]

# select rows 3 to 49 and columns 1 to 3
```

```
df.iloc[3:50, 1:4]

# randomly select 10 rows
df.sample(10)

# find rows with specific strings
df[df["species"].isin(["setosa"])]

# conditional filtering
df[df.sepal_length >= 5]

# filtering rows with multiple values e.g. 0.2, 0.3
df[df["petal_width"].isin([0.2, 0.3])]

# multi-conditional filtering
df[(df.petal_length > 1) & (df.species=="setosa") | (df.sepal_width < 3)]

# drop rows
df.drop(df.index[1]) # 1 is row index to be deleted

"""**Grouping**"""

# data grouped by column "species"
X = df.groupby("species")
X

# return mean values of a column ("sepal_length" ) grouped by "species" column
df.groupby("species")["sepal_length"].mean()

# return mean values of ALL columns grouped by "species" category
df.groupby("species").mean()

# get counts in different categories
df.groupby("species").nunique()
```

Employee Dataset:

Employee dataset contains columns such as first name, gender, start date, last login, salary bonus, senior management and team. Some of the fields are null in the dataset.

```
import pandas as pd
df = pd.read_csv("C:/Users/Vaishali/Desktop/AI-DS/employees.csv")
print(df)

print(df.describe())

#print(df.isNull())

print(pd.isnull(df['Team']))
```

```
print(pd.notnull(df['Team']))

print(df.fillna(1111))

print(df.fillna(method='pad'))

#import pandas as pd
#df=pd.read_csv("employees.csv")
print(df)

df.fillna(method='bfill') # check the output
print(df)

df['Gender'].fillna("No Gender",inplace=True)
print(df)

import numpy as np
print(df.replace(to_replace=np.NaN,value="SFIT"))

print(df)

print(df.interpolate(method='linear',limit_direction='forward'))

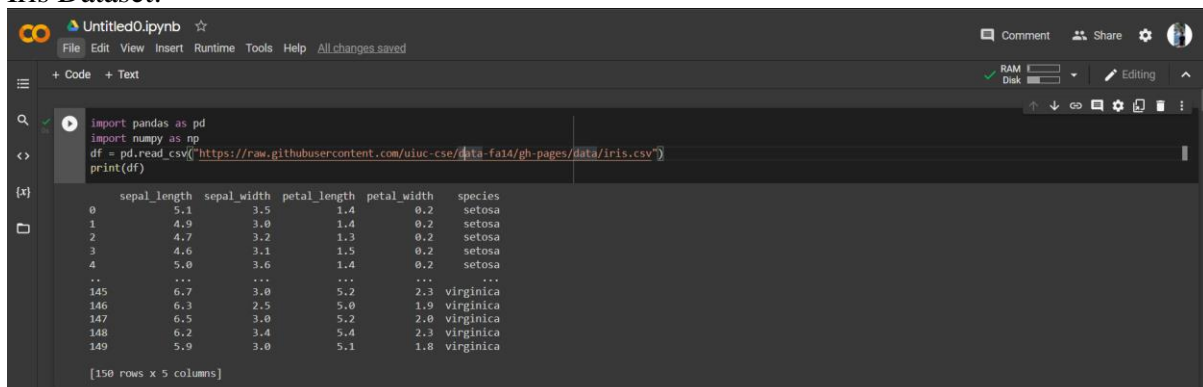
df1=pd.DataFrame({"A":[12,23,5,6,6,6],
                  "B":[34,2,34,5,67],
                  "C":[67,54,33,77,98],
                  "D":[45,87,65,33,23,23]
                  })
print(df1)

print(df1.interpolate(method='linear',limit_direction='forward'))

print(df1.dropna())
```

Paste Screenshots of above commands.

Iris Dataset:



```
import pandas as pd
import numpy as np
df = pd.read_csv("https://raw.githubusercontent.com/uiuc-cse/data-fa14/gh-pages/data/iris.csv")
print(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]

```
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+ Code + Text
[ ] 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.head(6)

   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
```

```
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+ Code + Text
[ ] df.shape

(150, 5)

[ ] 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

[ ] df.columns

Index(['sepal_length', 'sepal_width', 'petal_length', 'petal_width',
       'species'],
      dtype='object')
```

```
Untitled0.ipynb
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+ Code + Text
[ ] df['sepal_length'].nunique()

35

[ ] df['sepal_length'].unique()

array([5.1, 4.9, 4.7, 4.6, 5. , 5.4, 4.4, 4.8, 4.3, 5.8, 5.7, 5.2, 5.5,
       4.5, 5.3, 7. , 6.4, 6.9, 6.5, 6.3, 6.6, 5.9, 6. , 6.1, 5.6, 6.7,
       6.2, 6.8, 7.1, 7.6, 7.3, 7.2, 7.7, 7.4, 7.9])

[ ] df.columns.nunique()

5

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

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

```
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+ Code + Text
[ ] 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
```

```
Untitled0.ipynb ☆
File Edit View Insert Runtime Tools Help All changes saved
+ Code + Text
[ ] df.isnull().sum()*100/len(df)

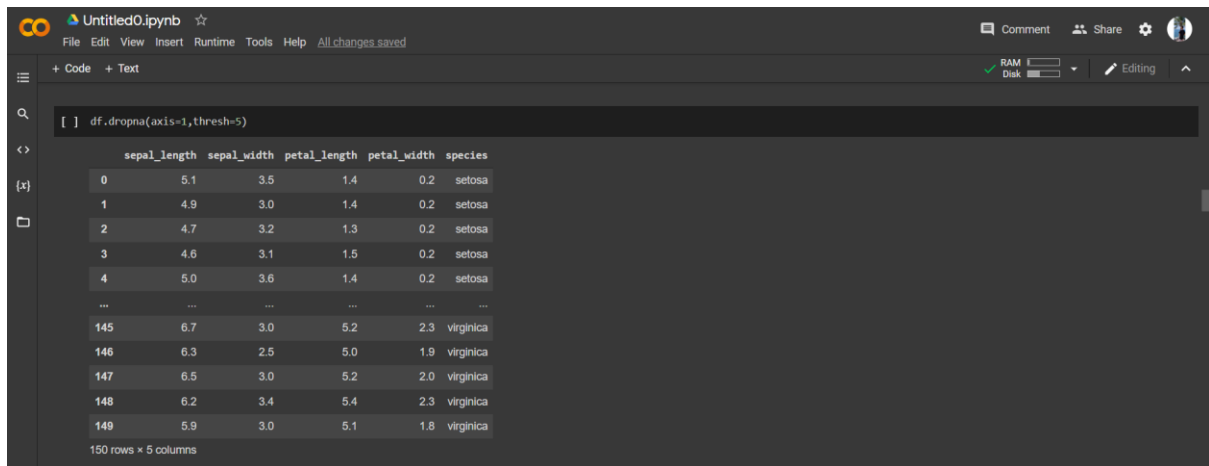
sepal_length 0.0
sepal_width 0.0
petal_length 0.0
petal_width 0.0
species 0.0
dtype: float64

