

Handling Missing Vaues

```
In [1]: import pandas as pd
```

```
In [2]: df=pd.read_csv(r'C:\Users\User\Downloads\archive (1)\diabetes.csv')
```

```
In [3]: df
```

```
Out[3]:
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome
0	6	148	72	35	0	33.6	0.627	50	1
1	1	85	66	29	0	26.6	0.351	31	0
2	8	183	64	0	0	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	0	36.8	0.340	27	0
765	5	121	72	23	112	26.2	0.245	30	0
766	1	126	60	0	0	30.1	0.349	47	1
767	1	93	70	31	0	30.4	0.315	23	0

768 rows × 9 columns

```
In [4]: df.head()
```

```
Out[4]:
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome
0	6	148	72	35	0	33.6	0.627	50	1
1	1	85	66	29	0	26.6	0.351	31	0
2	8	183	64	0	0	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

```
In [5]: df.tail()
```

```
Out[5]:
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome
763	10	101	76	48	180	32.9	0.171	63	0
764	2	122	70	27	0	36.8	0.340	27	0
765	5	121	72	23	112	26.2	0.245	30	0
766	1	126	60	0	0	30.1	0.349	47	1
767	1	93	70	31	0	30.4	0.315	23	0

```
In [6]: df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 768 entries, 0 to 767
Data columns (total 9 columns):
#   Column                Non-Null Count  Dtype
---  -
0   Pregnancies            768 non-null    int64
1   Glucose                768 non-null    int64
2   BloodPressure          768 non-null    int64
3   SkinThickness          768 non-null    int64
4   Insulin                768 non-null    int64
5   BMI                    768 non-null    float64
6   DiabetesPedigreeFunction 768 non-null    float64
7   Age                    768 non-null    int64
8   Outcome                768 non-null    int64
dtypes: float64(2), int64(7)
memory usage: 54.1 KB
```

```
In [7]: df.describe()
```

Out[7]:

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome
count	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000
mean	3.845052	120.894531	69.105469	20.536458	79.799479	31.992578	0.471876	33.240885	0.350277
std	3.369578	31.972618	19.355807	15.952218	115.244002	7.884160	0.331329	11.760232	0.486568
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	0.243750	24.000000	0.000000
50%	3.000000	117.000000	72.000000	23.000000	30.500000	32.000000	0.372500	29.000000	0.000000
75%	6.000000	140.250000	80.000000	32.000000	127.250000	36.600000	0.626250	41.000000	1.000000
max	17.000000	199.000000	122.000000	99.000000	846.000000	67.100000	2.420000	81.000000	1.000000

In [11]:

```
data_colmns=['Glucose','BloodPressure','SkinThickness','Insulin','BMI']
data_colmns
```

Out[11]:

['Glucose', 'BloodPressure', 'SkinThickness', 'Insulin', 'BMI']

In [12]:

```
(df[data_colmns]==0).sum()
```

Out[12]:

Glucose5
BloodPressure35
SkinThickness227
Insulin374
BMI11
dtype: int64

In [14]:

```
df.isnull().sum()
```

Out[14]:

Pregnancies0
Glucose0
BloodPressure0
SkinThickness0
Insulin0
BMI0
DiabetesPedigreeFunction0
Age0
Outcome0
dtype: int64

In [17]:

```
df[data_colmns]=df[data_colmns].replace(0,nan)
```

In [16]:

```
from numpy import nan
```

In [18]:

