

Group Name: **Attack on Data**

Team Members:

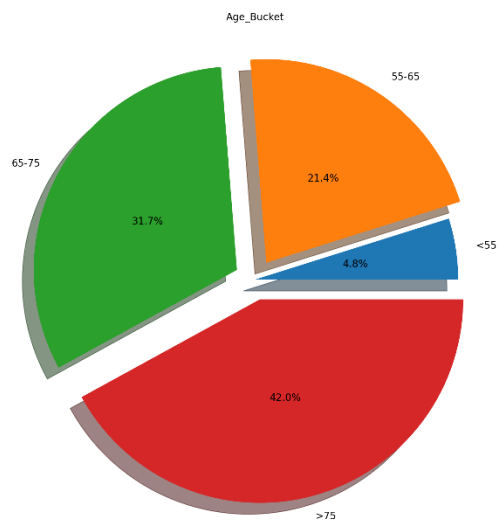
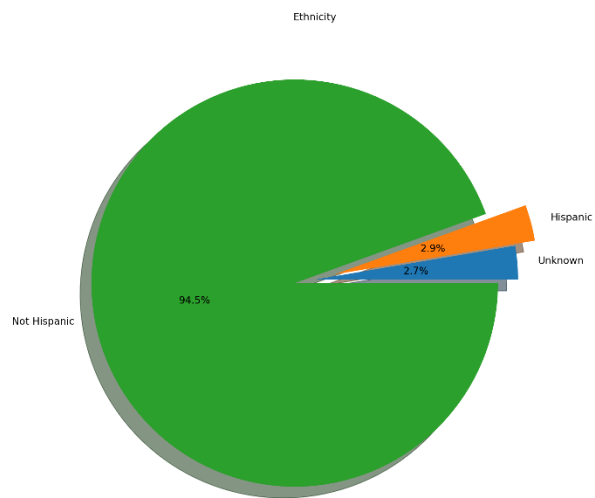
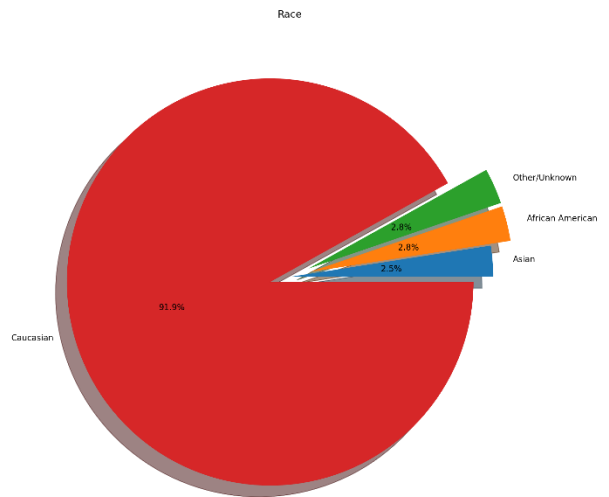
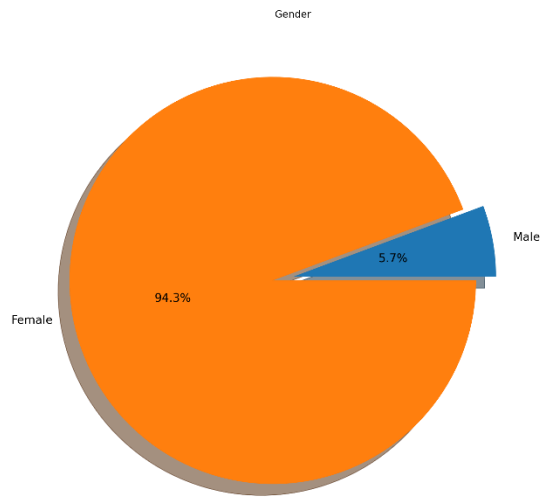
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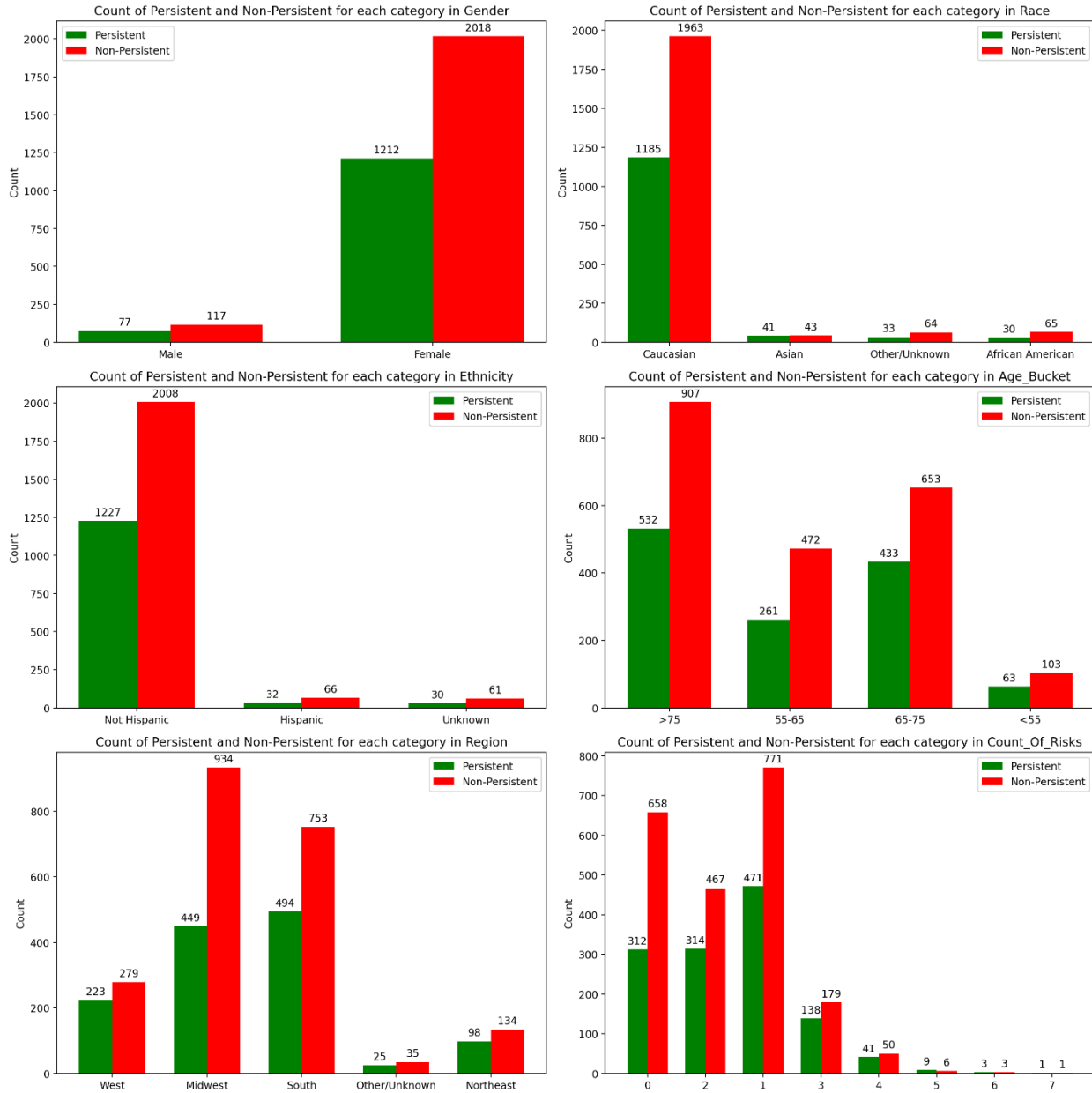
Problem description

