

# Predicting Chronic Obstructive Pulmonary Disease (COPD) in Nepal Project

The approaches mentioned below were followed to develop the system for COPD, including data collection, preprocessing, feature engineering, model development, and deployment.

## Step 1: Problem Statement and Objectives

### Problem Statement:

- To predict the possibility of a patient developing COPD based on various risk factors and patient characteristics.

### Objectives:

1. Collect and preprocess data relevant to COPD.
2. Identify and engineer significant features contributing to COPD.
3. Develop a predictive model to estimate the risk of COPD.
4. Evaluate the model's performance and refine it.
5. Deploy the model for practical use in a clinical or public health setting.

## Step 2: Data Collection

Various sites were visited and research was conducted to find good datasets, but all of them did not have a target variable: whether the patient has COPD or doesn't have COPD, so the “***synthetic\_COPD\_data.csv***” was used for the project.

### Data sources researched and identified

- Health Data Portals: WHO, World Bank, Nepal's Ministry of Health
- Research Papers: Google Scholar
- Public Datasets: Kaggle, Data.gov, Open Data Nepal
- Hospitals and Clinics

## Step 3: Data Preprocessing

The initially found data, i.e. **“finalalldata.csv”** and **“Dataset\_PowerBI.xlsx”** were studied and merged as shown in the **“data\_prep.ipynb”** file by reducing the columns and only using the necessary ones. But, later **“synthetic\_COPD\_data.csv”** was used for the project.

- Sample of **finalalldata.csv**

```
csv_data.head()
```

	uid	label	sex	age	bmi	smoke	location	rs10007052	rs8192288	rs20541	rs12922394	rs2910164	rs161976	rs473892	rs159497
0	copdcontrol1	0	2	28	19.22	0	4.63	1.671	1.0	0.448632	0.42328	1.000	1.0	1.473	1.000000
1	copdcontrol69	0	1	53	20.44	0	4.63	1.671	1.0	1.000000	0.65060	1.416	1.0	1.000	1.000000
2	copdcontrol68	0	1	58	20.45	1	4.63	1.000	1.0	0.669800	1.00000	1.416	1.0	1.473	1.000000
3	copdcontrol85	0	2	30	20.70	0	4.63	1.671	1.0	0.448632	1.00000	1.416	NaN	1.473	1.000000
4	copdcontrol78	0	1	55	20.76	1	4.63	1.671	1.0	0.669800	0.65060	1.000	1.0	1.473	2.088025

- Sample of **“Dataset\_PowerBI.xlsx”**

```
excel_data.head()
```

	Patient ID	Gender	Age	Height(cm)	Height(in)	weight kg	weight (lb)	BMI	Smoker	Comorbidity1	...	RR During vigorous exercise	Baseline FEV1	Current FEV1	Baseline VO2	Rd
0	3	Male	88	178	69.954	75	165.300	23.746689	No	NaN	...	74	0.729397	0.89 L	2.146	3.1
1	4	Female	60	167	65.631	79	174.116	28.416852	No	NaN	...	41	1.358489	1.179 L	1.780	13.1
2	8	Male	55	178	69.954	115	253.460	36.411591	No	Pulmonary Hypertension	...	47	1.588316	1.744 L	3.083	8.3
3	9	Female	55	170	66.810	81	178.524	28.117001	No	NaN	...	59	0.549915	0.526 L	1.963	5.0
4	10	Male	65	169	66.417	75	165.300	26.343269	No	NaN	...	42	1.337281	1.149 L	2.616	5.1

5 rows × 59 columns

- Sample of the merged dataset with relevant columns

	sex	bmi	age	location	smoke	Height(cm)	weight kg	BMI	Smoker	Gender
0	1	20.44	53	4.63	0	172	69	23.397745	Yes	Male
1	1	20.44	53	4.63	0	179	120	37.571367	No	Male
2	1	20.44	53	4.63	0	173	96	32.178133	Yes	Female
3	1	20.44	53	4.63	0	175	98	32.101978	No	Male
4	1	20.44	53	4.63	0	170	89	30.893988	Yes	Male

## Step 4: Exploratory Data Analysis (EDA)

Then, EDA was performed on the datasets following the mentioned steps:

1. Loading the datasets and studying them

Sample of the dataset used:

```
#Load the datasets
path = r"../Data/synthetic_COPD_data.csv"
df = pd.read_csv(path)
df.head()
```

	Age	Gender	Smoking_Status	Biomass_Fuel_Exposure	Occupational_Exposure	Family_History_COPD	BMI	Location	Air_Pollution_Level	Respiratory_Infections_Childhood
0	31	Male	Former	1	1	1	27.56	Lalitpur	84	1
1	60	Male	Never	1	0	0	30.30	Pokhara	131	0
2	33	Male	Former	0	0	1	28.45	Pokhara	123	0
3	36	Female	Current	1	0	0	27.49	Kathmandu	253	1
4	58	Male	Never	0	0	0	25.49	Pokhara	117	0

Then, a plot was set for better visibility and the data was studied along with the columns and values that were present using **df.head()** and **df.info()**.

