

Data Science HealthCare Project

Drug Persistence



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Problem description

ABC Pharma is looking for an automated way better than the traditional debilitating methods currently used to assess persistence of drugs as per the physician prescription, in order to have a deeper understanding on the factors impacting the persistence of their drug. The aim is to know if a patient, based on his/her information, will follow the prescription of the physician and continue taking the drug for all the treatment time. We have been provided with a dataset which contains patients' details.

Business understanding

We will create a classification model as a solution that divides patients into categories depending on their information, to determine if a patient was persistent or not.

Our goal is to create a web application that might be used as an automated solution to this process of identification.

Project Lifecycle Along with Deadline

The entire project, including all requirements, must be submitted by the **30th of August 2022**. The project has been split into several subtasks.



Figure 1 : Project Lifecycle

Data Intake Report

Name: Persistency of a Drug

Report Date: 09/08/2022

Internship Batch: LISUM10: 30

Data Intake: Sourour Cherif

Data Storage Location: https://github.com/Sururrrr/Drug_Persistence.git

Tabular Data Details: Healthcare_dataset.xlsx	
Total number of Observations	3424
Total number of File(s)	1
Total number of Features (Independent Variables or Predictors)	68
Base format of the File	.xlsx
Size of the dataset	899 KB

Data understanding

To fit any predictive model on a dataset, we need to understand the complexity of the dataset before deciding which predictive model to use to get optimal performance.

```
data.head()
```

	Persistency_Flag	Gender	Race	Ethnicity	Region	Age_Bucket	Ntm_Speciality	Ntm_Specialist_Flag	Ntm_Speciality_Bucket	Gluco_Record_Prior_Ntm	...
0	Persistent	Male	Caucasian	Not Hispanic	West	>75	GENERAL PRACTITIONER	Others	OB/GYN/Others/PCP/Unknown	N	...
1	Non-Persistent	Male	Asian	Not Hispanic	West	55-65	GENERAL PRACTITIONER	Others	OB/GYN/Others/PCP/Unknown	N	...
2	Non-Persistent	Female	Other/Unknown	Hispanic	Midwest	65-75	GENERAL PRACTITIONER	Others	OB/GYN/Others/PCP/Unknown	N	...
3	Non-Persistent	Female	Caucasian	Not Hispanic	Midwest	>75	GENERAL PRACTITIONER	Others	OB/GYN/Others/PCP/Unknown	N	...
4	Non-Persistent	Female	Caucasian	Not Hispanic	Midwest	>75	GENERAL PRACTITIONER	Others	OB/GYN/Others/PCP/Unknown	Y	...