[ ] df.dropna()

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
150 rows x 5 columns
```

```
Untitled0.ipynb ☆
File Edit View Insert Runtime Tools Help All changes saved
+ Code + Text
150 rows x 5 columns
[ ] df.dropna(axis=1)

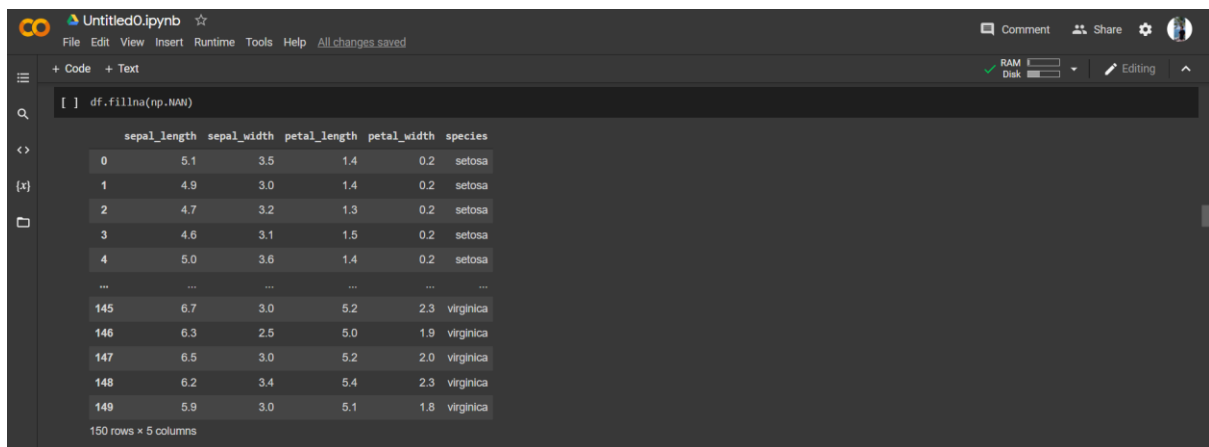
sepal_length sepal_width petal_length petal_width species
0 5.1 3.5 1.4 0.2 setosa
1 4.9 3.0 1.4 0.2 setosa
2 4.7 3.2 1.3 0.2 setosa
3 4.6 3.1 1.5 0.2 setosa
4 5.0 3.6 1.4 0.2 setosa
...
145 6.7 3.0 5.2 2.3 virginica
146 6.3 2.5 5.0 1.9 virginica
147 6.5 3.0 5.2 2.0 virginica
148 6.2 3.4 5.4 2.3 virginica
149 5.9 3.0 5.1 1.8 virginica
150 rows x 5 columns
```



```
[ ] df.dropna(axis=1, thresh=5)
```

| | 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 |

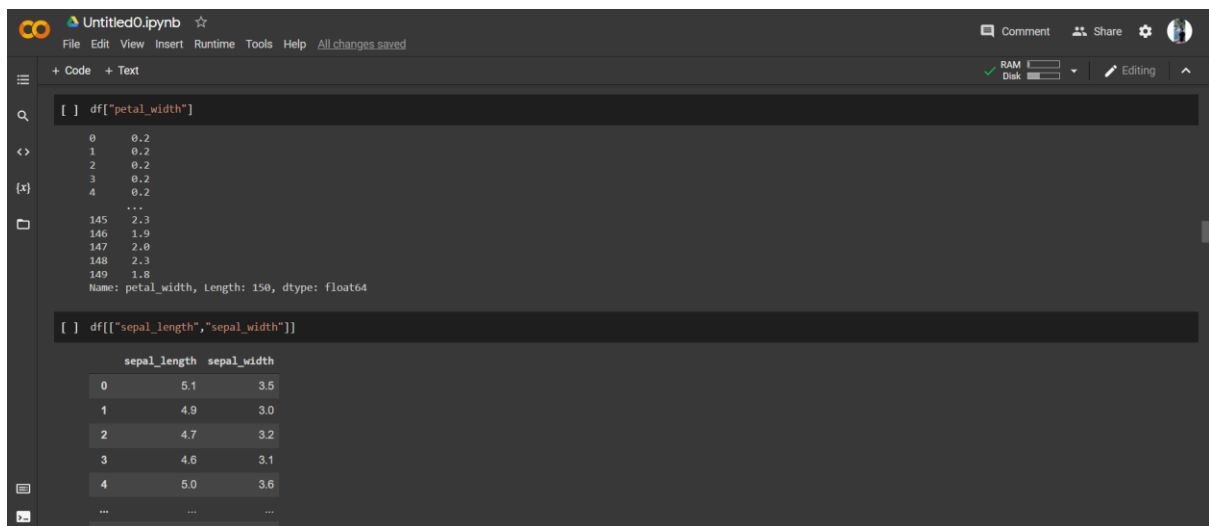
150 rows x 5 columns



```
[ ] df.fillna(np.NaN)
```

| | 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[\"petal_width\"]
```

0 0.2
1 0.2
2 0.2
3 0.2
4 0.2
...
145 2.3
146 1.9
147 2.0
148 2.3
149 1.8
Name: petal_width, Length: 150, dtype: float64

```
[ ] df[[\"sepal_length\", \"sepal_width\"]]
```

| | sepal_length | sepal_width |
|-----|--------------|-------------|
| 0 | 5.1 | 3.5 |
| 1 | 4.9 | 3.0 |
| 2 | 4.7 | 3.2 |
| 3 | 4.6 | 3.1 |
| 4 | 5.0 | 3.6 |
| ... | ... | ... |
| 145 | 6.7 | 3.0 |
| 146 | 6.3 | 2.5 |
| 147 | 6.5 | 3.0 |
| 148 | 6.2 | 3.4 |
| 149 | 5.9 | 3.0 |

