

## Data Collection and Preprocessing Phase

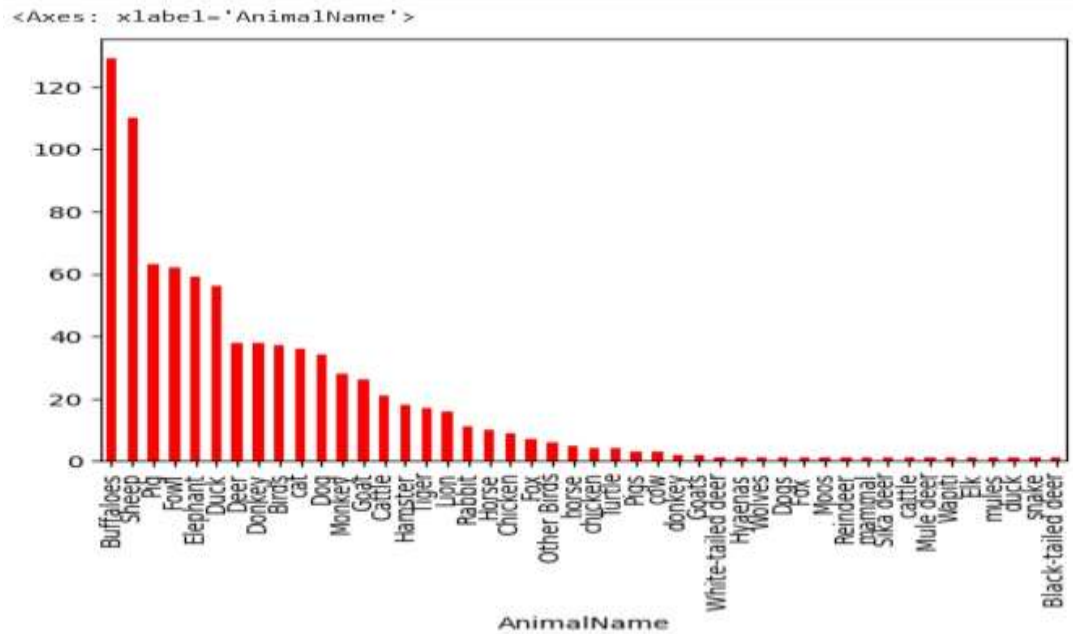
|               |  |
|---------------|--|
| Date          | 10 June 2024   |
| Team ID       | 740056   |
| Project Title | Beyond The Veil Of Wellness: Machine Learning's Unique Journey In Animal Health Classification |
| Maximum Marks | 6 Marks  |

## Data Exploration and Preprocessing Report

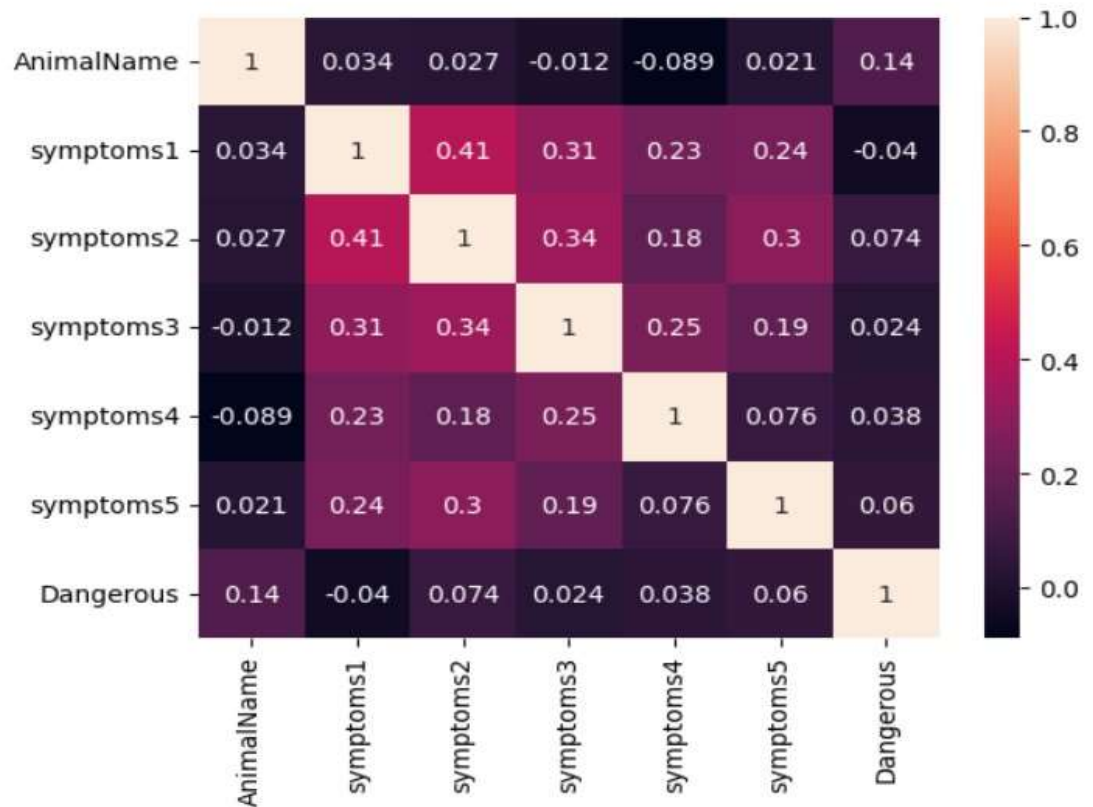
Dataset variables will be statistically analyzed to identify patterns and outliers, with Python employed for preprocessing tasks like normalization and feature engineering. Data cleaning will address missing values and outliers, ensuring quality for subsequent analysis and modeling, and forming a strong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