5 rows x 68 columns

Type of data

```
[ ] data.info()
```

<class 'pandas.core.frame.DataFrame'>			
RangeIndex: 3424 entries, 0 to 3423			
Data columns (total 68 columns):			
#	Column	Non-Null Count	Dtype
0	Persistency_Flag	3424 non-null	object
1	Gender	3424 non-null	object
2	Race	3424 non-null	object
3	Ethnicity	3424 non-null	object
4	Region	3424 non-null	object
5	Age_Bucket	3424 non-null	object
6	Ntm_Speciality	3424 non-null	object
7	Ntm_Specialist_Flag	3424 non-null	object
8	Ntm_Speciality_Bucket	3424 non-null	object
9	Gluco_Record_Prior_Ntm	3424 non-null	object
10	Gluco_Record_During_Rx	3424 non-null	object
11	Dexa_Freq_During_Rx	3424 non-null	int64
12	Dexa_During_Rx	3424 non-null	object
13	Frag_Frac_Prior_Ntm	3424 non-null	object
14	Frag_Frac_During_Rx	3424 non-null	object
15	Risk_Segment_Prior_Ntm	3424 non-null	object
16	Tscore_Bucket_Prior_Ntm	3424 non-null	object
17	Risk_Segment_During_Rx	3424 non-null	object
18	Tscore_Bucket_During_Rx	3424 non-null	object
19	Change_T_Score	3424 non-null	object
20	Change_Risk_Segment	3424 non-null	object
21	Adherent_Flag	3424 non-null	object
22	Idn_Indicator	3424 non-null	object
23	Injectable_Experience_During_Rx	3424 non-null	object
24	Comorb_Encounter_For_Screening_For_Malignant_Neoplasms	3424 non-null	object
25	Comorb_Encounter_For_Immunization	3424 non-null	object
26	Comorb_Encntr_For_General_Exam_W_O_Complaint,_Susp_Or_Reprtd_Dx	3424 non-null	object
27	Comorb_Vitamin D Deficiency	3424 non-null	object

```

35 Comorb_Osteoporosis_without_current_pathological_fracture      3424 non-null object
36 Comorb_Personal_history_of_malignant_neoplasm                  3424 non-null object
37 Comorb_Gastro_esophageal_reflux_disease                        3424 non-null object
38 Concom_Cholesterol_And_Triglyceride_Regulating_Preparations    3424 non-null object
39 Concom_Narcotics                                                3424 non-null object
40 Concom_Systemic_Corticosteroids_Plain                          3424 non-null object
41 Concom_Anti_Depressants_And_Mood_Stabilisers                  3424 non-null object
42 Concom_Fluoroquinolones                                         3424 non-null object
43 Concom_Cephalosporins                                           3424 non-null object
44 Concom_Macrolides_And_Similar_Types                            3424 non-null object
45 Concom_Broad_Spectrum_Penicillins                              3424 non-null object
46 Concom_Anaesthetics_General                                     3424 non-null object
47 Concom_Viral_Vaccines                                           3424 non-null object
48 Risk_Type_1_Insulin_Dependent_Diabetes                         3424 non-null object
49 Risk_Osteogenesis_Imperfecta                                    3424 non-null object
50 Risk_Rheumatoid_Arthritis                                       3424 non-null object
51 Risk_Untreated_Chronic_Hyperthyroidism                        3424 non-null object
52 Risk_Untreated_Chronic_Hypogonadism                           3424 non-null object
53 Risk_Untreated_Early_Menopause                                 3424 non-null object
54 Risk_Patient_Parent_Fractured_Their_Hip                       3424 non-null object
55 Risk_Smoking_Tobacco                                            3424 non-null object
56 Risk_Chronic_Malnutrition_Or_Malabsorption                     3424 non-null object
57 Risk_Chronic_Liver_Disease                                      3424 non-null object
58 Risk_Family_History_Of_Osteoporosis                           3424 non-null object
59 Risk_Low_Calcium_Intake                                         3424 non-null object
60 Risk_Vitamin_D_Insufficiency                                    3424 non-null object
61 Risk_Poor_Health_Frailty                                       3424 non-null object
62 Risk_Excessive_Thinness                                         3424 non-null object
63 Risk_Hysterectomy_Oophorectomy                                 3424 non-null object
64 Risk_Estrogen_Deficiency                                        3424 non-null object
65 Risk_Immobilization                                              3424 non-null object
66 Risk_Recurring_Falls                                           3424 non-null object
67 Count_Of_Risks                                                  3424 non-null int64
dtypes: int64(2), object(66)
memory usage: 1.8+ MB

```

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

	Dexa_Freq_During_Rx	Count_Of_Risks
count	3424.000000	3424.000000
mean	3.016063	1.239486
std	8.136545	1.094914
min	0.000000	0.000000
25%	0.000000	0.000000
50%	0.000000	1.000000
75%	3.000000	2.000000
max	146.000000	7.000000