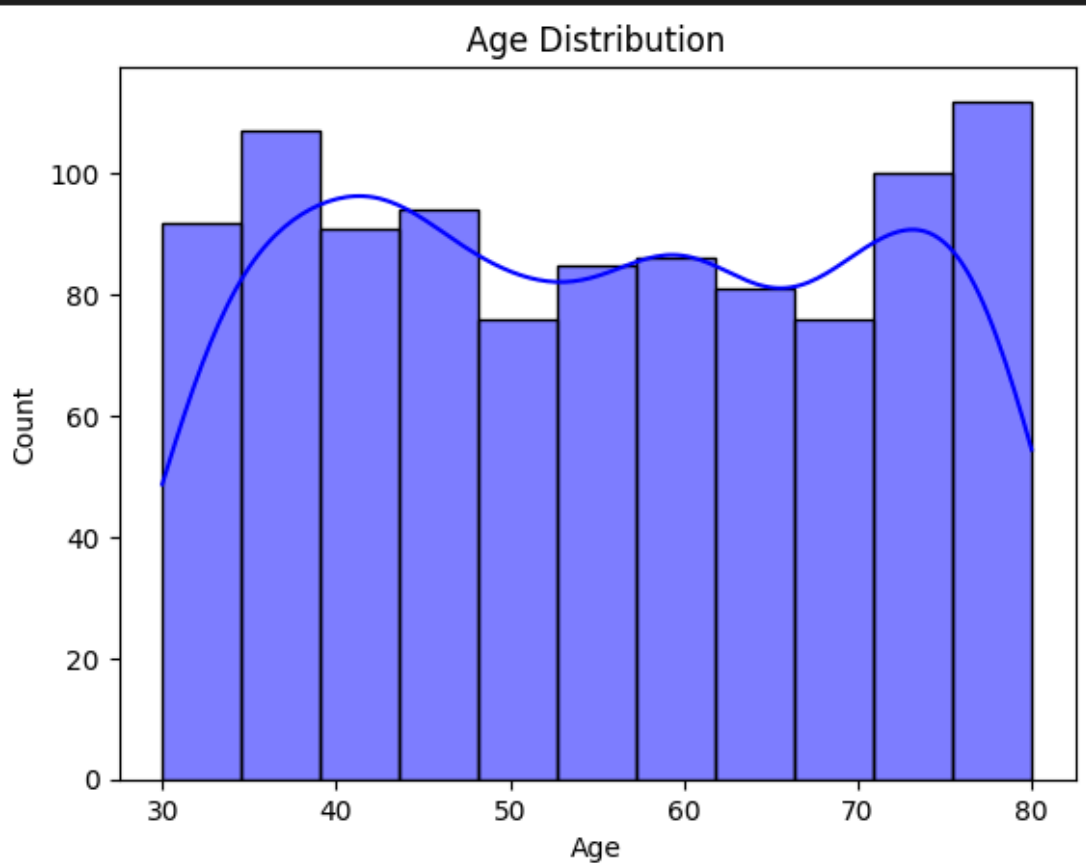
```
#Get info about the data
df.info()
```

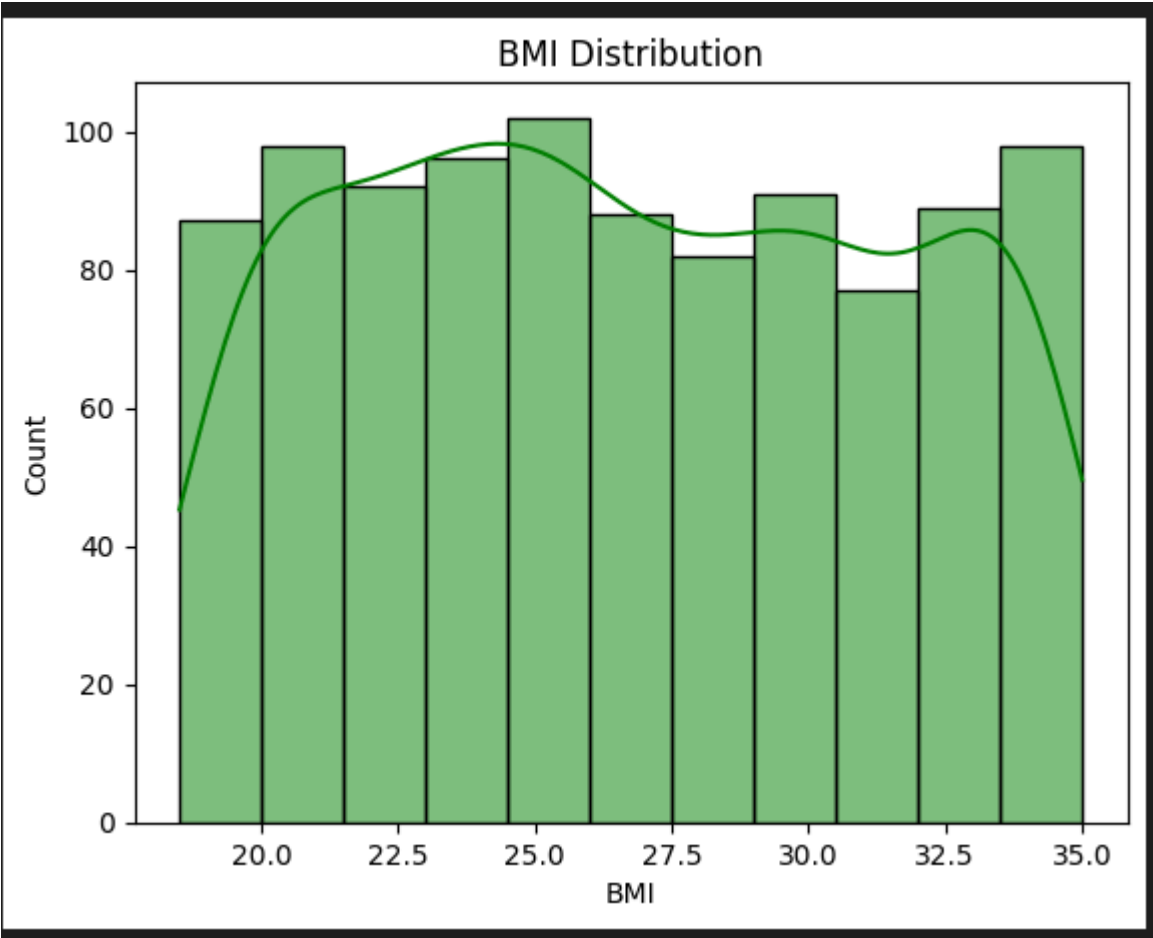
```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1000 entries, 0 to 999
Data columns (total 11 columns):
 #   Column                                     Non-Null Count  Dtype  
---  -
 0   Age                                       1000 non-null   int64  
 1   Gender                                   1000 non-null   object  
 2   Smoking_Status                           1000 non-null   object  
 3   Biomass_Fuel_Exposure                    1000 non-null   int64  
 4   Occupational_Exposure                    1000 non-null   int64  
 5   Family_History_COPD                     1000 non-null   int64  
 6   BMI                                       1000 non-null   float64 
 7   Location                                  1000 non-null   object  
 8   Air_Pollution_Level                     1000 non-null   int64  
 9   Respiratory_Infections_Childhood         1000 non-null   int64  
10  COPD_Diagnosis                           1000 non-null   int64  
dtypes: float64(1), int64(7), object(3)
memory usage: 86.1+ KB
```