One of the challenges for all pharmaceutical companies is to understand the persistency of drug as per the physician prescription. To solve this problem ABC pharma company approached an analytics company to automate this process of identification.

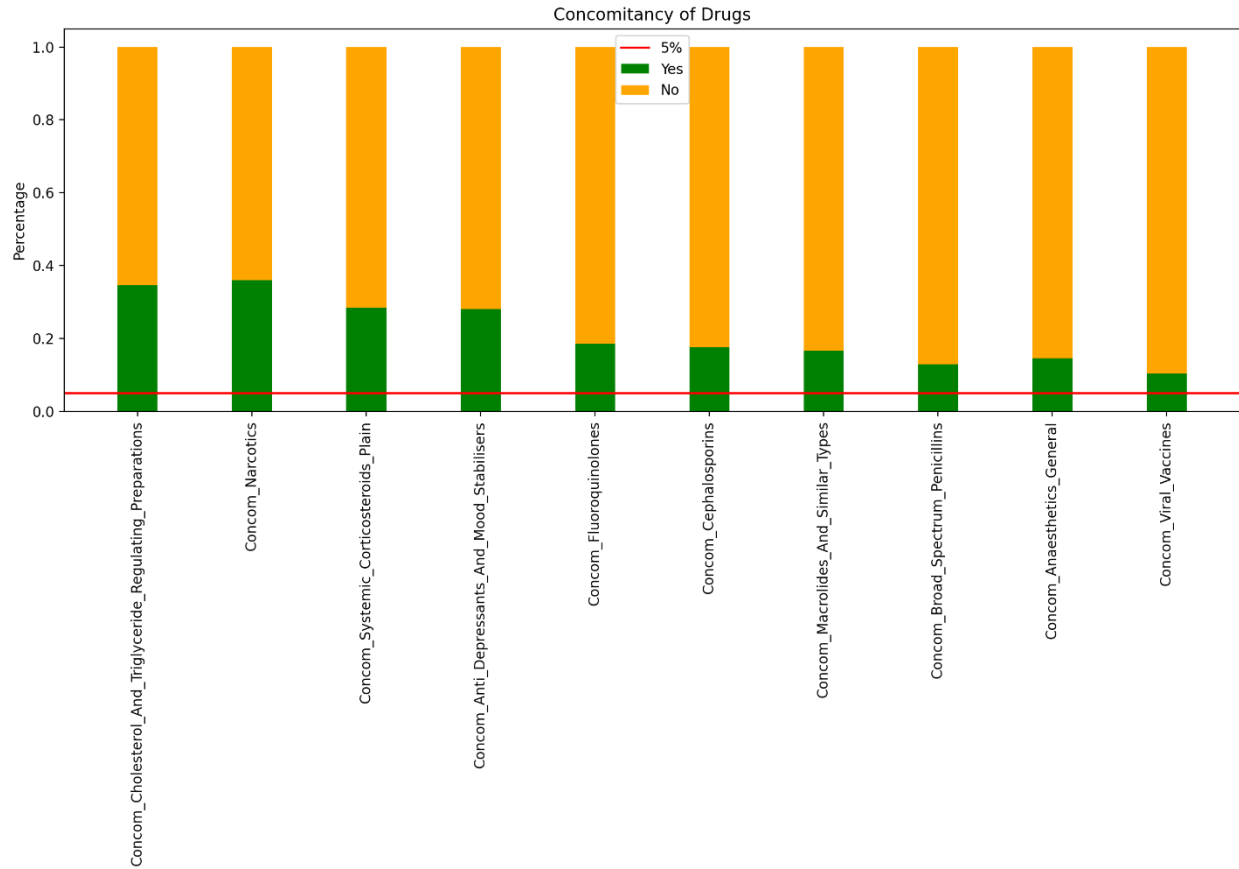
Exploratory data analysis (EDA)

