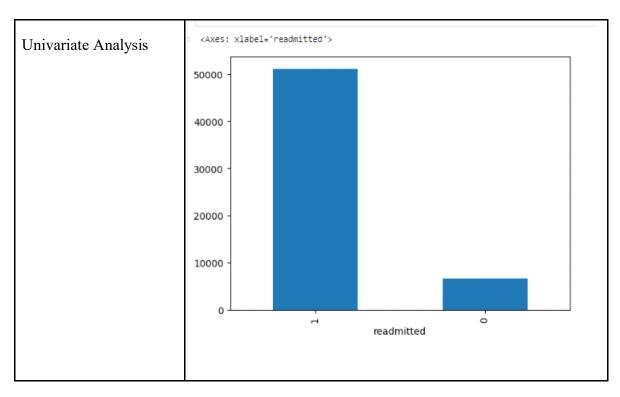
## **Data Collection and Preprocessing Phase**

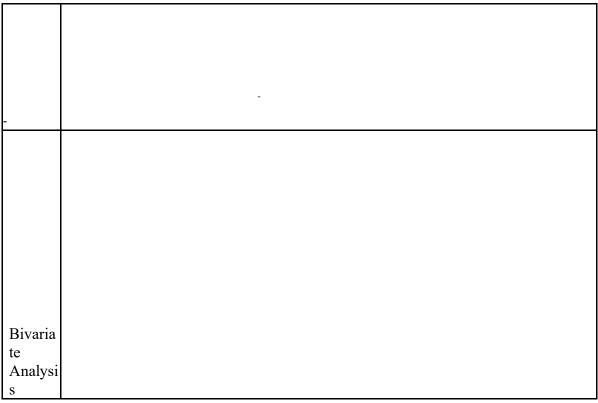
Date	20 JUNE 2024
Team ID	team-739701
Project Title	HOSPITAL READMISSION PREDICTOR USING MACHINE LEARNING
Maximum Marks	6 Marks

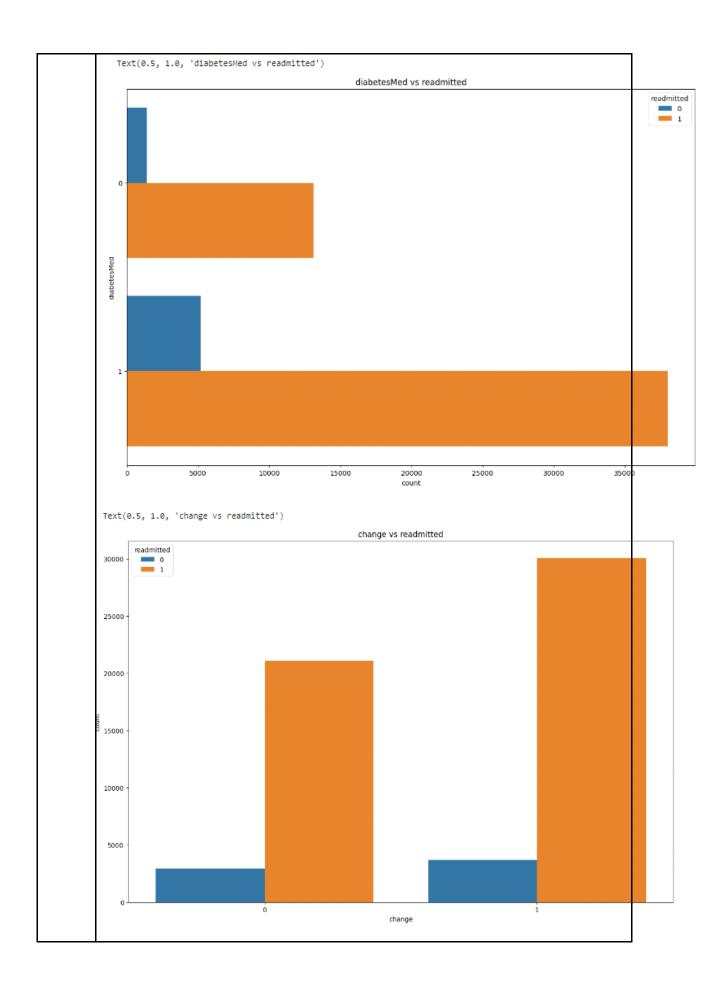
## **Data Exploration and Preprocessing Report**

