

## **Data Science Intern at Data Glacier**

# Project: Healthcare – Persistency of drug

#### Week 9

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### **Problem Description**

The objective of this project is to **understand and predict the persistency of a drug prescribed by physicians**. Persistency, in this context, refers to whether a patient, based on his/her information, will follow the prescribed medication regimen over a certain period.

Identifying factors that influence persistency is crucial for pharmaceutical companies to improve patient outcomes, reduce healthcare costs, and enhance their product offerings.

### **Business Understanding**

Persistency of medication is a critical factor in the effectiveness of treatment plans. High persistency rates generally correlate with better health outcomes, as patients are more likely to follow their treatment plans. Conversely, non-persistence can lead to worsening health conditions, increased hospitalizations, and higher overall healthcare costs.

Pharmaceutical company ABC Pharma is interested in identifying the key factors that influence whether patients persist with their medication. By leveraging machine learning to predict persistency, the company can develop strategies to improve adherence rates.

### Data Understanding

Healthcare dataset has 3424 observations and 69 features. Our intention is to build a model that predicts if a given patient will persist on his/her treatment or not. Having this, out target is the "Persistency Flag" variable, which is a binary data having values True or False depending on the other features.

Besides individual identificators and the target variable, there are other 4 buckets:

• Demographics

- Provider attributes
- Clinical factors
- Disease an treatment factors

Bucket	Variable	Variable Description	Information
Unique Row Id	Ptid	Unique ID of each patient	Type: object
-			Missing values: 0%
			Unique values: 3424
Target	Persistency_F	Flag indicating if a patient	Type: object
Variable	lag	was persistent or not	Missing values: 0%
			Unique values: 2
			Values: ['Persistent', 'Non-Persistent']
			Mode: 'Non-Persistent'
Demographics	Age_Bucket	Age of the patient during	Type: object
0 1		their therapy	Missing values: 0%
			Unique values: 4
			Values: ['>75', '55-65', '65-75', '<55']
			Mode: '>75' (42.03%)
	Race	Race of the patient from the	Type: object
		patient table	Missing values: 2.83% as
		1	'Other/Unknown'
			Unique values: 4
			Values: ['Caucasian', 'Asian',
			'Other/Unknown', 'African American']
			Mode: 'Caucasian' (91.94%)
	Region	Region of the patient from the	Type: object
	C	patient table	Missing values: 1.75% as
		1	'Other/Unknown'
			Unique values: 5
			Values: ['West', 'Midwest', 'South',
			'Other/Unknown', 'Northeast']
			Mode: 'Midwest' (40.39%)
	Ethnicity	Ethnicity of the patient from	Type: object
		the patient table	Missing values: 2.66% as 'Unknown'
		1	Unique values: 3
			Values: ['Not Hispanic', 'Hispanic',
			'Unknown']
			Mode: 'Not Hispanic' (94.48%)
	Gender	Gender of the patient from	Type: object
		the patient table	Missing values: 0%
			Unique values: 2
			Values: ['Male', 'Female']
			Mode: 'Female' (94.43%)
	Idn_Indicator	Flag indicating patients	Type: object
		mapped to IDN	Missing values: 0%
		PP-0-0-12-1,	Unique values: 2