Our dataset has 69 columns, 67 of them are categorical and two of them are numerical. There are no nan values in the dataset. The main problem of it is that some columns have rare labels. As we can see in the bellow charts rare categories is the main issue.

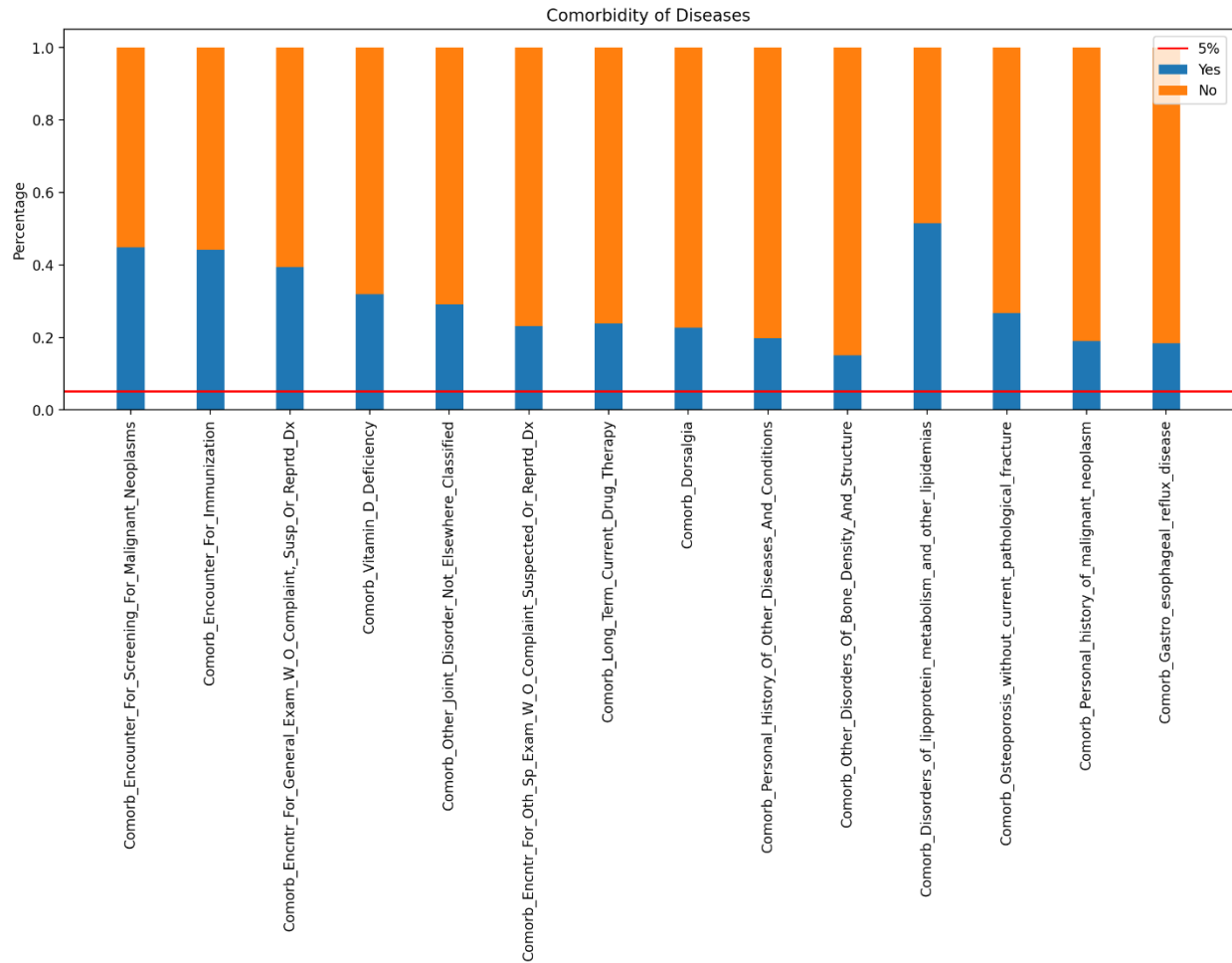




By a look at Concomitancy of Drugs bellow we can see that Cholesterol drug is more used than others beside the main drug. For other drugs, Concomitancy percentage is very low.



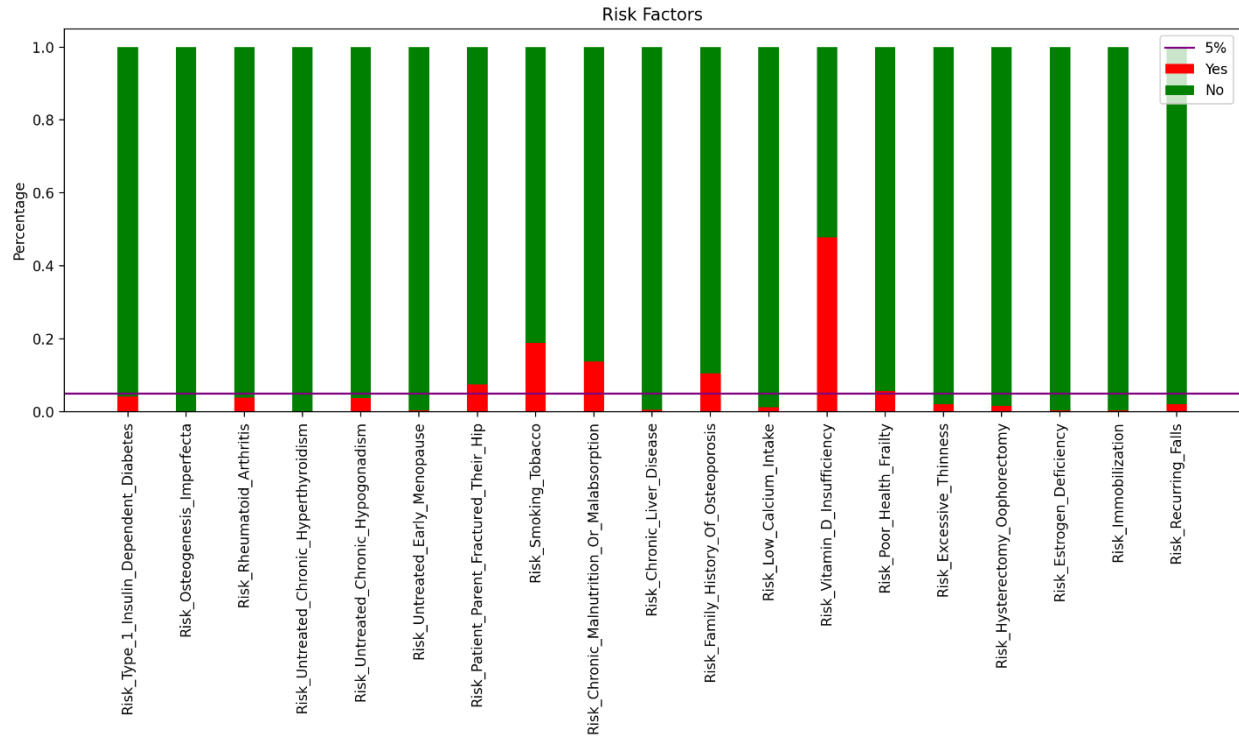
Also for Comorbidity of Diseases, we can see that disorders of lipoprotein have higher percentage to comorbid with main disease than other diseases and disorders of bone density is the lowest among all.



And among the risk factors most of them have less than 5% chance to endanger treatment. The risk factor with highest chance is Vitamin D Insufficiency and others above 5% are:

- Poor Health Frailty
- Family History of Osteoporosis
- Chronic Malnutrition or Malabsorption
- Smoking Tobacco
- Patient Parent Fractured Their Hip

Rest of the factors have less than 5% risk to endanger treatment.



