

Week 8 – Healthcare Project

Group Name: Cool Data Scientists Team

Team Members Details:

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Problem Description

One of the challenges for Pharmaceutical companies is to understand the persistency of drug as per the physician prescription. This issue results in a bad impact on the pharmacies for all the categories; patients, physicians, and administration. However, the team of data scientist is capable of discovering the analyzing the dataset and detecting the factors that are impacting the primary factor which is the "persistency". By building a classification machine learning model, we will be able to classify the dataset and find the variables that affect the target variables "Persistency Flag".

Data understanding

As a first step, we imported the dataset and copied it. Then we've looked at the first five and the last five entries.

The following pictures show how our dataset looks like:

```
df.head()
```

	Ptid	Persistence_Flag	Gender	Race	Ethnicity	Region	Age_Bucket	Ntm_Speciality	Ntm_Specialist_Flag	Ntm_Speciality_Bucket	...
0	P1	Persistent	Male	Caucasian	Not Hispanic	West	>75	GENERAL PRACTITIONER	Others	OB/GYN/Others/PCP/Unknown	...
1	P2	Non-Persistent	Male	Asian	Not Hispanic	West	55-65	GENERAL PRACTITIONER	Others	OB/GYN/Others/PCP/Unknown	...
2	P3	Non-Persistent	Female	Other/Unknown	Hispanic	Midwest	65-75	GENERAL PRACTITIONER	Others	OB/GYN/Others/PCP/Unknown	...
3	P4	Non-Persistent	Female	Caucasian	Not Hispanic	Midwest	>75	GENERAL PRACTITIONER	Others	OB/GYN/Others/PCP/Unknown	...
4	P5	Non-Persistent	Female	Caucasian	Not Hispanic	Midwest	>75	GENERAL PRACTITIONER	Others	OB/GYN/Others/PCP/Unknown	...

```
df.tail()
```

	Ptid	Persistence_Flag	Gender	Race	Ethnicity	Region	Age_Bucket	Ntm_Speciality	Ntm_Specialist_Flag	Ntm_Speciality_Bucket	...
3419	P3420	Persistent	Female	Caucasian	Not Hispanic	South	>75	GENERAL PRACTITIONER	Others	OB/GYN/Others/PCP/Unknown	...
3420	P3421	Persistent	Female	Caucasian	Not Hispanic	South	>75	Unknown	Others	OB/GYN/Others/PCP/Unknown	...
3421	P3422	Persistent	Female	Caucasian	Not Hispanic	South	>75	ENDOCRINOLOGY	Specialist	Endo/Onc/Uro	...
3422	P3423	Non-Persistent	Female	Caucasian	Not Hispanic	South	55-65	Unknown	Others	OB/GYN/Others/PCP/Unknown	...
3423	P3424	Non-Persistent	Female	Caucasian	Not Hispanic	South	65-75	Unknown	Others	OB/GYN/Others/PCP/Unknown	...

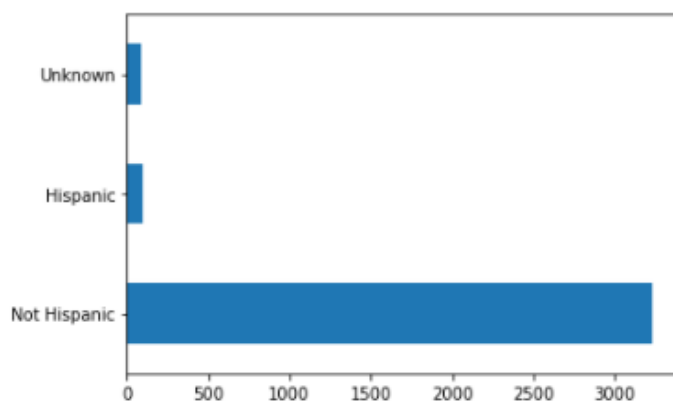
Totally we have 3424 observations and 69 features.

```
df.shape
```