			Values: ['Y', 'N']
			Mode: 'Y' (74.68%)
Provider	Ntm_Speciail	Specialty of the HCP that	Type: object
Attributes	ty	prescribed the NTM Rx	Missing values: 9.05% as 'Unknown'
			Unique values: 36
			Values: ['GENERAL PRACTITIONER',
			'Unknown', 'ENDOCRINOLOGY',
			'RHEUMATOLOGY', 'ONCOLOGY',
			'PATHOLOGY', [], 'VASCULAR
			SURGERY', 'CARDIOLOGY',
			'NUCLEAR MEDICINE']
			Mode: 'GENERAL PRACTITIONER'
			(44.83%)
	Ntm_Speciali	Specialty flag of the HCP that	Type: object
	st_Flag	prescribed the NTM Rx	Missing values: 0%
			Unique values: 2
			Values: ['Others', 'Specialist']
			Mode: 'Others' (58.79%)
	Ntm_Speciali	Specialty bucket of the HCP	Type: object
	ty_Bucket	that prescribed the NTM Rx	Missing values: 0%
			Unique values: 3
			Values:
			['OB/GYN/Others/PCP/Unknown',
			'Endo/Onc/Uro', 'Rheum']
			Mode: 'OB/GYN/Others/PCP/Unknown',
			(61.45%)
Clinical	Tscore_Buck	T Score of the patient prior of	Type: object
Factors	et_Prior_Ntm	the NTM Rx	Missing values: 0%
			Unique values: 2
			Values: ['>-2.5', '<=-2.5']
	Tscore_Buck	T Score of the patient at the	Type: object
	et_During_R	time of the NTM Rx (within	Missing values: 43% as 'Unknown'
	X	2 years prior from rxdate)	Unique values: 3
			Values: ['<=-2.5', 'Unknown', '>-2.5']
	Change_T_S	Change in Tscore before	Type: object
l	core	starting with any therapy and	Missing values: 43% as 'Unknown'
		after receiving therapy	Unique values: 4
			Values: ['No change', 'Unknown',
			'Worsened', 'Improved']
	Risk_Segmen	Risk Segment of the patient	Type: object
	t_Prior_Ntm	prior of the NTM Rx	Missing values: 0%
			Unique values: 2
			Values: ['VLR_LR', 'HR_VHR']
	Risk_Segmen	Risk Segment of the patient at	Type: object
	t_ During_Rx	the time of the NTM Rx	Missing values: 43% as 'Unknown'
		(within 2 years days prior	Unique values: 3
		from rxdate)	Values: ['VLR_LR, 'Unknown',
			'HR_VHR']

	Change_Risk _Segment	Change in Risk Segment before starting with any therapy and after receiving therapy	Type: object Missing values: 65% as 'Unknown' Unique values: 4 Values: ['No change', 'Unknown', 'Worsened', 'Improved']
	NTM - Multiple Risk Factors	Flag indicating if patient falls under multiple risk category (having more than 1 risk) at the time of the NTM Rx (within 365 days prior from rxdate)	Type: object Missing values: 0% Unique values: 2 Values: ['Y', 'N']
	Dexa_Freq_ During_Rx	Number of DEXA scans taken prior to the first NTM Rx date (within 365 days prior from rxdate)	Type: integer Missing values: 0% Unique values: 58 Values info: mean – 3.01, std – 8.14, min – 0, 50% – 0, max – 146 Mode: 0 (72.66%)
	Dexa_During _Rx	Flag indicating if the patient had a Dexa Scan during their first continuous therapy	Type: object Missing values: 0% Unique values: 2 Values: ['Y', 'N'] Mode: 'N' (72.66%)
	Frag_Frac_Pr ior_Ntm	Flag indicating if the patient had a recent fragility fracture (within 365 days prior from rxdate)	Type: object Missing values: 0% Unique values: 2 Values: ['Y', 'N'] Mode: 'N' (83.88%)
	Frag_Frac_D uring_Rx	Flag indicating if the patient had fragility fracture during their first continuous therapy	Type: object Missing values: 0% Unique values: 2 Values: ['Y', 'N'] Mode: 'N' (87.82%)
	Gluco_Recor d_Prior_Ntm	Flag indicating usage of Glucocorticoids (>=7.5mg strength) in the one year lookback from the first NTM Rx	Type: object Missing values: 0% Unique values: 2 Values: ['Y', 'N'] Mode: 'N' (87.82%)
	Gluco_Recor d_During_Rx	Flag indicating if the patient had a Glucocorticoid usage during the first continuous therapy	Type: object Missing values: 0% Unique values: 2 Values: ['Y', 'N'] Mode: 'N' (76.49%)
Disease/Treat ment Factor	Injectable_Ex perience_Dur ing_Rx	Flag indicating any injectable drug usage in the recent 12 months before the NTM OP Rx	Type: object Missing values: 0% Unique values: 2 Values: ['Y', 'N'] Mode: 'Y' (73.66%)