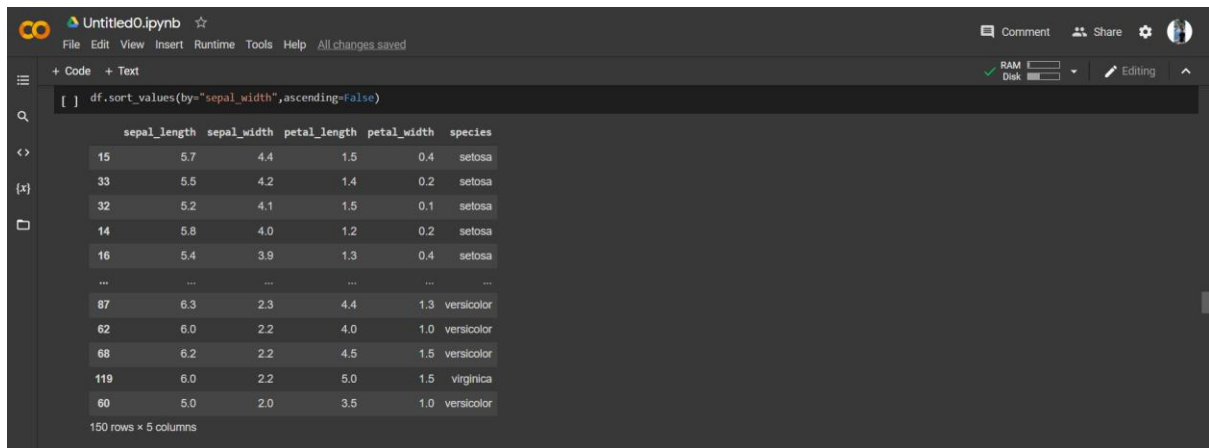
The screenshot shows a Jupyter Notebook window titled 'Untitled0.ipynb'. The top bar includes a menu (File, Edit, View, Insert, Runtime, Tools, Help) and a status indicator 'All changes saved'. The left sidebar contains navigation icons. The top right corner shows system status (RAM, Disk) and user options (Comment, Share, Settings, Profile). The main area displays a code cell with the following Python code:

```
df.sort_values(by="sepal_width", ascending=True)
```

Below the code cell, the resulting DataFrame is displayed as a table with 5 columns: sepal_length, sepal_width, petal_length, petal_width, and species. The rows are sorted by sepal_width in ascending order. The table shows 15 rows of data, with the last row indicating '150 rows x 5 columns'.