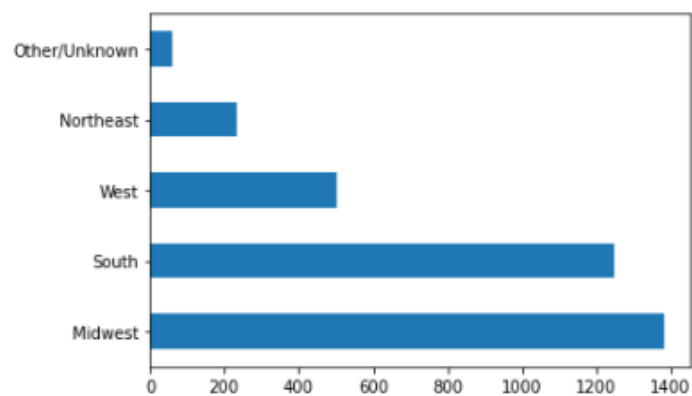
(3424, 69)

For Demographics, we have the followings:

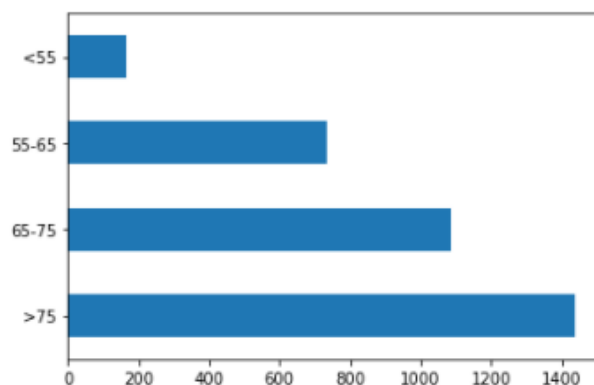
If we examine "Ethnicity", we see that "Non-Hispanic" people dominates the "Hispanic" people and also we have unknown values.



If we examine the “Region”, we see that patients are mostly “Midwest” and “South” region:

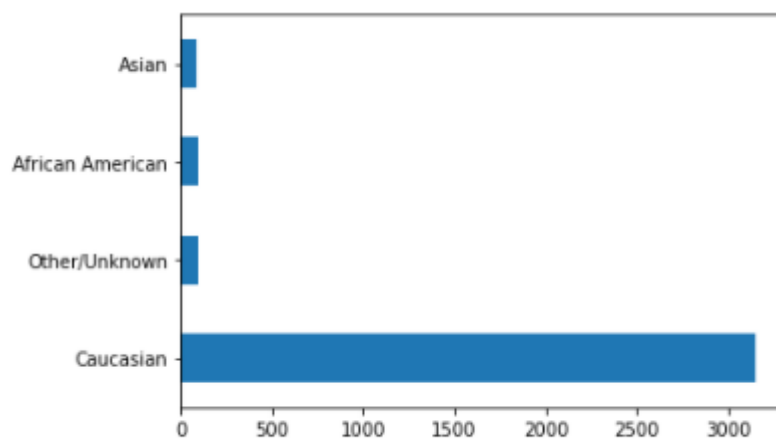


If we look at the “Age”, we see the following:

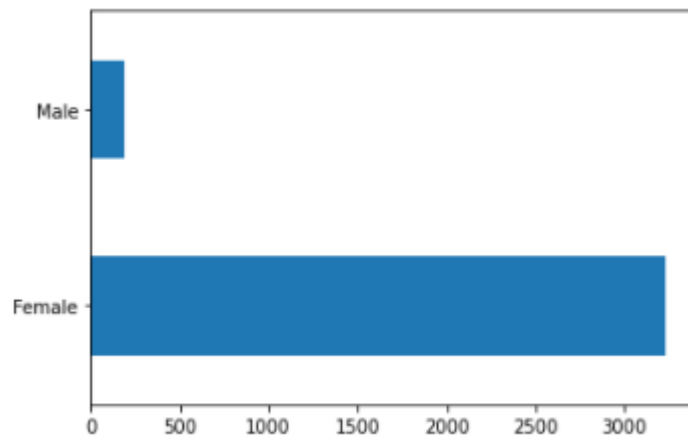


By looking at the above picture, it can be thought that being of age “>55” can be related to have persistency to drug.

If we look at the “Race”, we see that the Caucasians are dominated the other races.