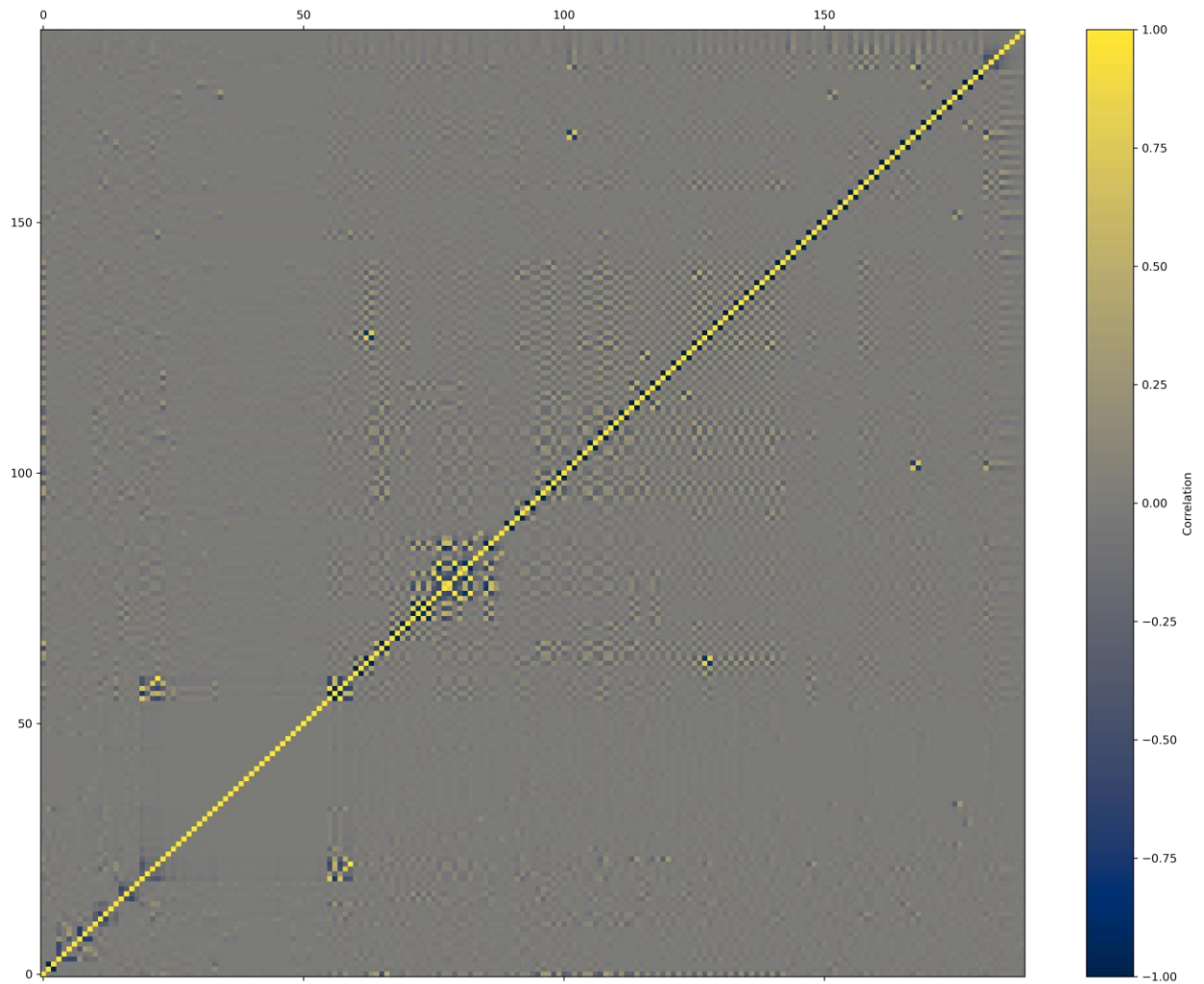
Recommendation

1. Handling **Unknown** values for Race, Region, and Ethnicity Variables
 - Using mode as an imputer as an imputer on **Race** and **Ethnicity** variables.
 - For **Region** variable, because most of the people with **Unknown** Region have **Not Hispanic** Ethnicity, and Most of people with Not Hispanic Ethnicity, have Midwest Region, we will replace Unknown Regions with **Midwest**.
2. Handling **Rare Labels**: Finding categories less than 5 percent in each variable, then merging those categories into one or drop them if the variable only has 2 categories (e.g., Y/N) and cardinality of one them is less than 5 percent.
3. Grouping integer values of Count_Of_Risks variable into two **bins**.

Group	Variable	Categories to Merge
Variables Chosen to Merge Categories	Race	African American, Asian
	Age_Bucket	<55, 55-65
	Ntm_Speciality	OBSTETRICS AND GYNECOLOGY , UROLOGY , ORTHOPEDIC SURGERY , CARDIOLOGY , PATHOLOGY , HEMATOLOGY & ONCOLOGY , OTOLARYNGOLOGY , PEDIATRICS , PHYSICAL MEDICINE AND REHABILITATION , PULMONARY MEDICINE , SURGERY AND SURGICAL SPECIALTIES , PSYCHIATRY AND NEUROLOGY , NEPHROLOGY , ORTHOPEDICS , PLASTIC SURGERY , VASCULAR SURGERY , HOSPICE AND PALLIATIVE MEDICINE , GERIATRIC MEDICINE , GASTROENTEROLOGY , TRANSPLANT SURGERY , CLINICAL NURSE SPECIALIST ,