| | 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

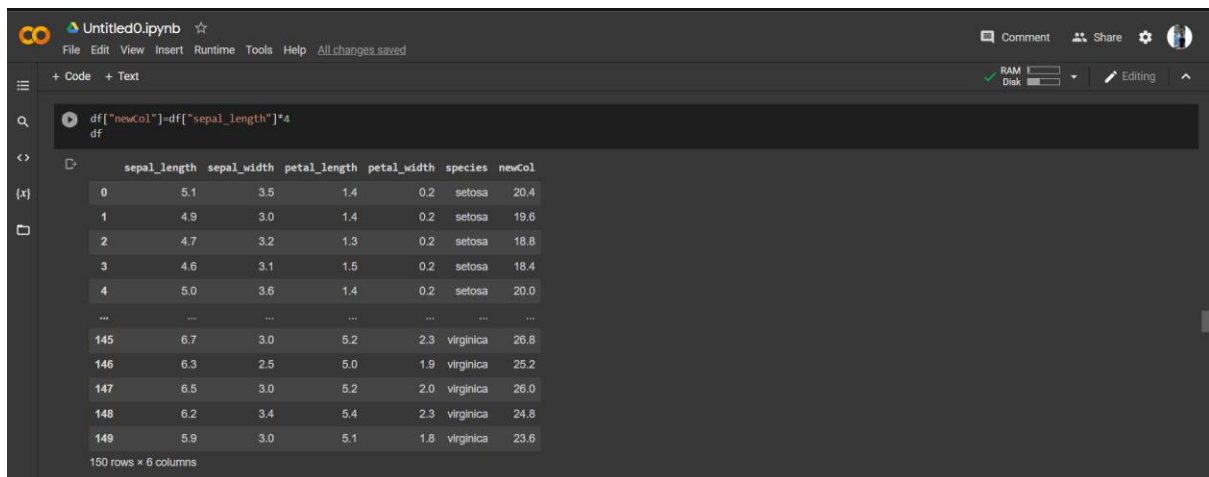


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```
[ ] df.sort_values(by="sepal_width",ascending=False)
```

| | sepal_length | sepal_width | petal_length | petal_width | species |
|-----|--------------|-------------|--------------|-------------|------------|
| 15 | 5.7 | 4.4 | 1.5 | 0.4 | setosa |
| 33 | 5.5 | 4.2 | 1.4 | 0.2 | setosa |
| 32 | 5.2 | 4.1 | 1.5 | 0.1 | setosa |
| 14 | 5.8 | 4.0 | 1.2 | 0.2 | setosa |
| 16 | 5.4 | 3.9 | 1.3 | 0.4 | setosa |
| ... | ... | ... | ... | ... | ... |
| 87 | 6.3 | 2.3 | 4.4 | 1.3 | versicolor |
| 62 | 6.0 | 2.2 | 4.0 | 1.0 | versicolor |
| 68 | 6.2 | 2.2 | 4.5 | 1.5 | versicolor |
| 119 | 6.0 | 2.2 | 5.0 | 1.5 | virginica |
| 60 | 5.0 | 2.0 | 3.5 | 1.0 | versicolor |

150 rows x 5 columns

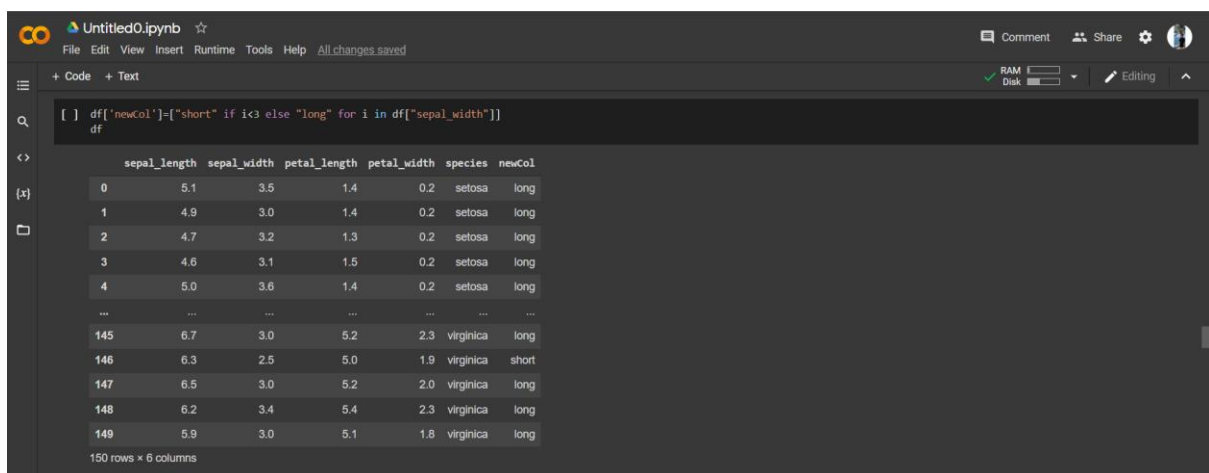


Untitled0.ipynb

```
df["newCol"]=df["sepal_length"]*4  
df
```

| | sepal_length | sepal_width | petal_length | petal_width | species | newCol |
|-----|--------------|-------------|--------------|-------------|-----------|--------|
| 0 | 5.1 | 3.5 | 1.4 | 0.2 | setosa | 20.4 |
| 1 | 4.9 | 3.0 | 1.4 | 0.2 | setosa | 19.6 |
| 2 | 4.7 | 3.2 | 1.3 | 0.2 | setosa | 18.8 |
| 3 | 4.6 | 3.1 | 1.5 | 0.2 | setosa | 18.4 |
| 4 | 5.0 | 3.6 | 1.4 | 0.2 | setosa | 20.0 |
| ... | ... | ... | ... | ... | ... | ... |
| 145 | 6.7 | 3.0 | 5.2 | 2.3 | virginica | 26.8 |
| 146 | 6.3 | 2.5 | 5.0 | 1.9 | virginica | 25.2 |
| 147 | 6.5 | 3.0 | 5.2 | 2.0 | virginica | 26.0 |
| 148 | 6.2 | 3.4 | 5.4 | 2.3 | virginica | 24.8 |
| 149 | 5.9 | 3.0 | 5.1 | 1.8 | virginica | 23.6 |

150 rows x 6 columns

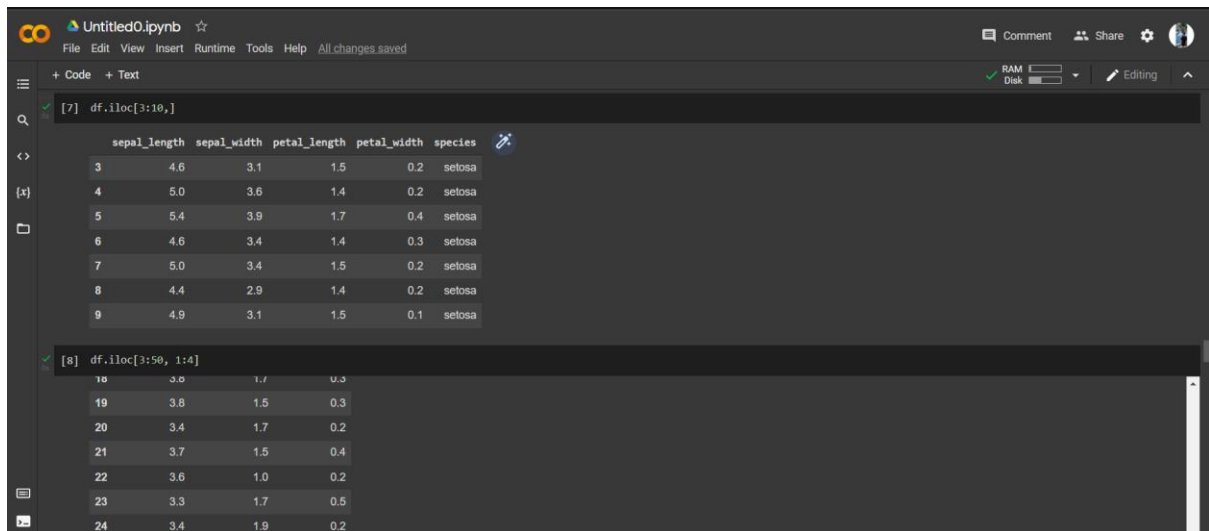


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```
[ ] df["newCol"]="short" if i<3 else "long" for i in df["sepal_width"]  
df
```