If we look at the “Gender”, by the following picture , the female patients are more than the male patients.



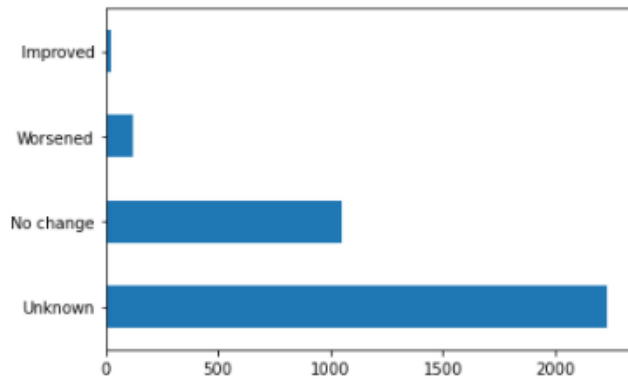
Ntm Speciality is the specialty of the HCP that prescribed the NTM Rx.

We see that General Practitioner, Rheumatology, Endocrinology and Oncology specialists prescribed the NTM Rx most.

GENERAL PRACTITIONER	1535
RHEUMATOLOGY	604
ENDOCRINOLOGY	458
Unknown	310
ONCOLOGY	225
OBSTETRICS AND GYNECOLOGY	90
UROLOGY	33
ORTHOPEDIC SURGERY	30
CARDIOLOGY	22
PATHOLOGY	16
HEMATOLOGY & ONCOLOGY	14
OTOLARYNGOLOGY	14
PEDIATRICS	13
PHYSICAL MEDICINE AND REHABILITATION	11
PULMONARY MEDICINE	8
SURGERY AND SURGICAL SPECIALTIES	8
PSYCHIATRY AND NEUROLOGY	4
NEPHROLOGY	3
ORTHOPEDICS	3
GERIATRIC MEDICINE	2
HOSPICE AND PALLIATIVE MEDICINE	2
PLASTIC SURGERY	2
GASTROENTEROLOGY	2
VASCULAR SURGERY	2
TRANSPLANT SURGERY	2
OCCUPATIONAL MEDICINE	1
OPHTHALMOLOGY	1
PAIN MEDICINE	1

Clinical Factors:

Risk Segment: We have compared the risk segments prior NTM and during NTM and examine how it changes:



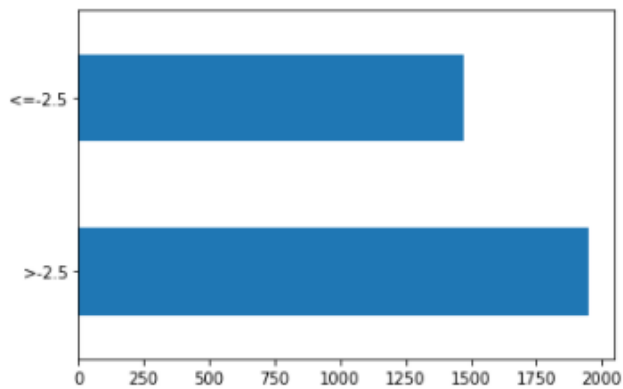
We have done similar computations for all other clinical factors.

For instance, we have examined the Fragility and we have obtained the following cross- table:

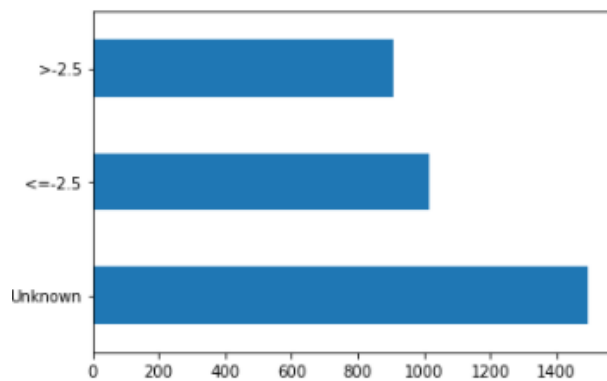
Frag_Frac_During_Rx		N	Y
Frag_Frac_Prior_Ntm			
	N	2691	181
	Y	316	236

T-scores:

We have compared the “T-scores”. The following picture shows the prior to NTM:



The following shows the “T-scores” during the Rx:



Besides, we have examined the Disease and Treatment Factors. They are all Yes/No information and we have decided which of the variables can affect the persistency to drug:

By comparing the results, we see that the followings can affect the target variable:

- 1) Comorb_Encounter_For_Screening_For_Malignant_Neoplasms
- 2) Comorb_Disorders_of_lipoprotein_metabolism_and_other_lipidemias
- 3) Comorb_Encounter_For_Immunization.