NTM - Risk	Risk Factors that the patient	Type: object
Factors	is falling into. For chronic	Missing values: 0%
	Risk Factors complete	Unique values: 2
	lookback to be applied and	Values: ['Y', 'N']
	for non-chronic Risk Factors,	. , ,
	one year lookback from the	
	date of first OP Rx	
NTM -	Comorbidities are divided	Type: object
Comorbidity	into two main categories -	Missing values: 0%
	Acute and chronic, based on	Unique values: 2
	the ICD codes. For chronic	Values: ['Y', 'N']
	disease we are taking	
	complete look back from the	
	first Rx date of NTM therapy	
	and for acute diseases, time	
	period before the NTM OP	
	Rx with one year lookback	
	has been applied	
NTM -	Concomitant drugs recorded	Type: object
Concomitanc	prior to starting with a	Missing values: 0%
у	therapy(within 365 days prior	Unique values: 2
	from first rxdate)	Values: ['Y', 'N']
Adherent_Fla	Adherence for the therapies	Type: object
g		Missing values: 0%
		Unique values: 2
		Values: ['Adherent', 'Non-Adherent']
		Mode: 'Adherent' (94.94%)
Count_Of_Ri	Total number of risks	Type: integer
sks		Missing values: 0%
		Unique values: 8
		Values: [0, 1, 2, 3, 4, 5, 6, 7]
		Mode: 1 (36.27%)

# Data Cleaning and Transformation

#### **Missing values:**

- *Race* using of mode as substitution, only 2.83% are 'Other/Unknown', so it's quite safe to use the mode to fill in the missing values.
- Region using of Region mode for 'Not Hispanic', because only 1.75% missing values and 100% of them are of Ethnicity 'Not Hispanic'.

- *Ethnicity* using of mode as substitution, only 2.66% missing values, so it's quite safe to use the mode to fill in the missing values.
- *Ntm\_Speciality* we will try 2 approaches:
- 1) keeping unknowns as a category since it accounts for less than 9.05% of data and see how it relates to other variables.
- 2) the categories that accounts for less than 0.01 of the number of observations will be replaced with 'OTHER' category.
- Risk\_Segment\_During\_Rx, Change\_T\_Score, Change\_Risk\_Segment these variables have more than 40% of missing values, consequently they will be removed.
- *Tscore\_Bucket\_During\_Rx* we will try 2 approaches:
- 1) remove this feature as it has more than 40% of missing values
- 2) replace unknown values with values using 'Tscore\_Bucket\_Prior\_Ntm' column.

#### **Outliers:**

- We will try 2 approaches:
- 1) detect and remove outliers using IQR method.
- 2) keep outliers.

#### Categorical data:

- *Injectable Experience, Risk Factors, Comorbidity and Concomitancy* (group of variables) 'Y' will be replaced with 1 and 'N' with 0.
- *Tscore\_Bucket\_Prior\_Ntm* '>-2.5' will be replaced with 1 and '<=-2.5' with 0.
- *Risk\_Segment\_Prior\_Ntm* 'VLR\_LR' will be replaced with 1 and 'HR\_VHR' with 0.
- *Ptid* will be removed, as it has the number of unique values equal to the number of observations and this won't help modeling.

- *Age\_Bucket* will be later encoded using Ordinal Encoder with the following categories: ['<55', '55-65', '65-75', '>75']
- All other categorical features will be encoded using Label Encoder or One-hot Encoder later.

## GitHub Repo Link

Project Link: <a href="https://github.com/kkudzelich/Data-Science-Intern">https://github.com/kkudzelich/Data-Science-Intern</a>

Submitted by: Kseniya Kudzelich

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