Unique elements in each Column

```
print(data.columns.unique())

<bound method Index.unique of Index(['Persistency_Flag', 'Gender', 'Race', 'Ethnicity', 'Region',
'Age_Bucket', 'Ntm_Speciality', 'Ntm_Specialist_Flag',
'Ntm_Speciality_Bucket', 'Gluco_Record_Prior_Ntm',
'Gluco_Record_During_Rx', 'Dexa_Freq_During_Rx', 'Dexa_During_Rx',
'Frag_Frac_Prior_Ntm', 'Frag_Frac_During_Rx', 'Risk_Segment_Prior_Ntm',
'Tscore_Bucket_Prior_Ntm', 'Risk_Segment_During_Rx',
'Tscore_Bucket_During_Rx', 'Change_T_Score', 'Change_Risk_Segment',
'Adherent_Flag', 'Idn_Indicator', 'Injectable_Experience_During_Rx',
'Comorb_Encounter_For_Screening_For_Malignant_Neoplasms',
'Comorb_Encounter_For_Immunization',
'Comorb_Encntr_For_General_Exam_W_O_Complaint', 'Susp_Or_Reprtd_Dx',
'Comorb_Vitamin_D_Deficiency',
'Comorb_Other_Joint_Disorder_Not_Elsewhere_Classified',
'Comorb_Encntr_For_Oth_Sp_Exam_W_O_Complaint_Suspected_Or_Reprtd_Dx',
'Comorb_Long_Term_Current_Drug_Therapy', 'Comorb_Dorsalgia',
'Comorb_Personal_History_Of_Other_Diseases_And_Conditions',
'Comorb_Other_Disorders_Of_Bone_Density_And_Structure',
'Comorb_Disorders_of_lipoprotein_metabolism_and_other_lipidemias',
'Comorb_Osteoporosis_without_current_pathological_fracture',
'Comorb_Personal_history_of_malignant_neoplasm',
'Comorb_Gastro_esophageal_reflux_disease',
'Concom_Cholesterol_And_Triglyceride_Regulating_Preparations',
'Concom_Narcotics', 'Concom_Systemic_Corticosteroids_Plain',
'Concom_Anti_Depressants_And_Mood_Stabilisers',
'Concom_Fluoroquinolones', 'Concom_Cephalosporins',
'Concom_Macrolides_And_Similar_Types',
'Concom_Broad_Spectrum_Penicillins', 'Concom_Anaesthetics_General',
'Concom_Viral_Vaccines', 'Risk_Type_1_Insulin_Dependent_Diabetes',
'Risk_Osteogenesis_Imperfecta', 'Risk_Rheumatoid_Arthritis',
'Risk_Untreated_Chronic_Hyperthyroidism',
'Risk_Untreated_Chronic_Hypogonadism', 'Risk_Untreated_Early_Menopause',
'Risk_Patient_Parent_Fractured_Their_Hip', 'Risk_Smoking_Tobacco',
'Risk_Chronic_Malnutrition_Or_Malabsorption',
'Risk_Chronic_Liver_Disease', 'Risk_Family_History_Of_Osteoporosis',
'Risk_Low_Calcium_Intake', 'Risk_Vitamin_D_Insufficiency',
'Risk_Poor_Health_Frailty', 'Risk_Excessive_Thinness',
'Risk_Hysterectomy_Oophorectomy', 'Risk_Estrogen_Deficiency',
'Risk_Immobilization', 'Risk_Recurring_Falls', 'Count_Of_Risks'],
dtype='object')>
```

Data Problems

Data problems such as irrelevant columns, Null values, duplicates, skewed data, outliers and many others may cause bad predictions ...

So we need to check if we have one of them to know then how to overcome it .

- Skewed Data :

```
✓ [497] def measure_skew_kurtosis(cols):  
0 s     for col in cols:  
         print(col)  
         result = data[[col]].agg(['skew', 'kurtosis']).transpose()  
         print(result)  
     measure_skew_kurtosis(numeric_col)
```

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

```
✓ [498] #skew and kurtosis values  
0 s     data.agg(['skew', 'kurtosis']).transpose()
```

	skew	kurtosis
Dexa_Freq_During_Rx	6.808730	74.758378
Count_Of_Risks	0.879791	0.900486

_ Outliers

```
▶ # creating a box plot of numerical columns against persistency flag to identify outliers  
def boxplot(data, cols):  
    for col in cols:  
        sns.set_style('whitegrid')  
        sns.boxplot(x='Persistency_Flag', y=col, data=data)  
        plt.title('Boxplot of ' + col)  
        plt.ylabel(col) #setting text for y axis  
        plt.show()  
    boxplot(data, numeric_col)
```

_ Duplicates

⇒ There is no duplicates, Having duplicates leads often to overfitting

_ Missing Values

```
[ ] # Total number of missing values
data.isnull().sum().sum()
```

0

⇒ No missing Values

Solutions

- Removing duplicates if they exists
- Dropping insignificant columns
- Eliminating Skewed data

```
✓ [501] #Checking skew after transformation
0s data.agg(['skew', 'kurtosis']).transpose()
```

	skew	kurtosis
Dexa_Freq_During_Rx	6.808730	74.758378
Count_Of_Risks	0.879791	0.900486
log_Dexa	1.405860	0.624570
log_Count_Risks	-0.091583	-1.006414