		OCCUPATIONAL MEDICINE , HOSPITAL MEDICINE , OPHTHALMOLOGY , PODIATRY , EMERGENCY MEDICINE , RADIOLOGY , OBSTETRICS & OBSTETRICS & GYNECOLOGY & OBSTETRICS & GYNECOLOGY , NEUROLOGY , PAIN MEDICINE , NUCLEAR MEDICINE
	Change_T_Score	Improved, Worsened
	Change_Risk_Segment	Improved, Worsened
	Count_Of_Risks	Bin 1 is [0,1,2,3] and bin 2 is [4,5,6,7]
Variables Chosen to Drop	Ethnicity	
	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_Chronic_Liver_Disease	
	Risk_Low_Calcium_Intake	
	Risk_Excessive_Thinness	
	Risk_Hysterectomy_Oophorectomy	
	Risk_Estrogen_Deficiency	
	Risk_Immobilization	
	Risk_Recurring_Falls	
	Dexa_Freq_During_Rx	

4. **One hot encoding** all the variables after doing above tasks
5. Removing variables with more than **98% correlation**.



6. Try different machine learning approaches and select the best model.

ML model evaluation and selection

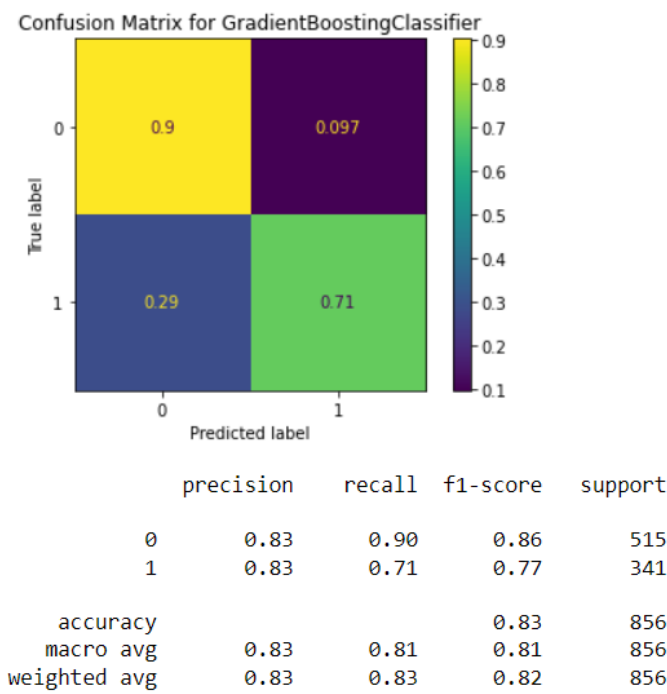
The following models were trained and tested on the processed dataset:

- 1. GradientBoostingClassifier
- 2. ExtraTreesClassifier
- 3. RandomForestClassifier
- 4. LogisticRegression
- 5. SVR (RBF kernel)
- 6. SVR (Linear kernel)
- 7. SVR (Polynomial kernel)
- 8. SVR (Sigmoid kernel)
- 9. XGBClassifier

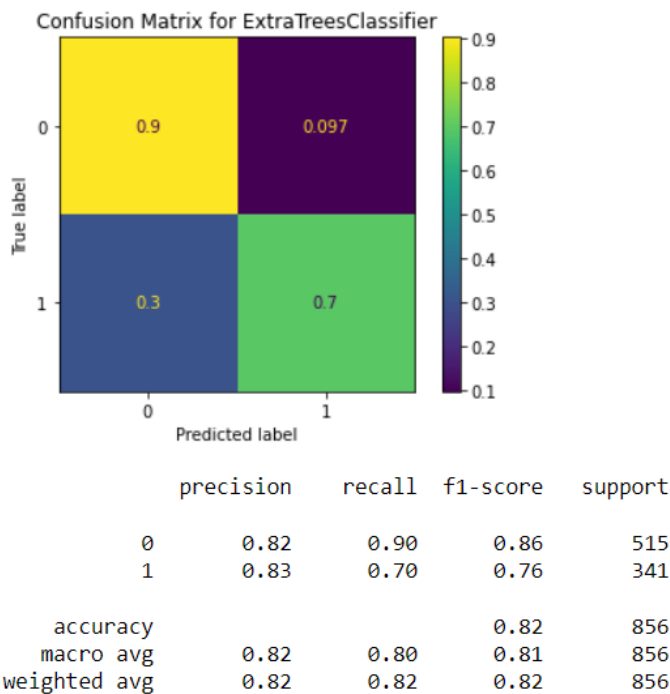
Results:

For each model, the accuracy, precision, recall, f1-score and support were noted. The figures below illustrate