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

150 rows x 6 columns

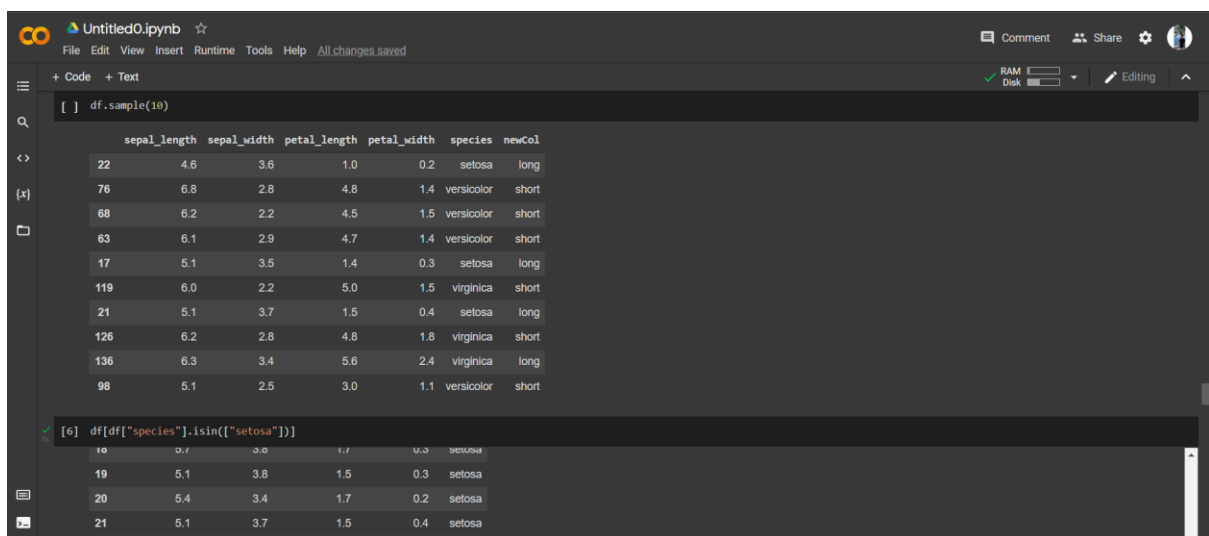


Code cell [7]: `df.iloc[3:10,]`

| | sepal_length | sepal_width | petal_length | petal_width | species |
|---|--------------|-------------|--------------|-------------|---------|
| 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 |

Code cell [8]: `df.iloc[3:50, 1:4]`

| | sepal_width | petal_length | petal_width |
|----|-------------|--------------|-------------|
| 10 | 3.0 | 1.7 | 0.3 |
| 19 | 3.8 | 1.5 | 0.3 |
| 20 | 3.4 | 1.7 | 0.2 |
| 21 | 3.7 | 1.5 | 0.4 |
| 22 | 3.6 | 1.0 | 0.2 |
| 23 | 3.3 | 1.7 | 0.5 |
| 24 | 3.4 | 1.9 | 0.2 |

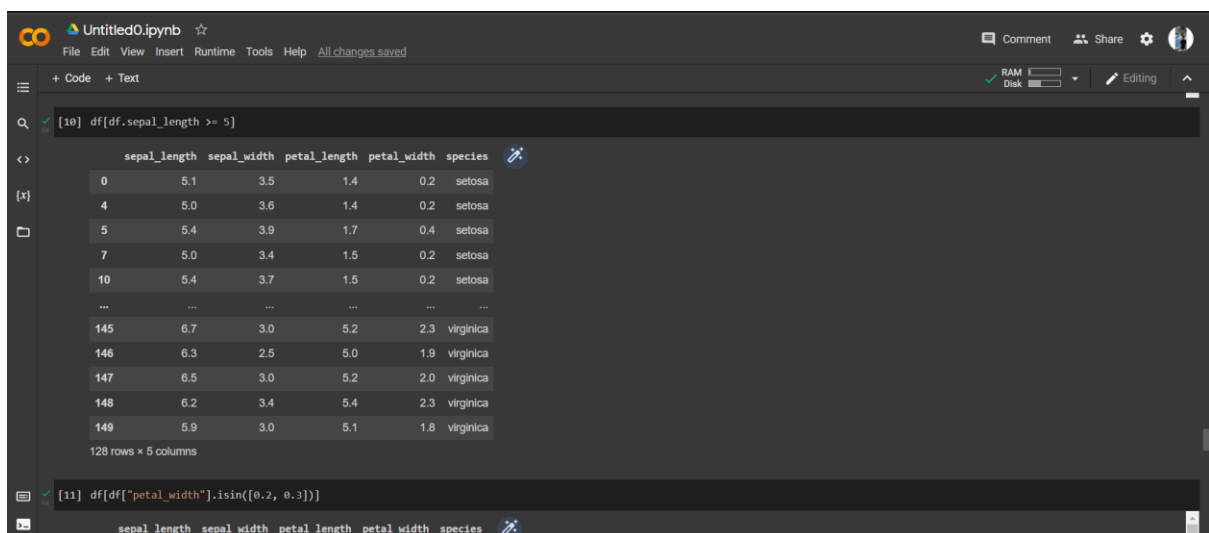


Code cell []: `df.sample(10)`

| | sepal_length | sepal_width | petal_length | petal_width | species | newCol |
|-----|--------------|-------------|--------------|-------------|------------|--------|
| 22 | 4.6 | 3.6 | 1.0 | 0.2 | setosa | long |
| 76 | 6.8 | 2.8 | 4.8 | 1.4 | versicolor | short |
| 68 | 6.2 | 2.2 | 4.5 | 1.5 | versicolor | short |
| 63 | 6.1 | 2.9 | 4.7 | 1.4 | versicolor | short |
| 17 | 5.1 | 3.5 | 1.4 | 0.3 | setosa | long |
| 119 | 6.0 | 2.2 | 5.0 | 1.5 | virginica | short |
| 21 | 5.1 | 3.7 | 1.5 | 0.4 | setosa | long |
| 126 | 6.2 | 2.8 | 4.8 | 1.8 | virginica | short |
| 136 | 6.3 | 3.4 | 5.6 | 2.4 | virginica | long |
| 98 | 5.1 | 2.5 | 3.0 | 1.1 | versicolor | short |

Code cell [6]: `df[df['species'].isin(['setosa'])]`

| | sepal_length | sepal_width | petal_length | petal_width | species |
|----|--------------|-------------|--------------|-------------|---------|
| 10 | 4.7 | 3.0 | 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 |



Code cell [10]: `df[df.sepal_length >= 5]`

| | sepal_length | sepal_width | petal_length | petal_width | species |
|-----|--------------|-------------|--------------|-------------|-----------|
| 0 | 5.1 | 3.5 | 1.4 | 0.2 | setosa |
| 4 | 5.0 | 3.6 | 1.4 | 0.2 | setosa |
| 5 | 5.4 | 3.9 | 1.7 | 0.4 | setosa |
| 7 | 5.0 | 3.4 | 1.5 | 0.2 | setosa |
| 10 | 5.4 | 3.7 | 1.5 | 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 |

