# **Data Science Project**

#### Healthcare – Persistency of a Drug

#### Week 9 works

### Team member's details:

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Specialization: Data Science

#### Problem Description

To identify the persistency of a drug, a pharmaceutical company approached to develop a model based on data analysis. Factors that affect the persistence of drugs should be identified, along with data insights with predictive analytics, to help the company for their smooth and efficient functioning, with the help of dataset provided by the company.

#### Github Repo Link

<u>Final Project DS SS/week 9 at master · Soniyasunny1/Final Project DS SS (github.com)</u>

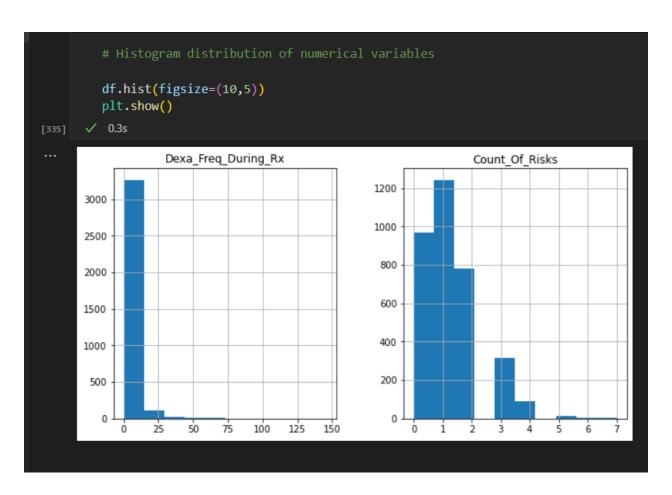
### Data Cleaning

- There are no null or nan values in the dataset. But there are some values like 'Unknown' or 'Others', which should be considered for transformation.
- In the 'Ntm\_Speciality' column, the values 'OBSTETRICS & OBSTETRICS & GYNECOLOGY & OBSTETRICS & GYNECOLOGY' and 'OBSTETRICS AND GYNECOLOGY' are changed to 'OBSTETRICS & GYNECOLOGY'. This helps to reduce the duplicates and to display charts properly when plotting count plot.
- Low frequency values were replaced with 'Other'. In 'Ntm\_Speciality' column, all values with value counts one or two were replaced with 'Other', to reduce the unique values in that column.

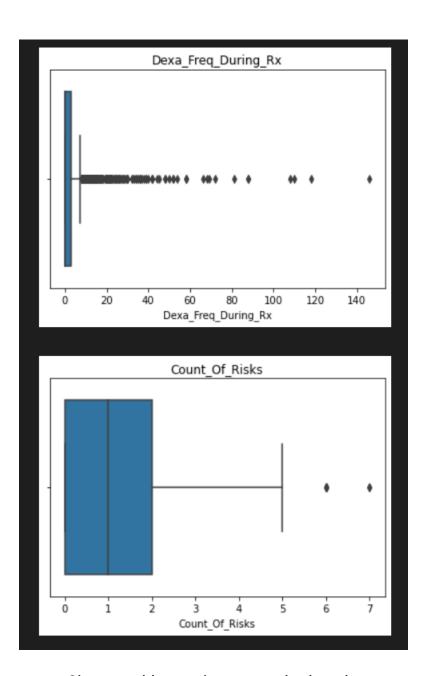
```
df["Ntm_Speciality"] = replace_low_freq(df, "Ntm_Speciality", 2, "Other")
        df["Ntm Speciality"].value counts()
[329] 	V 0.6s
    GENERAL PRACTITIONER
                                            1535
     RHEUMATOLOGY
                                             604
     ENDOCRINOLOGY
                                             458
     Unknown
                                             310
     ONCOLOGY
                                             225
     OBSTETRICS & GYNECOLOGY
                                              91
     UROLOGY
                                              33
     ORTHOPEDIC SURGERY
                                              30
     CARDIOLOGY
                                              22
     0ther
                                              22
     PATHOLOGY
                                              16
     HEMATOLOGY & ONCOLOGY
                                              14
     OTOLARYNGOLOGY
                                              14
     PEDIATRICS
                                              13
     PHYSICAL MEDICINE & REHABILITATION
                                              11
     SURGERY AND SURGICAL SPECIALTIES
                                               8
     PULMONARY MEDICINE
                                               8
     PSYCHIATRY AND NEUROLOGY
                                               4
     NEPHROLOGY
     ORTHOPEDICS
     Name: Ntm Speciality, dtype: int64
```