Similarly, we think that Vitamin D-insufficiency can affect the target variable.

What type of data you have got for analysis?

When we've checked the types of the variables, we obtained the following result:

```
df.info()

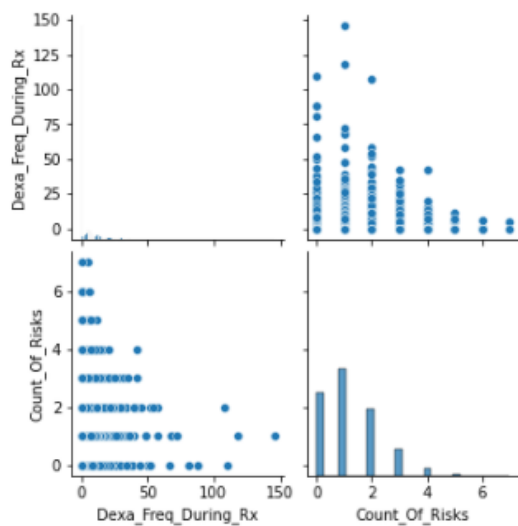
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 3424 entries, 0 to 3423
Data columns (total 69 columns):
#   Column                                     Non-Null Count  Dtype
---  -
0   Ptid                                       3424 non-null   object
1   Persistency_Flag                         3424 non-null   object
2   Gender                                   3424 non-null   object
3   Race                                     3424 non-null   object
4   Ethnicity                               3424 non-null   object
5   Region                                   3424 non-null   object
6   Age_Bucket                              3424 non-null   object
7   Ntm_Speciality                          3424 non-null   object
8   Ntm_Specialist_Flag                     3424 non-null   object
9   Ntm_Speciality_Bucket                   3424 non-null   object
10  Gluco_Record_Prior_Ntm                  3424 non-null   object
11  Gluco_Record_During_Rx                  3424 non-null   object
12  Dexa_Freq_During_Rx                     3424 non-null   int64
13  Dexa_During_Rx                          3424 non-null   object
14  Frag_Frac_Prior_Ntm                     3424 non-null   object
15  Frag_Frac_During_Rx                     3424 non-null   object
16  Risk_Segment_Prior_Ntm                  3424 non-null   object
17  Tscore_Bucket_Prior_Ntm                  3424 non-null   object
18  Risk_Segment_During_Rx                  3424 non-null   object
19  Tscore_Bucket_During_Rx                  3424 non-null   object
20  Change_T_Score                          3424 non-null   object
21  Change_Risk_Segment                     3424 non-null   object
22  Adherent_Flag                           3424 non-null   object
```