128 rows x 5 columns

Code cell [11]: `df[df['petal_width'].isin([0.2, 0.3])]`

| | sepal_length | sepal_width | petal_length | petal_width | species |
|--|--------------|-------------|--------------|-------------|---------|
|--|--------------|-------------|--------------|-------------|---------|

```
Untitled0.ipynb
File Edit View Insert Runtime Tools Help All changes saved
+ Code + Text
[14] df[(df.petal_length > 1) & (df.species=="setosa") | (df.sepal_width < 3)]

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
...
132 6.4 2.8 5.6 2.2 virginica
133 6.3 2.8 5.1 1.5 virginica
134 6.1 2.6 5.6 1.4 virginica
142 5.8 2.7 5.1 1.9 virginica
146 6.3 2.5 5.0 1.9 virginica
104 rows x 5 columns

[15] df.drop(df.index[1])

sepal_length sepal_width petal_length petal_width species
```

```
Untitled0.ipynb
File Edit View Insert Runtime Tools Help All changes saved
+ Code + Text
[15] 149 rows x 5 columns

[19] X = df.groupby("species")
X
<pandas.core.groupby.generic.DataFrameGroupBy object at 0x7fc703c49850>

[20] df.groupby("species").mean()

sepal_length sepal_width petal_length petal_width
species
setosa 5.006 3.418 1.464 0.244
versicolor 5.936 2.770 4.260 1.326
virginica 6.588 2.974 5.552 2.026

[22] df.groupby("species").nunique()

sepal_length sepal_width petal_length petal_width
species
setosa 15 16 9 6
versicolor 21 14 19 9
```

Employee

```
employeedataset.ipynb
File Edit View Insert Runtime Tools Help All changes saved
+ Code + Text
[18] import pandas as pd
df = pd.read_csv("/employees.csv")
print(df)

First Name Gender ... Senior Management Team
0 Douglas Male ... True Marketing
1 Thomas Male ... True NaN
2 Maria Female ... False Finance
3 Jerry Male ... True Finance
4 Larry Male ... True NaN
...
995 Henry NaN ... False Distribution
996 Phillip Male ... False Finance
997 Russell Male ... False Product
998 Larry Male ... False Business Development
999 Albert Male ... True Sales
[1000 rows x 8 columns]

[ ] from google.colab import drive
drive.mount('/content/drive')

[19] print(df.describe())

Salary Bonus %
count 1000.000000 999.000000
mean 98662.181000 10.198672
std 32923.693342 5.524105
min 35813.000000 1.015000
25% 62613.000000 5.390500
50% 98428.000000 9.828000
75% 118740.250000 14.837000
```

```
employeedataset.ipynb
File Edit View Insert Runtime Tools Help All changes saved
+ Code + Text
[20] print(pd.isnull(df['Team']))
0      False
1       True
2      False
3      False
4       True
...
995     False
996     False
997     False
998     False
999     False
Name: Team, Length: 1000, dtype: bool

[21] print(pd.notnull(df['Team']))
0       True
1      False
2       True
3       True
4      False
...
995       True
996       True
997       True
998       True
999       True
Name: Team, Length: 1000, dtype: bool
```

```
employeedataset.ipynb
File Edit View Insert Runtime Tools Help All changes saved
+ Code + Text
[23] print(df.fillna(1111))
   First Name  Gender  ... Senior Management  Team
0   Douglas    Male  ...          True      Marketing
1   Thomas    Male  ...          True      1111
2   Maria     Female  ...         False      Finance
3   Jerry     Male  ...          True      Finance
4   Larry     Male  ...          True      1111
...
995   Henry    1111  ...         False      Distribution
996  Phillip    Male  ...         False      Finance
997  Russell    Male  ...         False      Product
998   Larry     Male  ...         False  Business Development
999  Albert     Male  ...          True        Sales

[1000 rows x 8 columns]

[22] print(df.fillna(method='pad'))
   First Name  Gender  ... Senior Management  Team
0   Douglas    Male  ...          True      Marketing
1   Thomas    Male  ...          True      Marketing
2   Maria     Female  ...         False      Finance
3   Jerry     Male  ...          True      Finance
4   Larry     Male  ...          True      Finance
...
995   Henry    Male  ...         False      Distribution
996  Phillip    Male  ...         False      Finance
997  Russell    Male  ...         False      Product
998   Larry     Male  ...         False  Business Development
999  Albert     Male  ...          True        Sales

[1000 rows x 8 columns]
```

```
employeedataset.ipynb
File Edit View Insert Runtime Tools Help All changes saved
+ Code + Text
[24] df.fillna(method='bfill')
print(df)
   First Name  Gender  ... Senior Management  Team
0   Douglas    Male  ...          True      Marketing
1   Thomas    Male  ...          True      NaN
2   Maria     Female  ...         False      Finance
3   Jerry     Male  ...          True      Finance
4   Larry     Male  ...          True      NaN
...
995   Henry     NaN  ...         False      Distribution
996  Phillip    Male  ...         False      Finance
997  Russell    Male  ...         False      Product
998   Larry     Male  ...         False  Business Development
999  Albert     Male  ...          True        Sales

[1000 rows x 8 columns]

[25] df['Gender'].fillna('No Gender', inplace=True)
print(df)
   First Name  Gender  ... Senior Management  Team
0   Douglas    Male  ...          True      Marketing
1   Thomas    Male  ...          True      NaN
2   Maria     Female  ...         False      Finance
3   Jerry     Male  ...          True      Finance
4   Larry     Male  ...          True      NaN
...
995   Henry  No Gender  ...         False      Distribution
996  Phillip    Male  ...         False      Finance
997  Russell    Male  ...         False      Product
998   Larry     Male  ...         False  Business Development
999  Albert     Male  ...          True        Sales
```

The screenshot shows a Jupyter Notebook with the following code and output:

```
[30] import numpy as np
print(df.replace(to_replace=np.NaN,value="SFIT"))
```

| | First Name | Gender | ... | Senior Management | Team |
|-----|------------|-----------|-----|-------------------|----------------------|
| 0 | Douglas | Male | ... | True | Marketing |
| 1 | Thomas | Male | ... | True | SFIT |
| 2 | Maria | Female | ... | False | Finance |
| 3 | Jerry | Male | ... | True | Finance |
| 4 | Larry | Male | ... | True | SFIT |
| ... | ... | ... | ... | ... | ... |
| 995 | Henry | No Gender | ... | False | Distribution |
| 996 | Phillip | Male | ... | False | Finance |
| 997 | Russell | Male | ... | False | Product |
| 998 | Larry | Male | ... | False | Business Development |
| 999 | Albert | Male | ... | True | Sales |