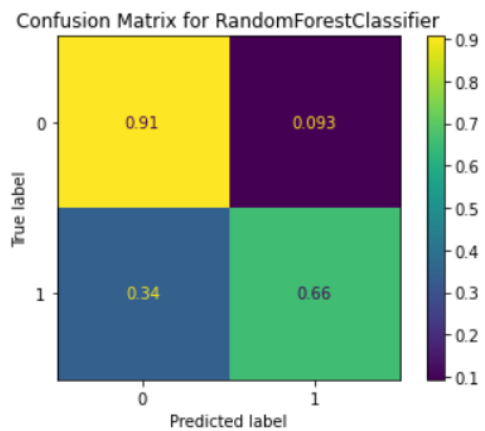
Accuracy GradientBoostingClassifier: 0.83



Accuracy ExtraTreesClassifier: 0.82

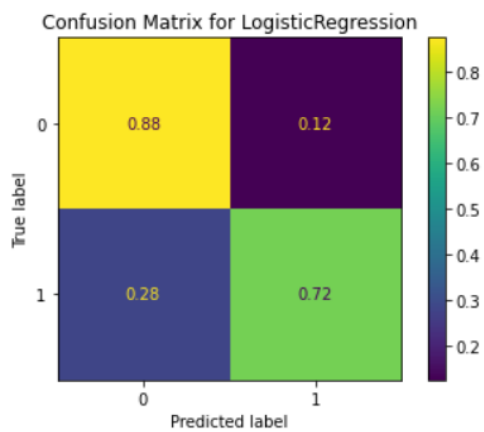


Accuracy RandomForestClassifier: 0.81



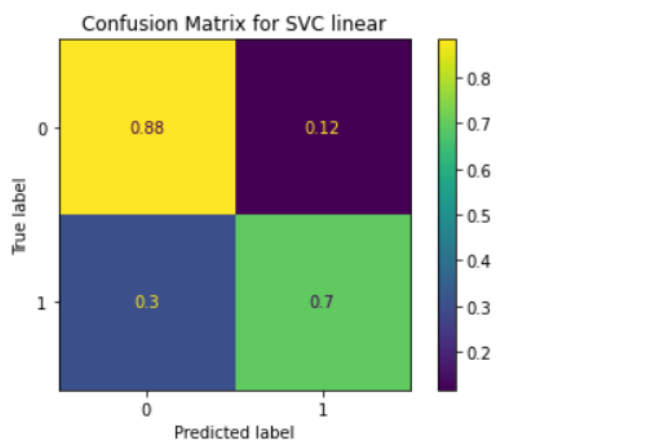
	precision	recall	f1-score	support
0	0.80	0.91	0.85	515
1	0.82	0.66	0.73	341
accuracy			0.81	856
macro avg	0.81	0.78	0.79	856
weighted avg	0.81	0.81	0.80	856

Accuracy LogisticRegression: 0.81



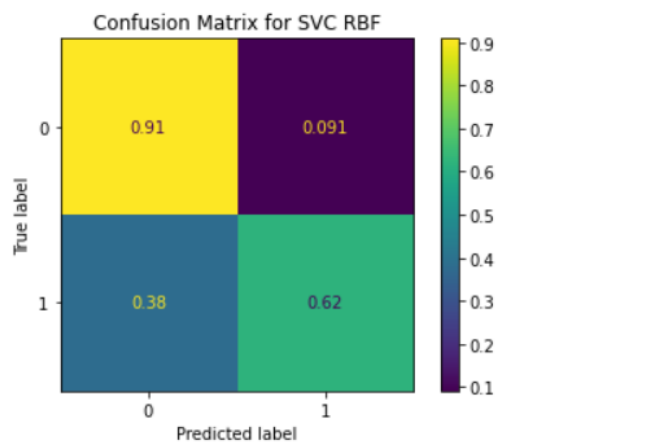
	precision	recall	f1-score	support
0	0.82	0.88	0.85	515
1	0.79	0.72	0.75	341
accuracy			0.81	856
macro avg	0.81	0.80	0.80	856
weighted avg	0.81	0.81	0.81	856

Accuracy SVC linear: 0.81



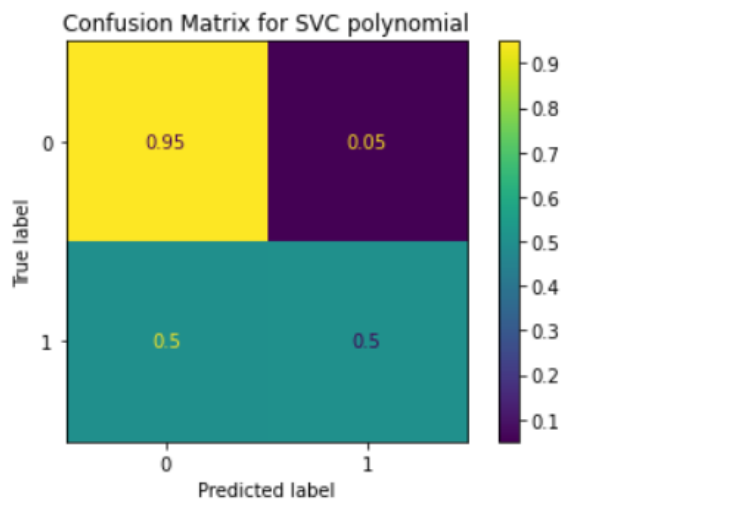
	precision	recall	f1-score	support
0	0.81	0.88	0.85	515
1	0.80	0.70	0.74	341
accuracy			0.81	856
macro avg	0.81	0.79	0.80	856
weighted avg	0.81	0.81	0.81	856