23	Idn_Indicator	3424	non-null	object
24	Injectable_Experience_During_Rx	3424	non-null	object
25	Comorb_Encounter_For_Screening_For_Malignant_Neoplasms	3424	non-null	object
26	Comorb_Encounter_For_Immunization	3424	non-null	object
27	Comorb_Encntr_For_General_Exam_W_O_Complaint,_Susp_Or_Reprtd_Dx	3424	non-null	object
28	Comorb_Vitamin_D_Deficiency	3424	non-null	object
29	Comorb_Other_Joint_Disorder_Not_Elsewhere_Classified	3424	non-null	object
30	Comorb_Encntr_For_Oth_Sp_Exam_W_O_Complaint_Suspected_Or_Reprtd_Dx	3424	non-null	object
31	Comorb_Long_Term_Current_Drug_Therapy	3424	non-null	object
32	Comorb_Dorsalgia	3424	non-null	object
33	Comorb_Personal_History_Of_Other_Diseases_And_Conditions	3424	non-null	object
34	Comorb_Other_Disorders_Of_Bone_Density_And_Structure	3424	non-null	object
35	Comorb_Disorders_of_lipoprotein_metabolism_and_other_lipidemias	3424	non-null	object
36	Comorb_Osteoporosis_without_current_pathological_fracture	3424	non-null	object
37	Comorb_Personal_history_of_malignant_neoplasm	3424	non-null	object
38	Comorb_Gastro_esophageal_reflux_disease	3424	non-null	object
39	Concom_Cholesterol_And_Triglyceride_Regulating_Preparations	3424	non-null	object
40	Concom_Narcotics	3424	non-null	object
41	Concom_Systemic_Corticosteroids_Plain	3424	non-null	object
42	Concom_Anti_Depressants_And_Mood_Stabilisers	3424	non-null	object
43	Concom_Fluoroquinolones	3424	non-null	object
44	Concom_Cephalosporins	3424	non-null	object
45	Concom_Macrolides_And_Similar_Types	3424	non-null	object
46	Concom_Broad_Spectrum_Penicillins	3424	non-null	object
47	Concom_Anaesthetics_General	3424	non-null	object
48	Concom_Viral_Vaccines	3424	non-null	object
49	Risk_Type_1_Insulin_Dependent_Diabetes	3424	non-null	object
50	Risk_Osteogenesis_Imperfecta	3424	non-null	object
51	Risk_Rheumatoid_Arthritis	3424	non-null	object
52	Risk_Untreated_Chronic_Hyperthyroidism	3424	non-null	object
53	Risk_Untreated_Chronic_Hypogonadism	3424	non-null	object
54	Risk_Untreated_Early_Menopause	3424	non-null	object
55	Risk_Patient_Parent_Fractured_Their_Hip	3424	non-null	object
56	Risk_Smoking_Tobacco	3424	non-null	object
57	Risk_Chronic_Malnutrition_Or_Malabsorption	3424	non-null	object
58	Risk_Chronic_Liver_Disease	3424	non-null	object
59	Risk_Family_History_Of_Osteoporosis	3424	non-null	object
60	Risk_Low_Calcium_Intake	3424	non-null	object
61	Risk_Vitamin_D_Insufficiency	3424	non-null	object
62	Risk_Poor_Health_Frailty	3424	non-null	object
63	Risk_Excessive_Thinness	3424	non-null	object
64	Risk_Hysterectomy_Oophorectomy	3424	non-null	object
65	Risk_Estrogen_Deficiency	3424	non-null	object
66	Risk_Immobilization	3424	non-null	object
67	Risk_Recurring_Falls	3424	non-null	object
68	Count_Of_Risks	3424	non-null	int64

dtypes: int64(2), object(67)
memory usage: 1.8+ MB

We have that those 67 features are of object type and just 2 of them are int64 type.

And we have determined the relation between these two numerical variables:



What are the problems in the data (number of NA values, outliers , skewed etc):

NA Values:

When we checked that whether there is any NA value, we have obtained the following:

```
In [8]: df.isnull().values.any()
Out[8]: False

In [9]: df.isnull().sum()
Out[9]: Ptid 0
Persistency_Flag 0
Gender 0
Race 0
Ethnicity 0
..
Risk_Hysterectomy_Oophorectomy 0
Risk_Estrogen_Deficiency 0
Risk_Immobilization 0
Risk_Recurring_Falls 0
Count_Of_Risks 0
Length: 69, dtype: int64
```

Even if we don't have any NA values, we have "Unknown" variables. The followings are only examples of some of them:


```
In [9]: df["Ethnicity"].value_counts()
```

```
Out[9]: Not Hispanic    3235  
        Hispanic        98  
        Unknown         91  
        Name: Ethnicity, dtype: int64
```

```
In [11]: df["Region"].value_counts()
```