Example of removing 99% Percentile

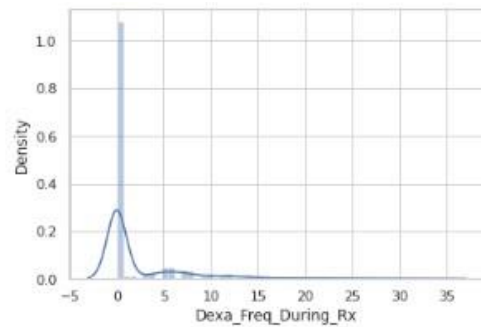
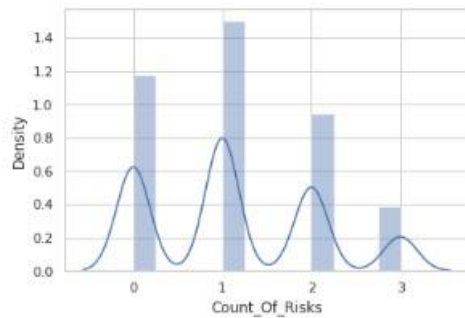

```

[505] # To remove the 99th percentile
q = data['Dexa_Freq_During_Rx'].quantile(0.99)
data_1 = data[data['Dexa_Freq_During_Rx'] < q]
data_1.describe()

```

	Dexa_Freq_During_Rx	Count_Of_Risks	log_Dexa	log_Count_Risks
count	3389.000000	3389.000000	3389.000000	3389.000000
mean	2.440248	1.240484	0.572915	0.685941
std	5.183446	1.095904	0.997375	0.499826
min	0.000000	0.000000	0.000000	0.000000
25%	0.000000	0.000000	0.000000	0.000000
50%	0.000000	1.000000	0.000000	0.693147
75%	3.000000	2.000000	1.386294	1.098612
max	34.000000	7.000000	3.555348	2.079442

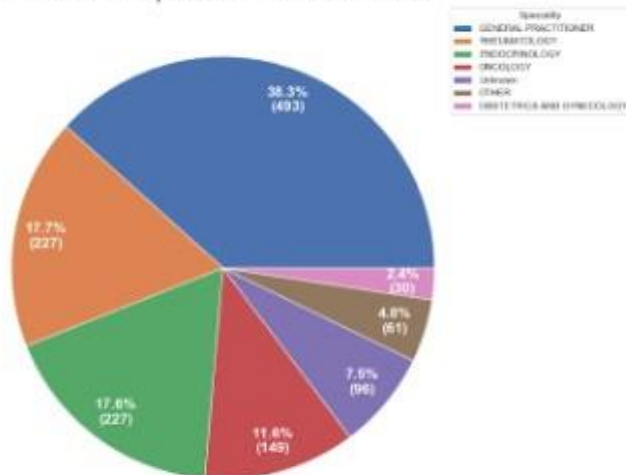
- Removing outliers



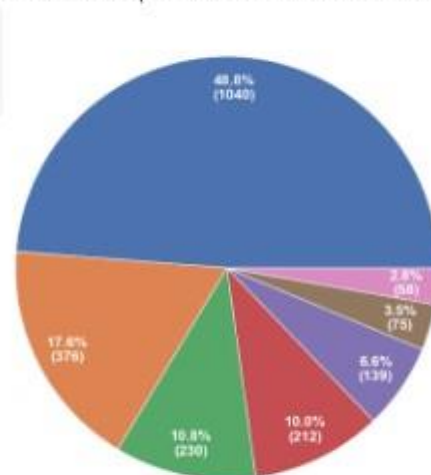
EDA

Does the speciality of the person who prescribed the drug have any effect on the persistent rate?

Distribution of Specialities for Persistent Cases



Distribution of Specialities for Non-Persistent Cases



We see that both pie charts are pretty similar in distribution of frequency for each speciality. Thus, we can rule out the possibility that one of the factors that the drug is persistent or not is the speciality that prescribed the drug in the first place.

Does 'Ntm_Specialist_Flag' and 'Ntm_Speciality_Bucket' Variables have useful information for the classification task?