- The column 'Ptid' is dropped, as it is not useful for our analysis.
- Columns were classified into numerical and categorical based on dtypes.

- Histogram distribution of numerical variables were plotted.



- Box plot of numerical variables



#### - Skew and kurtosis were calculated

```
Dexa_Freq_During_Rx
skew kurtosis

Dexa_Freq_During_Rx 6.80873 74.758378

Count_Of_Risks
skew kurtosis

Count_Of_Risks 0.879791 0.900486
```

#### **Outlier Removal**

- Outliers were dropped based on inter-quartile range

```
Dexa_Freq_During_Rx
0.0
3.0
3.0
-4.5 7.5
Count_Of_Risks
0.0
2.0
2.0
-3.0 5.0
```

- After outlier removal, there is difference in skew and kurtosis.

```
Dexa_Freq_During_Rx

skew kurtosis

Dexa_Freq_During_Rx 2.249892 3.486526

Count_Of_Risks

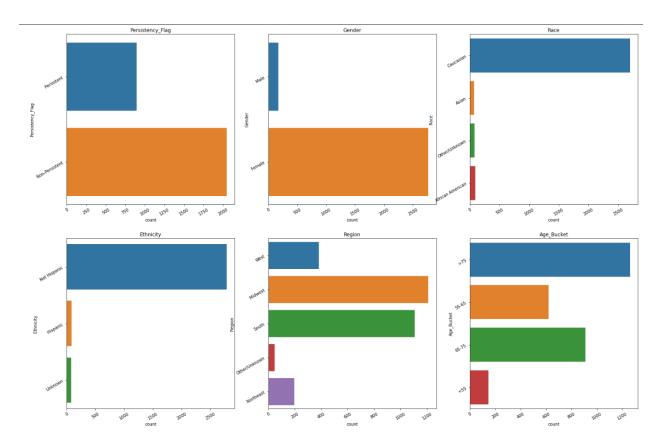
skew kurtosis

Count_Of_Risks 0.651706 -0.194379
```

- Minimum maximum normalization is also tried.

```
min_max_Dexa_Freq_During_Rx
skew kurtosis
min_max_Dexa_Freq_During_Rx 2.249892 3.486526
```

- Count plots of categorical features were also plotted in subplots.



# Mode based approach

Replace 'Unknown' values with mode.

```
templist=[]
# Replace unknown values with mode if it is less than 40% of total values
# Drop if it is greater than 40%
for l in unknown:
    val = tempdf[l].value_counts().Unknown
    if val>len(tempdf)*0.4:
        tempdf.drop(l , axis=1 , inplace=True)
        templist.append(l)
    else:
        tempdf[l].replace(to_replace='Unknown', value=tempdf[l].mode()[0], inplace=True)
    tempdf['Race'].replace(to_replace='Other/Unknown', value=tempdf['Race'].mode()[0], inplace=True)

> 0.1s
```

## Data Intake Report

Name: Data Science Final Project - 'Healthcare - Persistency of a Drug'

Report date: July 30, 2022

Internship Batch: LISUM10: 30

Version:<1.0>

Data intake by: Soniya Sunny

Data intake reviewer:<intern who reviewed the report>

Data storage location: <u>Healthcare dataset.xlsx - Google Drive</u>

#### Tabular data details:

Total number of observations	3425
Total number of files	1
Total number of features	69
Base format of the file	.xlsx
Size of the data	899 KB