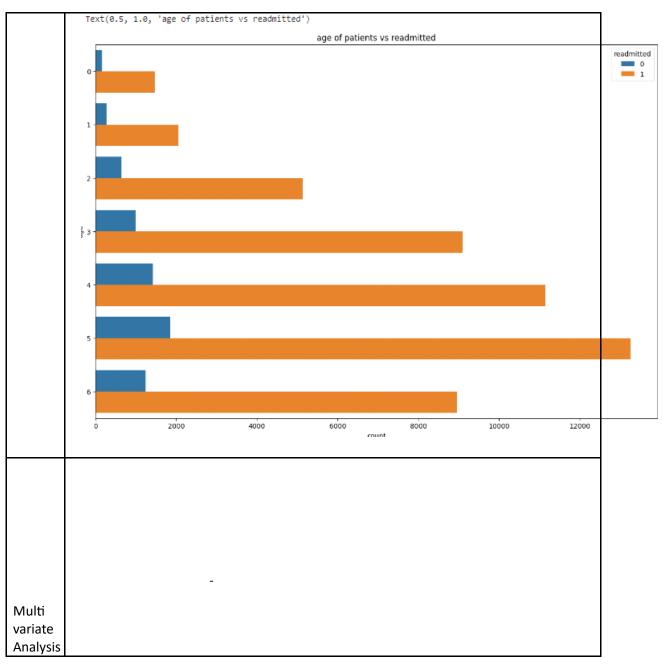
Dataset variables will be statistically analyzed to identify patterns and outliers, with Python employed for preprocessing tasks like normalization and feature engineering. Data cleaning will address missing values and outliers, ensuring quality for subsequent analysis and modeling, and forming a strong foundation for insights and predictions.

Section	Description									
	5 rc	nension ows × 2 istics:	_	lun	nns <u>Desc</u>	<u>riptive</u>				
		race	gender	age	admission_type_id	discharge_disposition_id	admission_source_id	time_in_hospital	num_lab_procedure	
	0	Caucasian	Female	[0- 30)	Others	Others	Referral	-1.632460	-0.10847	
Data Overview	1	Caucasian	Female	[0- 30)	Emergency	Discharged to home	Emergency Room	-0.271144	0.80852	
	<b>2</b> A	fricanAmerican	Female	[0- 30)	Emergency	Discharged to home	Emergency Room	-0.825587	-1.62961	
	3	Caucasian	Male	[30- 40)	Emergency	Discharged to home	Emergency Room	-0.825587	0.04419	
	4	Caucasian	Male	[40- 50)	Emergency	Discharged to home	Emergency Room	-1.632460	0.40069	
	5 row	5 rows × 29 columns								









Outliers and Anomalies	-								
Data Preproc	es				shots admission_type_id	discharge disposition id	admission source id	time in hospital	num lab proced
	0	Caucasian		[0-	Others	Others	Referral	-1,632460	-0.108
	1	Caucasian	Female	[0- 30)	Emergency	Discharged to home	Emergency Room	-0.271144	0.808
Loading Data	2	AfricanAmerican	Female	[0- 30)	Emergency	Discharged to home	Emergency Room	-0.825587	-1.625
	3	Caucasian	Male	[30- 40)	Emergency	Discharged to home	Emergency Room	-0.825587	0.044
	4	Caucasian	Male	[40- 50)	Emergency	Discharged to home	Emergency Room	-1.632460	0.400

	Γ		1
		<pre>data['num_medications'].fillna(data['num_medications'].mean(),inplace=True)</pre>	_
		data['number_outpatient'].fillna(data['number_outpatient'].mean(),inplace=True)	
		data['number_emergency'].fillna(data['number_emergency'].mean(),inplace=True)	
Handling		data['number_diagnoses'].fillna(data['number_diagnoses'].mean(),inplace=True)	
Missing Data		data['age_derived'].fillna(data['age_derived'].mean(),inplace=True)	
		data['count_Steady'].fillna(data['count_Steady'].mean(),inplace=True)	
		data['count_Down'].fillna(data['count_Down'].mean(),inplace=True)	
		data['count_Up'].fillna(data['count_Up'].mean(),inplace=True)	
		<pre>data['number_inpatient'].fillna(data['number_inpatient'].mean(),inplace=True)</pre>	
	:	le=LabelEncoder()	1
	:	data['gender']=le.fit_transform(data['gender'])	
Data Transformatio n	:	<pre>data['age']=le.fit_transform(data['age'])</pre>	
	:	data['admission_type_id']=le.fit_transform(data['admission_type_id'])	
	:	data['discharge_disposition_id']=le.fit_transform(data['discharge_disposition_id'])	
	:	data['admission_source_id']=le.fit_transform(data['admission_source_id'])	
	:	<pre>data['diag_1']=le.fit_transform(data['diag_1'])</pre>	
	:	<pre>data['diag_2']=le.fit_transform(data['diag_2'])</pre>	
	:	<pre>data['diag_3']=le.fit_transform(data['diag_3'])</pre>	
	:	data['metformin']=le.fit_transform(data['metformin'])	
	:	data['repaglinide']=le.fit_transform(data['repaglinide'])	
	:	<pre>data['glipizide']=le.fit_transform(data['glipizide'])</pre>	
	:	<pre>data['insulin']=le.fit_transform(data['insulin'])</pre>	
	:	<pre>data['change']=le.fit_transform(data['change'])</pre>	
	:	data['diabetesMed']=le.fit_transform(data['diabetesMed'])	
	:	<pre>data['readmitted']=le.fit_transform(data['readmitted'])</pre>	

Feature Engineering	Attached the codes in final submission