Persistency_Flag	0	1
Ntm_Specialist_Flag		
Others	0.686214	0.313786
Specialist	0.552553	0.447447

Persistency_Flag	0	1
Ntm_Speciality_Bucket		
Endo/Onc/Uro	0.473134	0.526866
OB/GYN/Others/PCP/Unknown	0.684884	0.315116
Rheum	0.629565	0.370435

It seems Rheum flag in Ntm_Speciality_Bucket have some useful information.

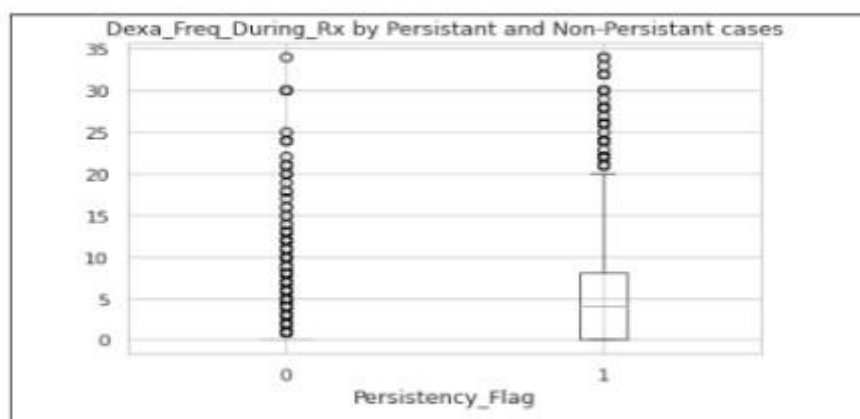
What about 'Gluko_Record_Prior_Ntm', 'Gluko_Record_During_Rx'?

Persistency_Flag	0	1
Gluko_Record_Prior_Ntm		
N	0.627879	0.372121
Y	0.645119	0.354881

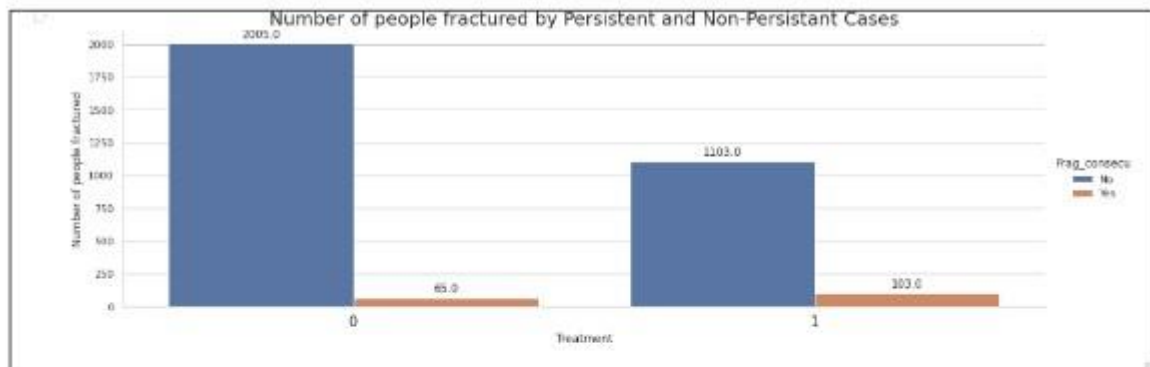
Persistency_Flag	0	1
Gluko_Record_During_Rx		
N	0.691044	0.308956
Y	0.460808	0.539192

Gluko_Record_During_Rx seems to be more useful than Gluko_Record_Prior_Ntm to predict the target

The distribution of Dexa_Freq_During_Rx numbers seems to be higher in the Persistent patients