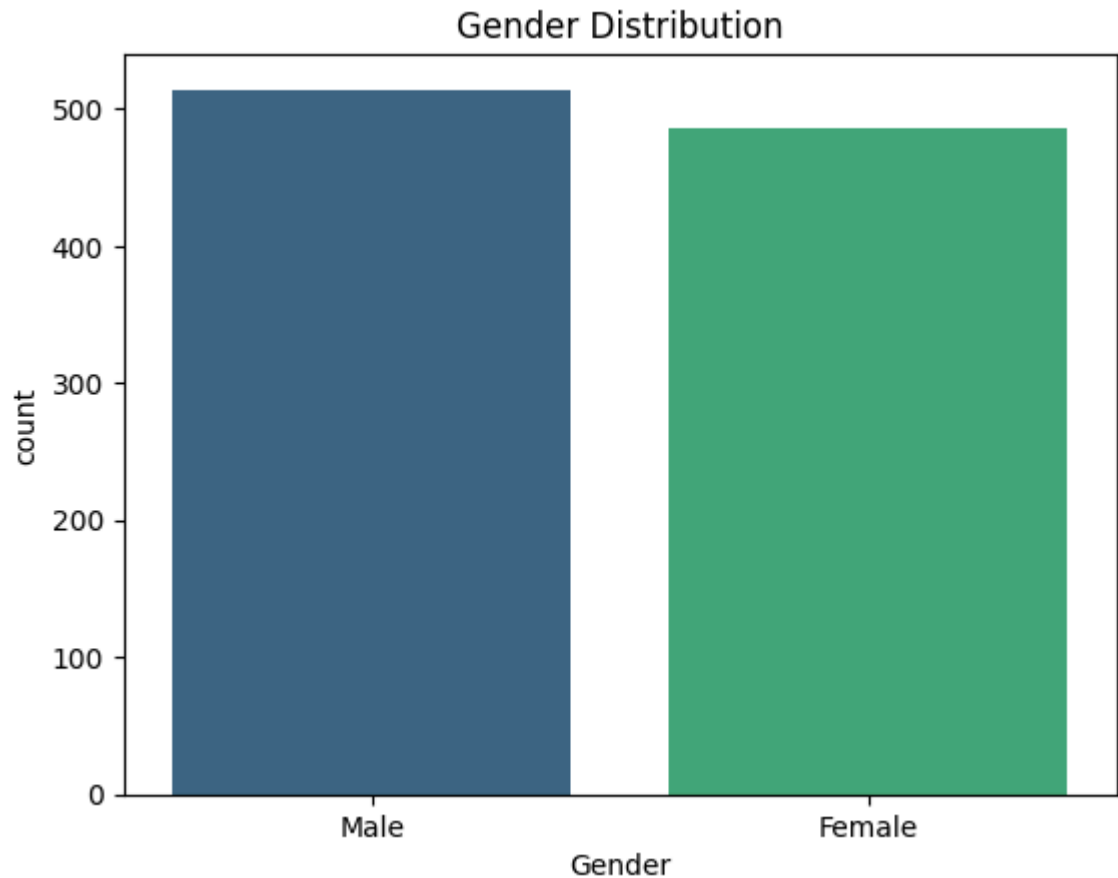
2. Then, **Univariate Analysis** was performed on the datasets to further study the data, the analysis was performed on various columns such as age, BMI, gender, smoking status, etc. Some examples:

```
#Age distributions in the data
sns.histplot(df['Age'], kde = True, color = 'blue')
plt.title('Age Distribution')

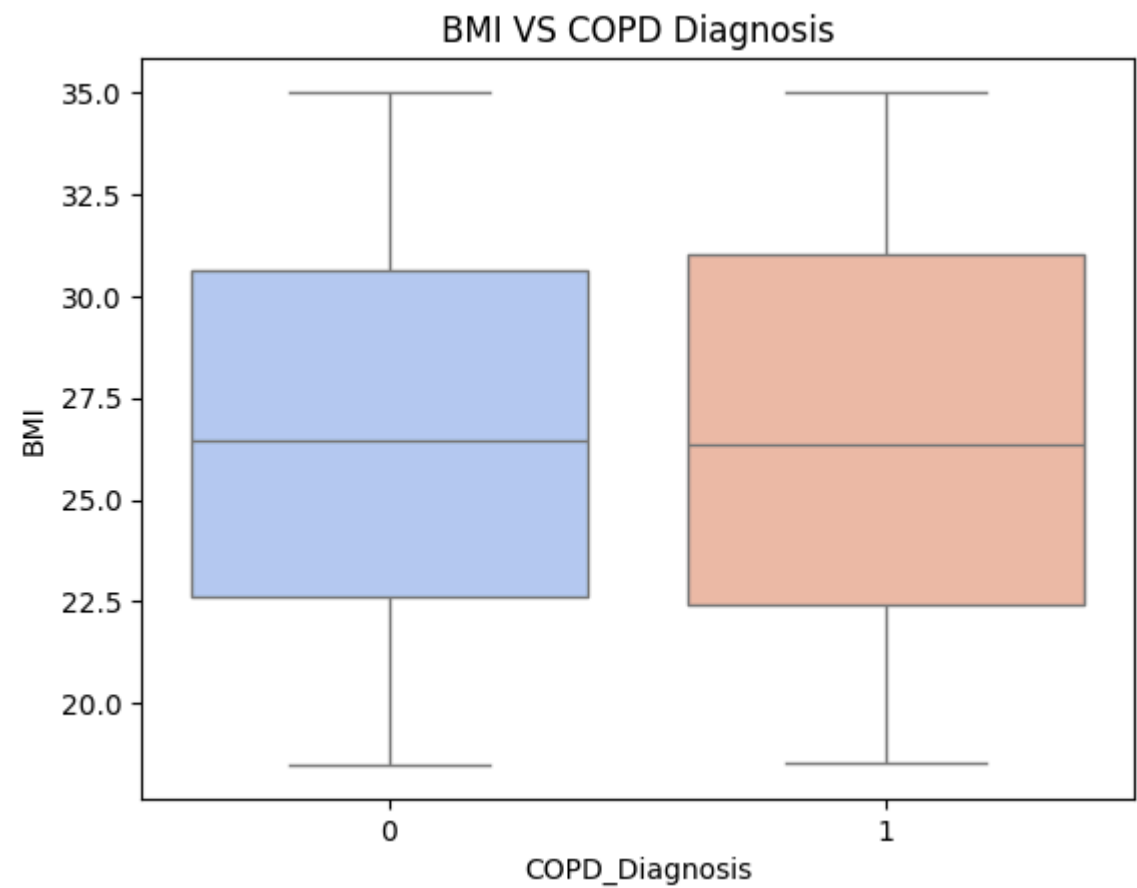
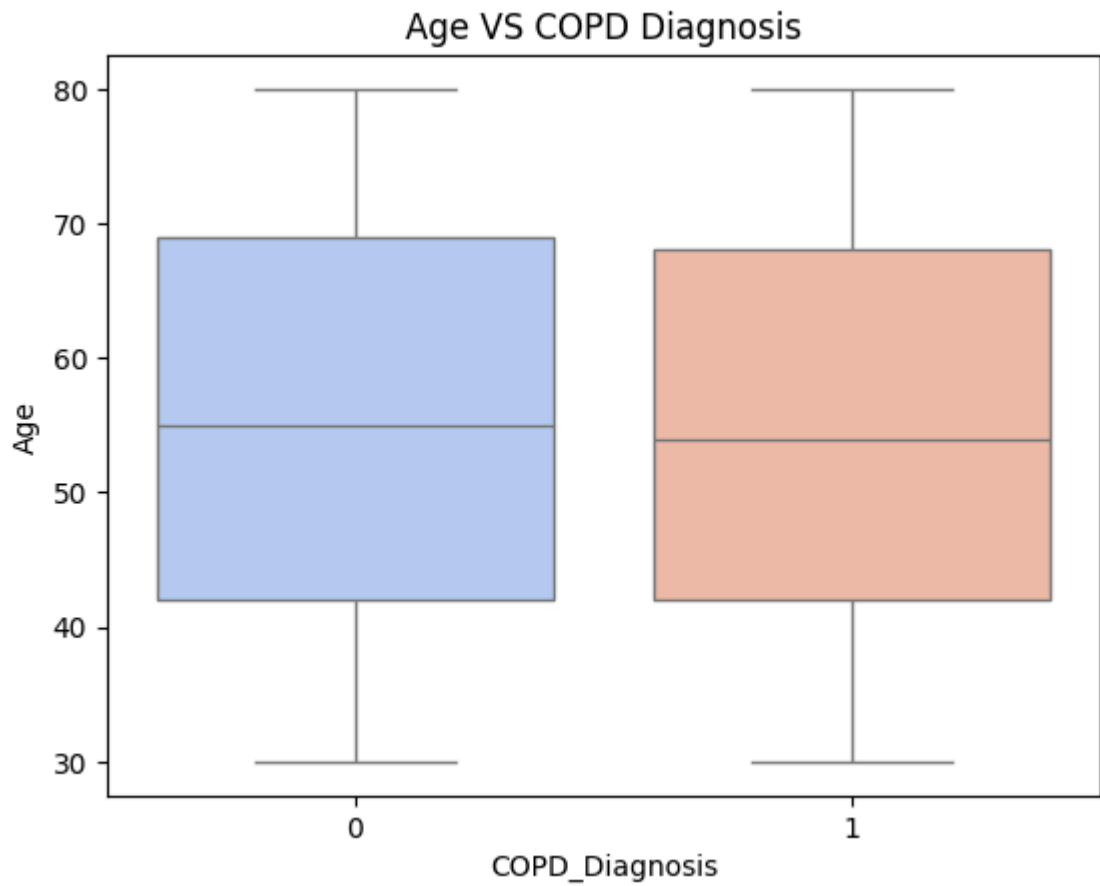
text(0.5, 1.0, 'Age Distribution')
```