Accuracy SVC RBF: 0.79



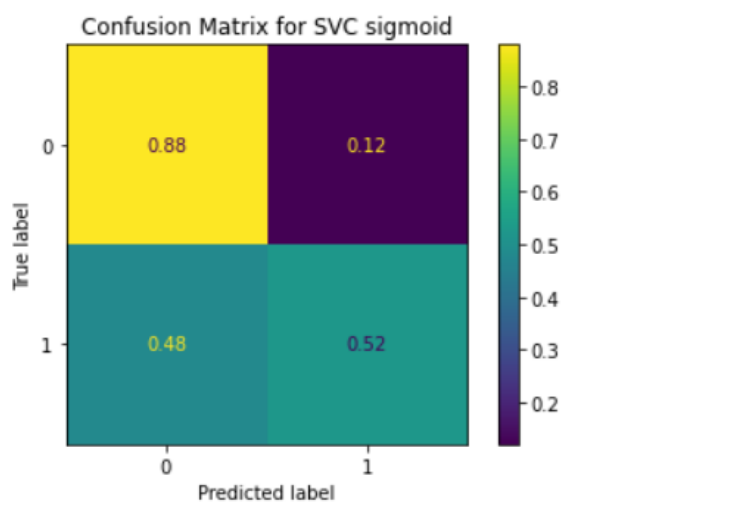
	precision	recall	f1-score	support
0	0.78	0.91	0.84	515
1	0.82	0.62	0.70	341
accuracy			0.79	856
macro avg	0.80	0.76	0.77	856
weighted avg	0.80	0.79	0.79	856

Accuracy SVC polynomial: 0.77



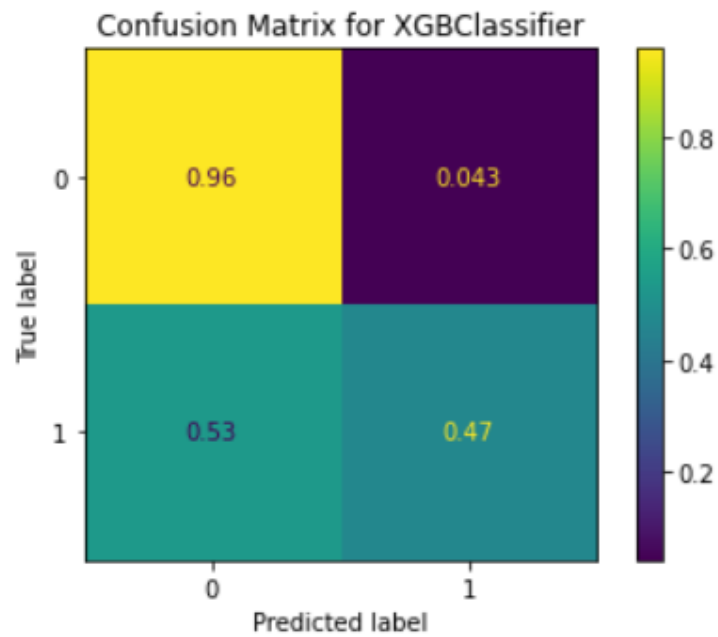
	precision	recall	f1-score	support
0	0.74	0.95	0.83	515
1	0.87	0.50	0.64	341
accuracy			0.77	856
macro avg	0.81	0.73	0.73	856
weighted avg	0.79	0.77	0.75	856

Accuracy SVC sigmoid: 0.74



	precision	recall	f1-score	support
0	0.74	0.88	0.80	515
1	0.74	0.52	0.61	341
accuracy			0.74	856
macro avg	0.74	0.70	0.71	856
weighted avg	0.74	0.74	0.73	856

Accuracy XGBClassifier: 0.76



	precision	recall	f1-score	support
0	0.73	0.96	0.83	515
1	0.88	0.47	0.61	341
accuracy			0.76	856
macro avg	0.80	0.71	0.72	856
weighted avg	0.79	0.76	0.74	856

Final Recommendations

The highest accuracies belonged to Gradient boosting, Extra Trees classifier and the random forest classifiers, having accuracies of 83%, 82% and 81% respectively. The random forest classifier is 1% more likely to make a false negative prediction on the drug persistency, which is favorable to the contrary. A physician can follow up on the patient that is predicted to stop the medication regardless of their actual drug persistency. The models also have the highest F1 scores: 0.85 0.85 and 0.84 respectively.

Github Repo:

https://github.com/Arminkhayati/dataglacier_internship