[1000 rows x 8 columns]

```
[31] print(df.interpolate(method="linear",limit_direction="forward"))
```

| | First Name | Gender | ... | Senior Management | Team |
|-----|------------|-----------|-----|-------------------|----------------------|
| 0 | Douglas | Male | ... | True | Marketing |
| 1 | Thomas | Male | ... | True | NaN |
| 2 | Maria | Female | ... | False | Finance |
| 3 | Jerry | Male | ... | True | Finance |
| 4 | Larry | Male | ... | True | NaN |
| ... | ... | ... | ... | ... | ... |
| 995 | Henry | No Gender | ... | False | Distribution |
| 996 | Phillip | Male | ... | False | Finance |
| 997 | Russell | Male | ... | False | Product |
| 998 | Larry | Male | ... | False | Business Development |
| 999 | Albert | Male | ... | True | Sales |

The screenshot shows a Jupyter Notebook with the following code and output:

```
[32] df1=pd.DataFrame({"A":[12,23,None,5,6,None],
                        "B":[34,None,2,34,5,67],
                        "C":[67,54,33,None,77,98],
                        "D":[45,87,65,33,23,None]
                      })
print(df1)
```

| | A | B | C | D |
|---|------|------|------|------|
| 0 | 12.0 | 34.0 | 67.0 | 45.0 |
| 1 | 23.0 | NaN | 54.0 | 87.0 |
| 2 | NaN | 2.0 | 33.0 | 65.0 |
| 3 | 5.0 | 34.0 | NaN | 33.0 |
| 4 | 6.0 | 5.0 | 77.0 | 23.0 |
| 5 | NaN | 67.0 | 98.0 | NaN |

```
[33] print(df1.interpolate(method="linear",limit_direction="forward"))
```

| | A | B | C | D |
|---|------|------|------|------|
| 0 | 12.0 | 34.0 | 67.0 | 45.0 |
| 1 | 23.0 | 18.0 | 54.0 | 87.0 |
| 2 | 14.0 | 2.0 | 33.0 | 65.0 |
| 3 | 5.0 | 34.0 | 55.0 | 33.0 |
| 4 | 6.0 | 5.0 | 77.0 | 23.0 |
| 5 | 6.0 | 67.0 | 98.0 | 23.0 |

```
print(df1.dropna())
```

| | A | B | C | D |
|---|------|------|------|------|
| 0 | 12.0 | 34.0 | 67.0 | 45.0 |
| 4 | 6.0 | 5.0 | 77.0 | 23.0 |

8. Post-Experiments Exercise

A. Extended Theory: (Soft Copy)

How to handle missing data in dataset? (Use Diabetes dataset & reference link)

Pandas treat None and NaN as essentially interchangeable for indicating missing or null values. To facilitate this convention, there are several useful functions for detecting,

removing, and replacing null values in Pandas DataFrame :

- isnull()
- notnull()
- dropna()
- fillna()
- replace
- interpolate()