```
df.head(20)
```

Out[18]:

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome
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
3	1	89.0	66.0	23.0	94.0	28.1	0.167	21	0
4	0	137.0	40.0	35.0	168.0	43.1	2.288	33	1
5	5	116.0	74.0	NaN	NaN	25.6	0.201	30	0
6	3	78.0	50.0	32.0	88.0	31.0	0.248	26	1
7	10	115.0	NaN	NaN	NaN	35.3	0.134	29	0
8	2	197.0	70.0	45.0	543.0	30.5	0.158	53	1
9	8	125.0	96.0	NaN	NaN	NaN	0.232	54	1
10	4	110.0	92.0	NaN	NaN	37.6	0.191	30	0
11	10	168.0	74.0	NaN	NaN	38.0	0.537	34	1
12	10	139.0	80.0	NaN	NaN	27.1	1.441	57	0
13	1	189.0	60.0	23.0	846.0	30.1	0.398	59	1
14	5	166.0	72.0	19.0	175.0	25.8	0.587	51	1
15	7	100.0	NaN	NaN	NaN	30.0	0.484	32	1
16	0	118.0	84.0	47.0	230.0	45.8	0.551	31	1
17	7	107.0	74.0	NaN	NaN	29.6	0.254	31	1
18	1	103.0	30.0	38.0	83.0	43.3	0.183	33	0
19	1	115.0	70.0	30.0	96.0	34.6	0.529	32	1

Deleting Missing Values

In [19]:

df.isnull().sum()

Out[19]:

Pregnancies	0
Glucose	5
BloodPressure	35
SkinThickness	227
Insulin	374
BMI	11
DiabetesPedigreeFunction	0
Age	0
Outcome	0
dtype: int64	

As we see above Glucose,Blood Pressure and BMI has less missing values, so we can use the dropna() method.It would not detrimental our dataset

In [21]:

df= df.dropna(subset=['Glucose', 'BloodPressure', 'BMI'])
df

Out[21]:

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome
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
3	1	89.0	66.0	23.0	94.0	28.1	0.167	21	0
4	0	137.0	40.0	35.0	168.0	43.1	2.288	33	1
...
763	10	101.0	76.0	48.0	180.0	32.9	0.171	63	0
764	2	122.0	70.0	27.0	NaN	36.8	0.340	27	0
765	5	121.0	72.0	23.0	112.0	26.2	0.245	30	0
766	1	126.0	60.0	NaN	NaN	30.1	0.349	47	1
767	1	93.0	70.0	31.0	NaN	30.4	0.315	23	0

724 rows × 9 columns

In [22]:

df.isnull().sum()

```
Out[22]: Pregnancies      0
         Glucose          0
         BloodPressure    0
         SkinThickness    192
         Insulin          332
         BMI              0
         DiabetesPedigreeFunction  0
         Age              0
         Outcome          0
         dtype: int64
```

As we see the null values have been gone in the Glucose,Bloodpressure,BMI. But in some columns like SkinThickness and Insulin have more missing values

Replacing the missing values

The column 'Insulin' has 332 missing values,since it is a big number we would like to perform mean value of that column to replace missing value

```
In [23]: mean_val=df['Insulin'].mean()
         mean_val
```

```
Out[23]: 156.05612244897958
```

```
In [24]: df['Insulin'].fillna(mean_val,inplace=True)
```

C:\Users\User\AppData\Local\Temp\ipykernel_1620\1580020127.py:1: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame

See the caveats in the documentation: https://pandas.pydata.org/pandas-docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy
df['Insulin'].fillna(mean_val,inplace=True)

```
In [25]: df.isnull().sum()
```

```
Out[25]: Pregnancies      0
         Glucose          0
         BloodPressure    0
         SkinThickness    192
         Insulin          0
         BMI              0
         DiabetesPedigreeFunction  0
         Age              0
         Outcome          0
         dtype: int64
```

```
In [26]: df['SkinThickness'].interpolate(inplace=True)
```

C:\Users\User\AppData\Local\Temp\ipykernel_1620\4019890499.py:1: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame

See the caveats in the documentation: https://pandas.pydata.org/pandas-docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy
df['SkinThickness'].interpolate(inplace=True)

```
In [27]: df.isnull().sum()
```

```
Out[27]: Pregnancies      0
         Glucose          0
         BloodPressure    0
         SkinThickness    0
         Insulin          0
         BMI              0
         DiabetesPedigreeFunction  0
         Age              0
         Outcome          0
         dtype: int64
```

```
In [ ]:
```