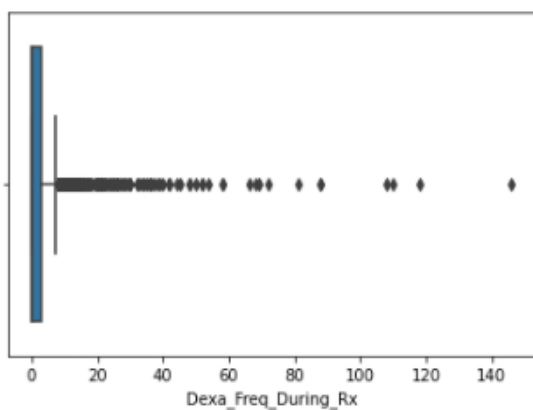
```
Out[11]: Midwest        1383  
        South          1247  
        West            502  
        Northeast       232  
        Other/Unknown    60  
        Name: Region, dtype: int64
```

```
In [20]: df["Risk_Segment_During_Rx"].value_counts()
```

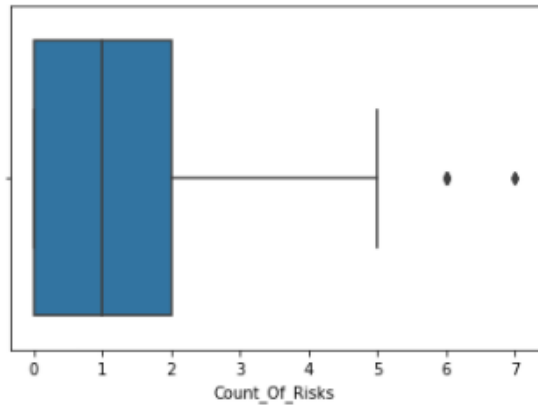
```
Out[20]: Unknown        1497  
        HR_VHR          965  
        VLR_LR           962  
        Name: Risk_Segment_During_Rx, dtype: int64
```

Outliers:

To detect the outliers, we've used boxplot.



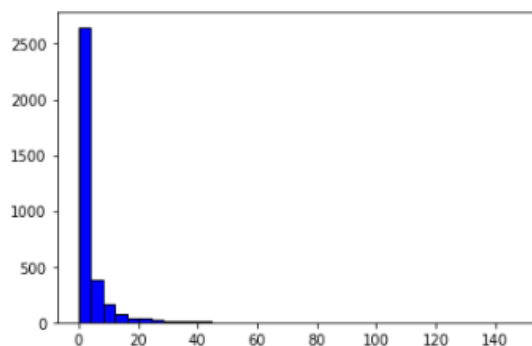
We have 460 outliers in "Dexa_Freq_During_Rx" variable.



We have 8 outliers in “Count_Of_Risks” variable.

Skewed Data:

We have the following histogram graphs:



As seen in the above, since the tail is on the right side, we can say that “Dexa_Freq_During_Rx” variable has right-skewed distribution. Hence, we can conclude that the mean value is greater than the mode.

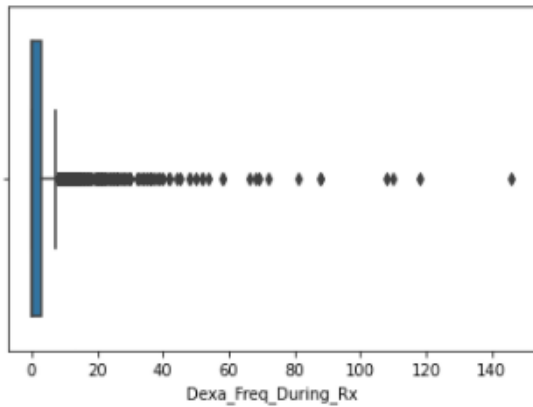
What approaches you are trying to apply on your data set to overcome problems like NA value, outlier etc and why?

For NA values: Since all of the NA values are in object types we prefer to ignore these values.

For instance, we have NA values in “Ethnicity”. If we change the Unknown values with “Hispanic” or “Non-Hispanic” it can change the result of the dataset.

For Outliers:

As seen in the following picture, the outliers of the “**Dexa_Freq_During_Rx**” variable are placed on the right-hand side of the upper bound. So, if we replace them with the mean value can change the type of the dataset. But instead, we have discussed on suppressing them with the upper bound.



On the other hand, the number of the outliers of the “**Count_Of_Risks**” variable is just 8. So, we can use mean value or suppress them with the upper bound.

Github Repo link

<https://github.com/melis-ta/Healthcare>