Experiment:1

Title: Data Pre-processing

Aim: To Collect, Clean, Integrate and Transform Healthcare Data based on specific disease < specify the name of the dataset here> using python.

Theory:

<Explanation about Data Pre processing tasks with example. >

Procedure:

Perform the following Pre -processing tasks on the chosen dataset

- 1. Identify duplicate rows
- 2. Find the missing values and remove/replace
- 3. Replace the missing values of nominal attribute (if any) to NULL
- 4. Identify columns with very few values
- 5. Rename the nominal values of nominal attribute
- 6. Identify columns that contain single value
- 7. Perform transformation as per data attributes

Output:

<Provide the snapshots for the above pre-processing steps>

Conclusion:

References:

https://www.javatpoint.com/data-preprocessing-machine-learning

https://serokell.io/blog/data-preprocessing

 $\underline{https://vitalflux.com/data-preprocessing\text{-}steps\text{-}in\text{-}machine\text{-}learning/}$

https://archive.ics.uci.edu/datasets

Sample pre-processing tasks solved using python

1. Load any dataset, identify the irrelevant columns for specific analysis and drop them.

Code:

```
import pandas as pd
df = pd.read_csv('hepatitis.csv')
irrelevant_cols = ['histology',]
df = df.drop(columns=irrelevant_cols)df.head(10)
```

2. Find the rows with missing values and replace with specific string or value

```
Code:
```

```
df = df.fillna(value=-1)
df.head(10)
```

3. Add any 2 dummy attributes in your dataset

Code:

```
import numpy as np
df['dummy1'] = np.random.rand(len(df))
df['dummy2'] = np.random.randint(0, 2, size=len(df))
df.head(10)
```

4. Apply any 2 standard normalization techniques on numeric attributes of the sampledataset chosen.

Code:

```
from sklearn.preprocessing import StandardScaler, MinMaxScaler
numeric_cols = ['alk_phosphate', 'sgot']
scaler1 = StandardScaler()
df[numeric_cols] = scaler1.fit_transform(df[numeric_cols])
print(df.head())
```