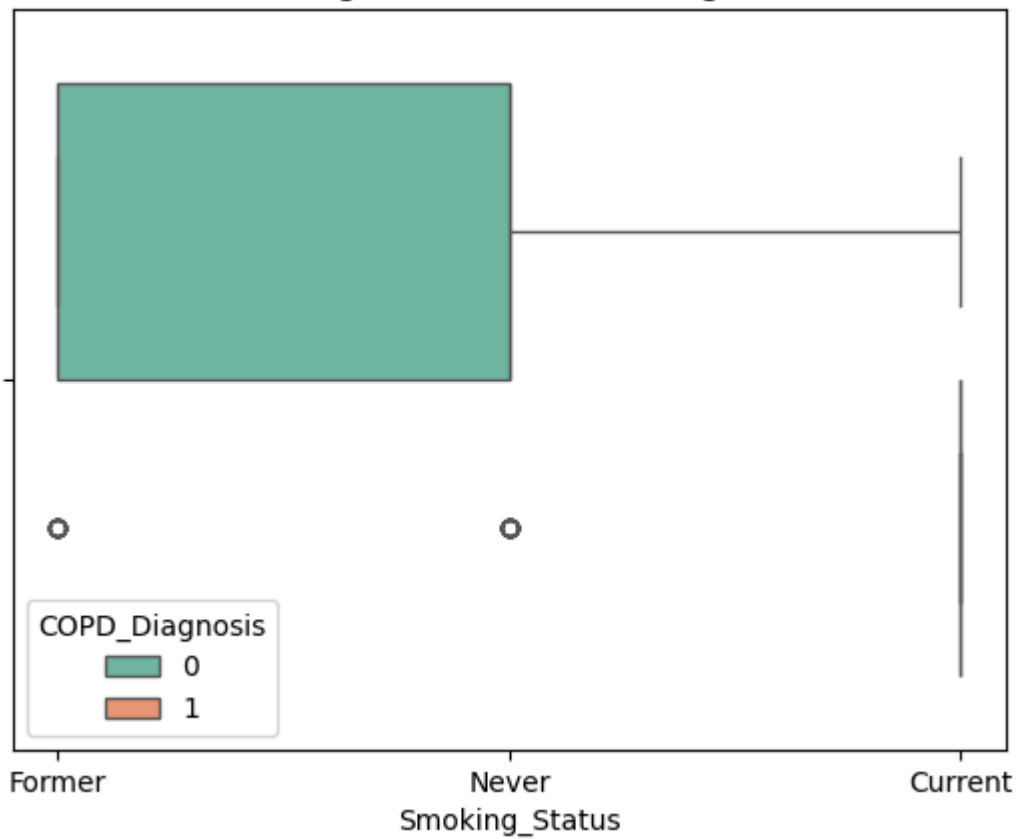




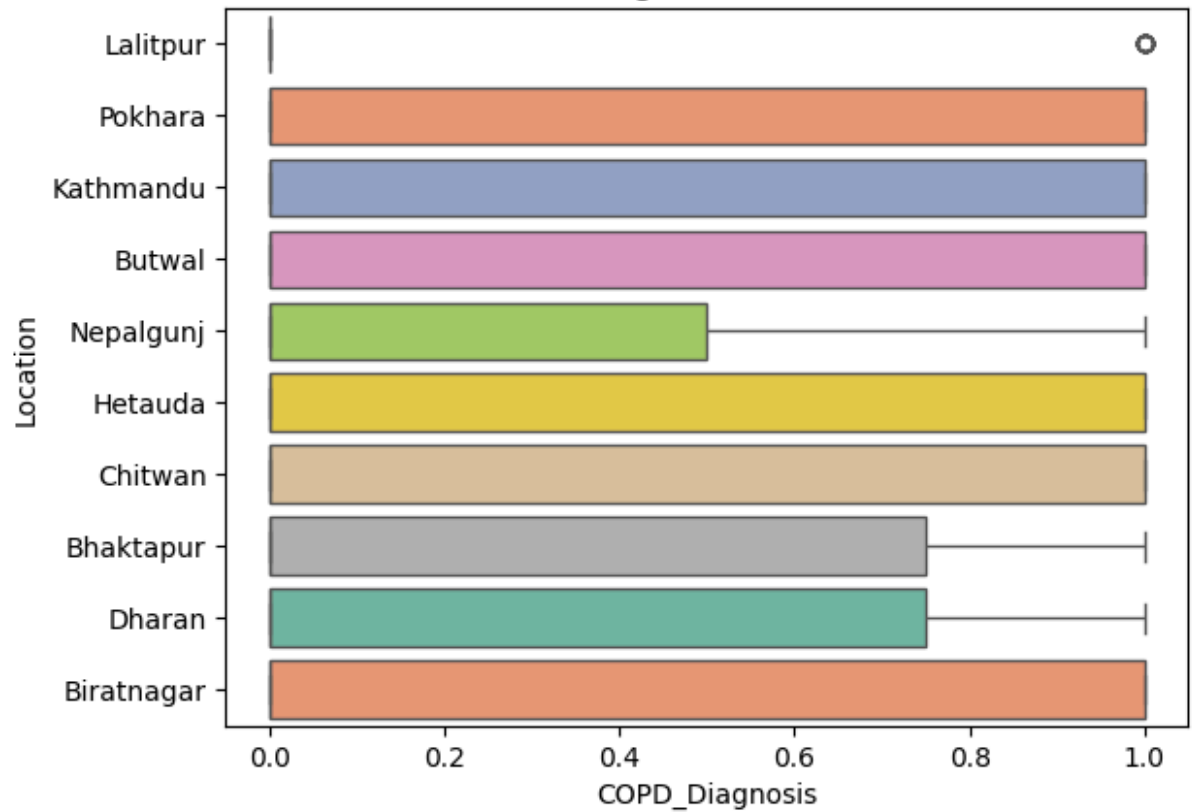
3. After that, **Bivariate Analysis** was done to study the relationship between various features of the datasets, i.e. Age versus COPD Diagnosis, BMI versus COPD Diagnosis, Smoking Status versus COPD Diagnosis, using boxplot and countplot etc. Some examples:



### Smoking Status VS COPD Diagnosis

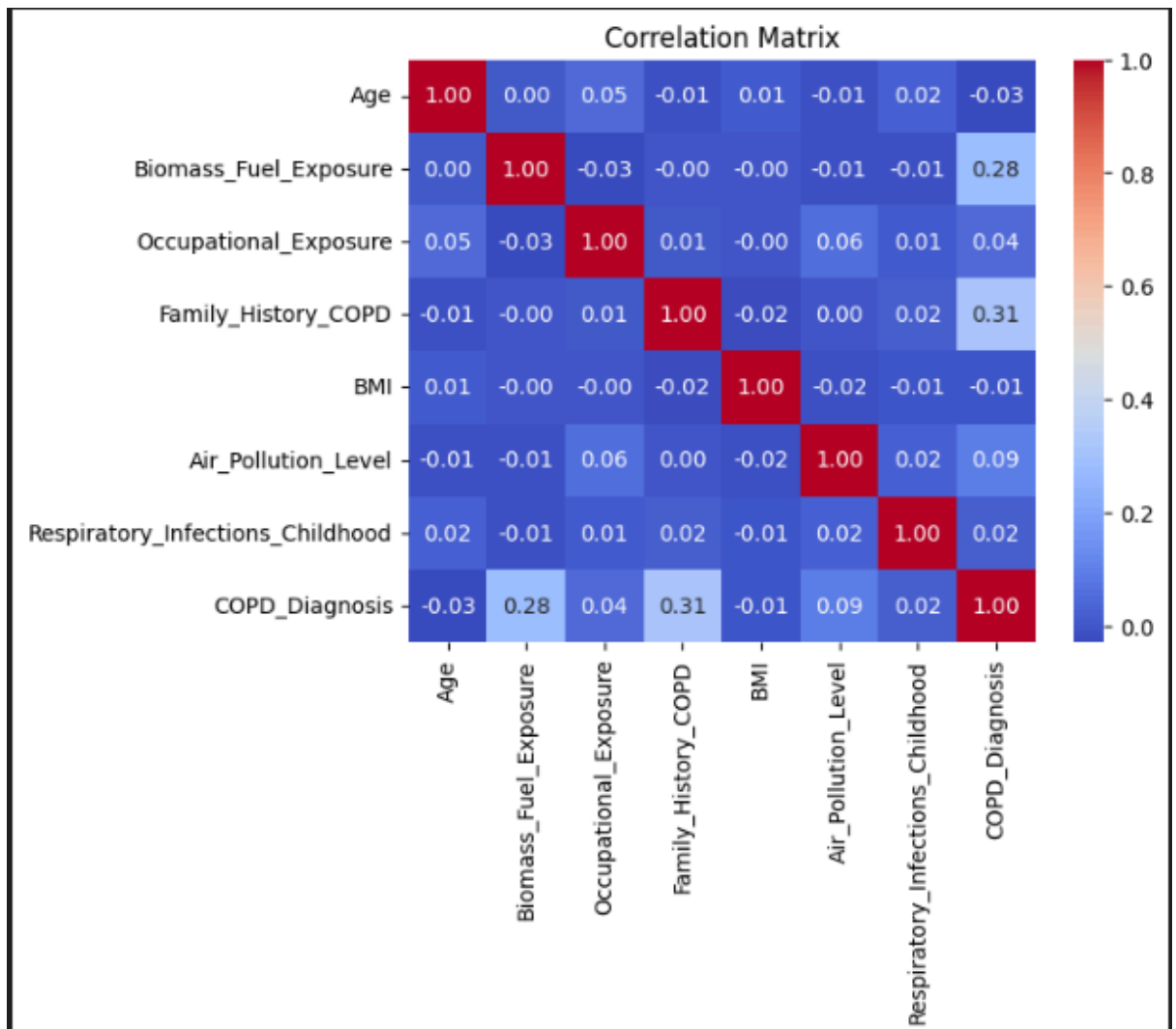


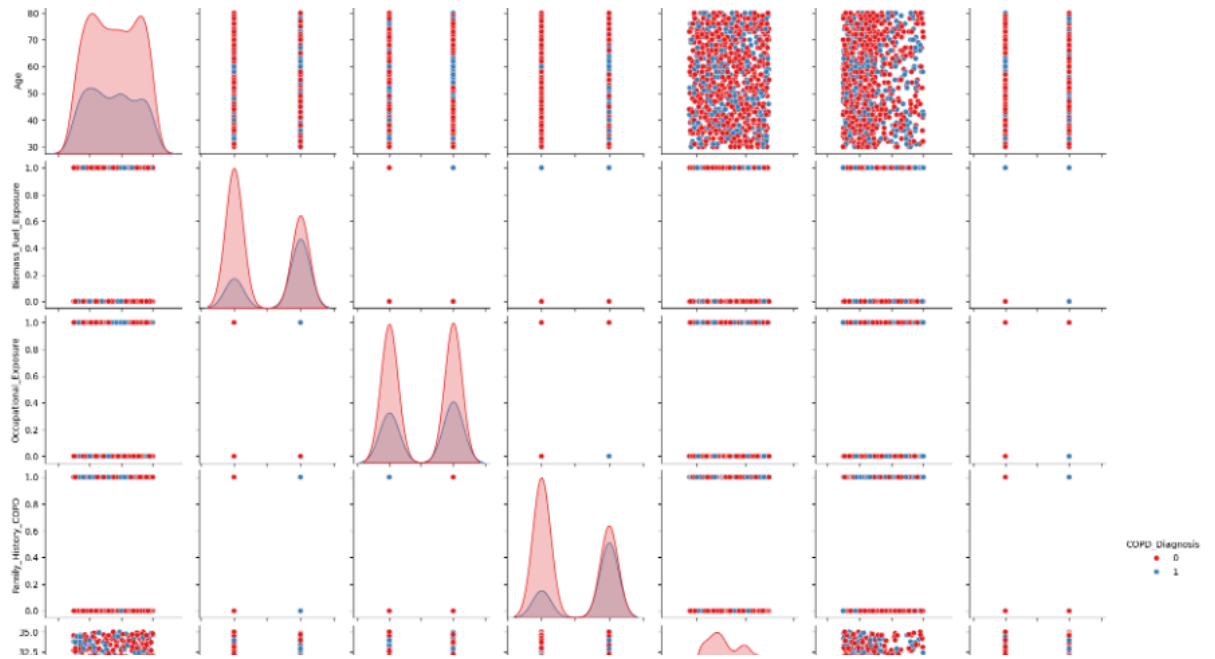
### COPD Diagnosis VS Location





4. Then, we perform **Multivariate Analysis** to see the relation between the target variable COPD\_Diagnosis and rest of the features using correlation matrix and pairplot. Some examples:





## Step 5: Feature Engineering

Based on the datasets and research for the domain that was being worked on, various new features were created and some of the existing ones were updated by using **One-Hot Encoding**.

- Age Category
- BMI Category - Overweight, Normal Weight, Underweight
- Pollution Risk Score - Based on Air Pollution Level
- Smoking Status Encoding
- Interaction Features - Smoking Pollution Interaction based on Smoking Status and Air Pollution Level
- Location Encoding - Categories have to be passed as numbers, changing categories to numerical values

#	Column	Non-Null Count	Dtype
0	Age	1000 non-null	int64
1	Gender	1000 non-null	object
2	Smoking_Status	1000 non-null	object
3	Biomass_Fuel_Exposure	1000 non-null	int64
4	Occupational_Exposure	1000 non-null	int64
5	Family_History_COPD	1000 non-null	int64
6	BMI	1000 non-null	float64
7	Air_Pollution_Level	1000 non-null	int64
8	Respiratory_Infections_Childhood	1000 non-null	int64
9	COPD_Diagnosis	1000 non-null	int64
10	Age_Category	980 non-null	category
11	BMI_Category	1000 non-null	category
12	Pollution_Risk_Score	1000 non-null	int32
13	Smoking_Status_Encoded	1000 non-null	float64
14	Gender_Encoded	1000 non-null	int64
15	Smoking_Pollution_Interaction	1000 non-null	float64
16	Location_Biratnagar	1000 non-null	bool