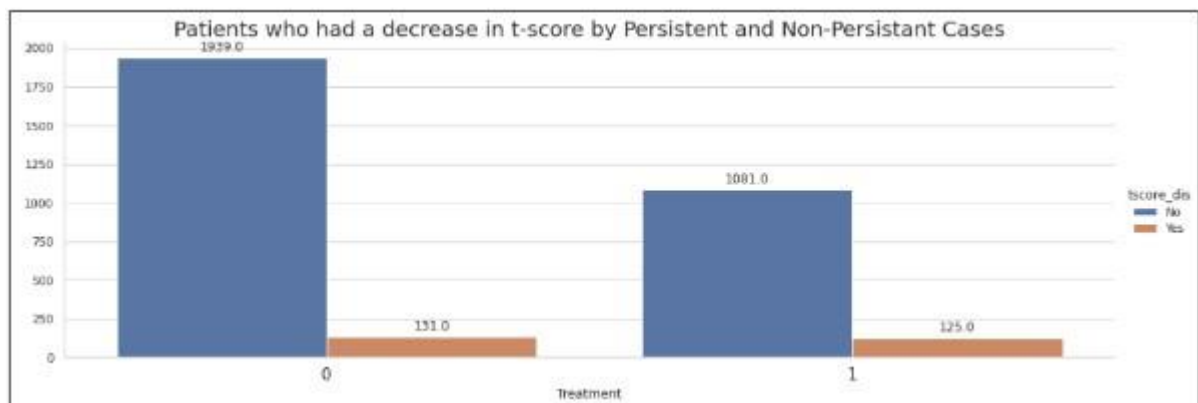


Variables that are recorded during the treatment have more useful information for the classification than others. It can be checked with the percentages shown by Dexa_During_Rx variable.



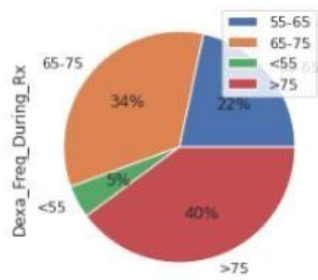
Of the total number of patients, 8% of people were affected by the treatment, weakening their bones

- The count of people affected by the treatment is small, and we can speculate that the treatment not affected considerably to the bones of the patients.

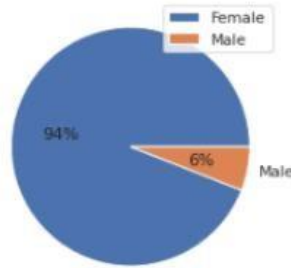


There is 10% of people with treatment who had a decrease in the t-score

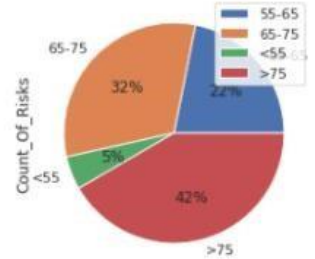
- Then there is 90% approximately of people who maintained or improved their t-score.
- In conclusion, the treatment is improving the t-score of the patients.



Dexa_Freq_During_Rx
by Age_Bucket



Count_Of_Risks
by Gender



Counts_Of_Risks
by Age-Bucket

Most of the patients already hold comorbidity factors, while holding risk factors is less common.

Some highlights:

- The main comorbidity factor is related to lipoproteins and metabolism (cholesterol).
- The main risk factor is deficiency in vitamin D.
- More than one third has been found to have taken narcotics.
- 99 % of our sample hold at least one risk, comorbidity and/or concomitant factor.

There are some significant differences between genders:

- Women seem to be more affected by **vitamin D deficiencies**.
- More than twice as many women as men have passed as screening for **malignant neoplasms**.
- Four times as many men as women suffer from **Hypogonadism** (untreated).

- As expected, patients **older than 65** are affected by the mentioned factors in a higher proportion.
- There are some risks and other factors that seem to be significantly higher in **South and West regions**. It might be interesting to find out about socioeconomic factors aside.
- There seem to be some remarkable differences between **Asian and other** races. They are probably due to cultural factors and other behaviours, like medical reviews on a more regular basis (this is just a hypothesis to be found out).

⇒ EDA Summary

The file contained information of 3,424 patients. For each patient it has demographic information, clinical records, others diseases as risk factor information and also about their physician's speciality.

There are some significant differences between genders (vitamin D deficiencies, screening for malignant neoplasms, Hypogonadism).

Most of the patients already hold comorbidity factors, while holding risk factors is less common.

Patients older than 65 are affected by the mentioned factors in a higher proportion.

There seem to be some remarkable differences between Asian and other races.

Variables that are recorded during the treatment like DEXA_Freq_During_Rx, DEXA_During_Rx and Gluco_Record_During_Rx have more useful information for the classification than others.

Modeling Techniques

Considering the nature of target variable the classification modeling techniques are most suitable for present study. This is a problem of binary classification and models logistic regression, decision tree can be used easily.