Diabetes dataset eg

```
diabetes.ipynb
File Edit View Insert Runtime Tools Help All changes saved
+ Code + Text
import pandas as pd
import numpy as np
df = pd.read_csv("https://raw.githubusercontent.com/jbrownlee/datasets/master/pima-indians-diabetes.csv")
print(df)

6 148 72 35 0 33.6 0.627 50 1
0 1 85 66 29 0 26.6 0.351 31 0
1 8 183 64 0 0 23.3 0.672 32 1
2 1 89 66 23 94 28.1 0.167 21 0
3 0 137 40 35 168 43.1 2.288 33 1
4 5 116 74 0 0 25.6 0.201 30 0
.. ..
762 10 101 76 48 180 32.9 0.171 63 0
763 2 122 70 27 0 36.8 0.340 27 0
764 5 121 72 23 112 26.2 0.245 30 0
765 1 126 60 0 0 30.1 0.349 47 1
766 1 93 70 31 0 30.4 0.315 23 0

[767 rows x 9 columns]

[4] df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 767 entries, 0 to 766
Data columns (total 9 columns):
 # column Non-Null count Dtype
---
0 6 767 non-null int64
1 148 767 non-null int64
2 72 767 non-null int64
3 35 767 non-null int64
4 0 767 non-null int64
5 33.6 767 non-null float64
```

```
diabetes.ipynb
File Edit View Insert Runtime Tools Help All changes saved
+ Code + Text
[5] df.shape
(767, 9)

[7] df.describe()

count 767.000000 767.000000 767.000000 767.000000 767.000000 767.000000 767.000000 767.000000 767.000000
mean 3.842243 120.859192 69.101695 20.517601 79.903520 31.990482 0.471674 33.219035 0.348110
std 3.370877 31.978468 19.368155 15.954059 115.283105 7.889091 0.331497 11.752296 0.476682
min 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.078000 21.000000 0.000000
25% 1.000000 99.000000 62.000000 0.000000 0.000000 27.300000 0.243500 24.000000 0.000000
50% 3.000000 117.000000 72.000000 23.000000 32.000000 32.000000 0.371000 29.000000 0.000000
75% 6.000000 140.000000 80.000000 32.000000 127.500000 36.600000 0.625000 41.000000 1.000000
max 17.000000 199.000000 122.000000 99.000000 846.000000 67.100000 2.420000 81.000000 1.000000

[8] df.head(20)

6 148 72 35 0 33.6 0.627 50 1
0 1 85 66 29 0 26.6 0.351 31 0
1 8 183 64 0 0 23.3 0.672 32 1
2 1 89 66 23 94 28.1 0.167 21 0
```

```
diabetes.ipynb
File Edit View Insert Runtime Tools Help All changes saved
+ Code + Text
[11] import pandas as pd
import numpy as np
df = pd.read_csv("https://raw.githubusercontent.com/jbrownlee/Datasets/master/pima-indians-diabetes.csv", header=None)
num_missing=(df[[1,2,3,4,5]]==0).sum()
print(num_missing)

1 5
2 35
3 227
4 374
5 11
dtype: int64

[16] df[[1,2,3,4,5]]=df[[1,2,3,4,5]].replace(0,np.nan)
print(df.isnull().sum())

0 0
1 5
2 35
3 227
4 374
5 11
6 0
7 0
8 0
dtype: int64

[15] print(df.head(20))

0 6 148.0 72.0 35.0 NaN 33.6 0.627 50 1
1 1 85.0 66.0 29.0 NaN 26.6 0.351 31 0
2 8 183.0 64.0 NaN NaN 23.3 0.672 32 1
```

```
diabetes.ipynb
File Edit View Insert Runtime Tools Help All changes saved
+ Code + Text
[22] df[6].nunique()
517
[23] df.columns.nunique()
9
[24] df[6].value_counts()
0.254    6
0.253    6
0.259    5
0.238    5
0.207    5
..
0.886    1
0.804    1
1.251    1
0.382    1
0.375    1
Name: 6, Length: 517, dtype: int64
[25] df.isnull()
   0  1  2  3  4  5  6  7  8
0  False False False False True False False False False
1  False False False False True False False False False
2  False False False True True False False False False
```

```
diabetes.ipynb
File Edit View Insert Runtime Tools Help All changes saved
+ Code + Text
[26] df.isnull().sum()
0    0
1     5
2    35
3   227
4   374
5     11
6      0
7      0
8      0
dtype: int64
[27] df.fillna("Data missing")
   0  1  2  3  4  5  6  7  8
0  6  148  72    35  Data missing  33.6  0.627  50  1
1  1  85  66    29  Data missing  26.6  0.351  31  0
2  8  183  64  Data missing  Data missing  23.3  0.672  32  1
3  1  89  66    23    94  28.1  0.167  21  0
4  0  137  40    35   168  43.1  2.288  33  1
...  ...  ...  ...  ...  ...  ...  ...  ...
763  10  101  76    48   180  32.9  0.171  63  0
764  2  122  70    27  Data missing  36.8  0.340  27  0
765  5  121  72    23   112  26.2  0.245  30  0
766  1  126  60  Data missing  Data missing  30.1  0.349  47  1
```

```
diabetes.ipynb
File Edit View Insert Runtime Tools Help All changes saved
+ Code + Text
[27] 767  1  93  70    31  Data missing  30.4  0.315  23  0
768 rows x 9 columns
[28] df.fillna(df.mean())
   0  1  2  3  4  5  6  7  8
0  6  148.0  72.0  35.000000  155.548223  33.6  0.627  50  1
1  1  85.0  66.0  29.000000  155.548223  26.6  0.351  31  0
2  8  183.0  64.0  29.15342  155.548223  23.3  0.672  32  1
3  1  89.0  66.0  23.000000  94.000000  28.1  0.167  21  0
4  0  137.0  40.0  35.000000  168.000000  43.1  2.288  33  1
...  ...  ...  ...  ...  ...  ...  ...  ...
763  10  101.0  76.0  48.000000  180.000000  32.9  0.171  63  0
764  2  122.0  70.0  27.000000  155.548223  36.8  0.340  27  0
765  5  121.0  72.0  23.000000  112.000000  26.2  0.245  30  0
766  1  126.0  60.0  29.15342  155.548223  30.1  0.349  47  1
767  1  93.0  70.0  31.000000  155.548223  30.4  0.315  23  0
768 rows x 9 columns
[ ]
```


B. Questions:

Mention types of data structures in Pandas.

Mention difference between Numpy and Pandas.

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| | |
|----------|--|
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| Date | |

Q.8. B M.

1) Mention types of data structures in Pandas.

→ Pandas is an open-source library that uses for working with relational or labeled both easily & intuitively. It supports two data structures.

1) Series

2) Dataframe.

1) Series

Series is one-dimensional labelled array capable of holding any data type (Integers, strings, floating point, numbers, etc) This axis labels are collectively referred to as index. The basic method to call

`s = pd.Series(data, index=index)`

Here, data can be many different things.

1) a python dict.

2) an ndarray

3) a scalar value (like 5)

The passed index, is a list of axis labels. Thus, this separates into a few cases depending on what data is:

2) Dataframe

Dataframe is a 2-dimensional labeled data structure with columns of potentially different types. You can think of it like a spreadsheet or SQL table or a dict of Series objects. It is generally the most commonly used Pandas object. Like Series

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| | |
|----------|--|
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Dataframes accepts many different kinds of Input

- Object of 1D ndarrays, dicts or Series
- 2-D numpy.ndarray
- A series
- Another Data frame.

2) Mention difference between Numpy & Pandas.

| Pandas | Numpy |
|---|---|
| 1) Pandas module works with the tabular data | 1) Numpy module works with numerical data |
| 2) Pandas has powerful tools like Series, DataFrames etc | 2) Numpy has a powerful tool like Arrays. |
| 3) Pandas is used in popular organizations like Instacart, Sendgrid & Sights. | 3) Numpy is used in popular organization like SweepSouth. |
| 4) Pandas has a better performance for 500k rows or more | 4) Numpy has a better performance for 50k rows or less. |
| 5) Pandas consume large memory as compared to numpy | 5) Numpy consumes less memory as compared to pandas. |
| 6) Pandas provides 2d table object called Dataframe | 6) Numpy provides a multi-dimensional array. |
| 7) Pandas uses R language as its reference language & hence provide many similar functions. | 7) Numpy is written in the C programming language & hence uses multiple function-ality from it. |

C. Conclusion:

Write the significance of the topic studied in the experiment.

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| | |
|----------|--|
| Page No. | |
| Date | |

Conclusion

In this experiment we learnt to how to use numpy and pandas libraries of Pandas and we ~~are~~ were able to gain understand and deal with missing data and inspect data. We also learnt rows and columns operations. We were able to apply the operations on various datasets and got the desired output.

CS Scanned with CamScanner

D. References:

1. [How to Handle Missing Data with Python \(machinelearningmastery.com\)](https://machinelearningmastery.com/how-to-handle-missing-data-with-python/)
 2. <https://www.w3schools.com/python/pandas>
 3. <https://www.geeksforgeeks.org/difference-between-pandas-vs-numpy/>
-