17	Location_Butwal	1000 non-null	bool
18	Location_Chitwan	1000 non-null	bool
19	Location_Dharan	1000 non-null	bool
...			
23	Location_Nepalgunj	1000 non-null	bool
24	Location_Pokhara	1000 non-null	bool

Then, after that, all the features or columns present were studied and then encoded, updated and dropped as necessary since we can only work with integer, float and boolean data-types as shown in the image below:

#	Column	Non-Null Count	Dtype
0	Age	1000 non-null	int64
1	Biomass_Fuel_Exposure	1000 non-null	int64
2	Occupational_Exposure	1000 non-null	int64
3	Family_History_COPD	1000 non-null	int64
4	BMI	1000 non-null	float64
5	Air_Pollution_Level	1000 non-null	int64
6	Respiratory_Infections_Childhood	1000 non-null	int64
7	COPD_Diagnosis	1000 non-null	int64
8	Pollution_Risk_Score	1000 non-null	int32
9	Smoking_Status_Encoded	1000 non-null	float64
10	Gender_Encoded	1000 non-null	int64
11	Smoking_Pollution_Interaction	1000 non-null	float64
12	Location_Biratnagar	1000 non-null	bool
13	Location_Butwal	1000 non-null	bool
14	Location_Chitwan	1000 non-null	bool
15	Location_Dharan	1000 non-null	bool
16	Location_Hetauda	1000 non-null	bool
17	Location_Kathmandu	1000 non-null	bool
18	Location_Lalitpur	1000 non-null	bool
19	Location_Nepalgunj	1000 non-null	bool
20	Location_Pokhara	1000 non-null	bool

dtypes: bool(9), float64(3), int32(1), int64(8)  
memory usage: 98.8 KB

## Step 6: Model Development

Then, this updated dataset was split into train and test data sets and then based on the data, COPD Diagnosis(the target variable) is a binary classification which means someone can have COPD(1) or cannot(0), so the following models were used after consideration.

- Logistic Regression
- Decision Trees
- Random Forest

After the datasets were model trained and saved, we evaluated them using accuracy score, precision score and F1 score to figure out the best model as shown below:

Logistic Regression Evaluation:					
	precision	recall	f1-score	support	
0	0.97	0.98	0.97	134	
1	0.95	0.94	0.95	66	
accuracy			0.96	200	
macro avg	0.96	0.96	0.96	200	
weighted avg	0.96	0.96	0.96	200	
Decision Tree Evaluation:					
	precision	recall	f1-score	support	
0	1.00	1.00	1.00	134	
1	1.00	1.00	1.00	66	
accuracy			1.00	200	
macro avg	1.00	1.00	1.00	200	
weighted avg	1.00	1.00	1.00	200	

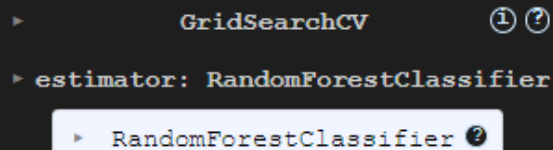
Random Forest Evaluation:					
	precision	recall	f1-score	support	
...					
accuracy			1.00	200	
macro avg	1.00	1.00	1.00	200	
weighted avg	1.00	1.00	1.00	200	

As we can see both “**Decision Trees and Random Forest**”, have a “1.00” score for accuracy, precision and F1 Score, so both can be nominated for best models. But, even if the score is “1.00” there are few things to consider such as “**Overfitting, Data Imbalance, Test Set Size, etc.**” before concluding it being perfect so any of the two can be chosen for refinement. In this case, **Random Forest** is chosen.

## Step 7: Model Tuning and Optimization

For random forest model refinement, we use GridSearchCV to create a refined model along with the best parameters and save it to a pickle file.

```
#Fit the Grid Search CV
grid_search_rf.fit(X_train, y_train)
```



```
GridSearchCV  ⓘ ⓘ
└─ estimator: RandomForestClassifier
   └─ RandomForestClassifier ⓘ
```

```
#Best Parameters
print(f"Best Parameteres: {grid_search_rf.best_params_}")
best_model = grid_search_rf.best_estimator_
```

```
Best Parameteres: {'max_depth': 10, 'min_samples_split': 2, 'n_estimators': 200}
```

```
#Save the model
with open('Best_Random_Forest_Model.pkl','wb') as f:
    pickle.dump(best_model, f)

print("Model refinement completed and best modelsaved!")
```

```
Model refinement completed and best modelsaved!
```

## Step 8: Model Deployment

After this, we used the recently refined and trained datasets to predict COPD. Then deploy the system it to a URL so that anyone will be able to access and use it for COPD prediction.

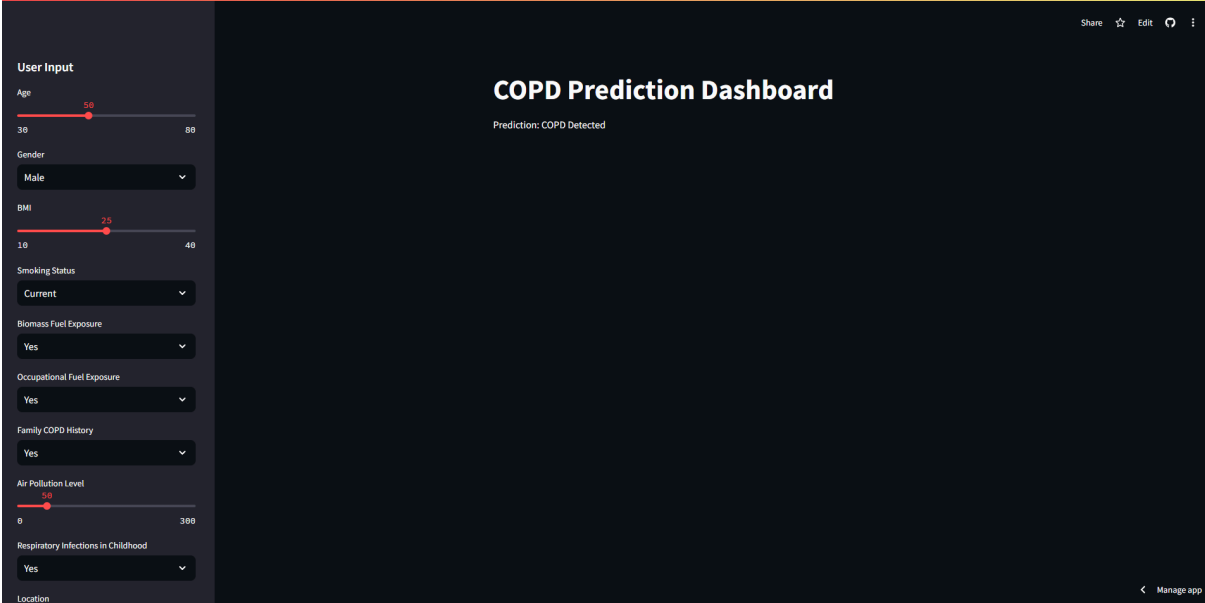
In the dashboard, various input fields are used to update the features which will predict the COPD and then display the output.

### Input Fields:

- Age
- Gender
- BMI
- Smoking Status
- Biomass Fuel Exposure
- Occupational Fuel Exposure
- Family COPD History
- Air Pollution Level
- Respiratory Infections in Childhood
- Location

We use the StreamLit Environment for deployment.

**Deployed URL:** <https://copdprediction-angela.streamlit.app/>



The screenshot displays the 'COPD Prediction Dashboard' interface. On the left, a sidebar titled 'User Input' contains various controls: a slider for 'Age' (range 30-80, value 50), a dropdown for 'Gender' (selected 'Male'), a slider for 'BMI' (range 10-40, value 25), dropdowns for 'Smoking Status' (selected 'Current'), 'Biomass Fuel Exposure' (selected 'Yes'), 'Occupational Fuel Exposure' (selected 'Yes'), 'Family COPD History' (selected 'Yes'), a slider for 'Air Pollution Level' (range 0-300, value 50), a dropdown for 'Respiratory Infections in Childhood' (selected 'Yes'), and a text input for 'Location'. The main panel on the right shows the title 'COPD Prediction Dashboard' and the prediction result 'Prediction: COPD Detected'. At the bottom right of the main panel, there is a link to 'Manage app'.