| Section       | Description  |           |       |          |          |             |       |
|---------------|--|-----------|-------|----------|----------|-------------|-------|
| Data Overview | <div> <div>AnimalName</div> <div>symptoms1</div> <div>symptoms2</div> <div>symptoms3</div> <div>symptoms4</div> <div>symptoms5</div> <div>Dangerous</div> </div> |           |       |          |          |             |       |
|               | count  | 871       | 871   | 871      | 871      | 871         | 869   |
|               | unique   | 46        | 232   | 230      | 229      | 217         | 203   |
|               | top  | Buffaloes | Fever | Diarrhea | Coughing | Weight loss | Pains |
|               | freq   | 129       | 257   | 119      | 95       | 117         | 99    |

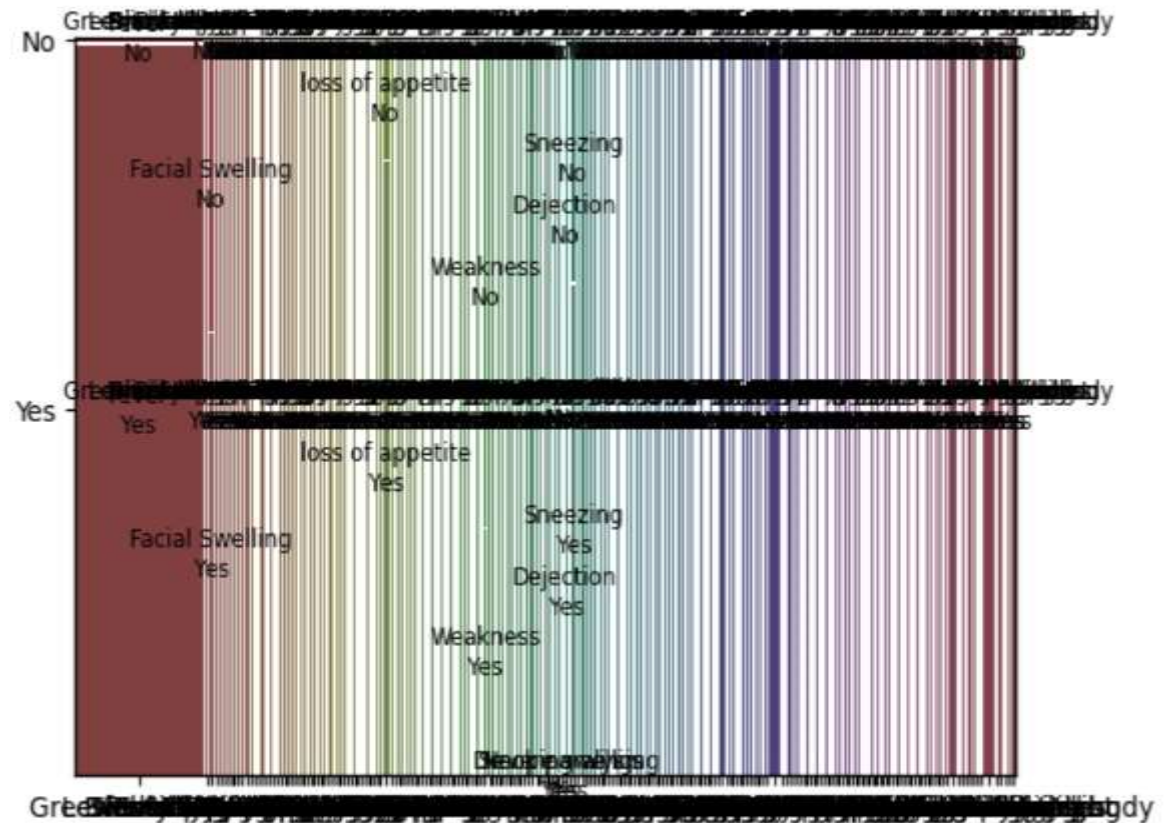
Univariate  
Analysis:



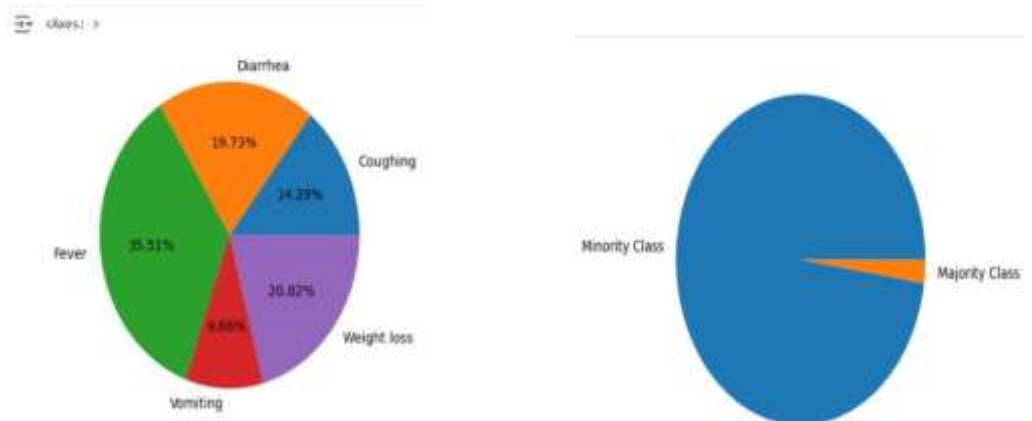
Bivariate  
Analysis



Multivariate  
Analysis



Pie chart



Data Preprocessing Code Screenshots :

## Loading Data

```
df=pd.read_csv("/content/data.csv")
df
```

|     | AnimalName | symptoms1                       | symptoms2            | symptoms3          | symptoms4            | symptoms5          | Dangerous |
|-----|------------|---------------------------------|----------------------|--------------------|----------------------|--------------------|-----------|
| 0   | Dog        | Fever                           | Diarrhea             | Vomiting           | Weight loss          | Dehydration        | Yes       |
| 1   | Dog        | Fever                           | Diarrhea             | Coughing           | Tiredness            | Pains              | Yes       |
| 2   | Dog        | Fever                           | Diarrhea             | Coughing           | Vomiting             | Anorexia           | Yes       |
| 3   | Dog        | Fever                           | Difficulty breathing | Coughing           | Lethargy             | Sneezing           | Yes       |
| 4   | Dog        | Fever                           | Diarrhea             | Coughing           | Lethargy             | Blue Eye           | Yes       |
| ... |            |                                 |                      |                    |                      |                    |           |
| 866 | Buffaloes  | Fever                           | Difficulty breathing | Poor Appetite      | Eye and Skin change  | Unable to exercise | Yes       |
| 867 | Buffaloes  | Fever                           | Loss of appetite     | Lesion on the skin | Lethargy             | Joint Pain         | Yes       |
| 868 | Buffaloes  | Lesions in the nasal cavity     | Lesions on nose      | Vomiting           | Noisy Breathing      | Lesions on nose    | Yes       |
| 869 | Buffaloes  | Hair loss                       | Dandruff             | Vomiting           | Crusting of the skin | Ulcerated skin     | Yes       |
| 870 | Buffaloes  | Greenish-yellow nasal discharge | Lack of pigmentation | Vomiting           | Lethargy             | Pain on face       | Yes       |

871 rows x 7 columns

## Handling Missing Data

```
df.isnull().sum()
```

```
AnimalName      0
symptoms1       0
symptoms2       0
symptoms3       0
symptoms4       0
symptoms5       0
Dangerous        2
dtype: int64
```

```
df['Dangerous'].unique()
```

```
array(['Yes', 'No', nan], dtype=object)
```

```
df['Dangerous'].value_counts()
```

```
Dangerous
Yes      849
No        20
Name: count, dtype: int64
```

```
df['Dangerous'].fillna('Yes',inplace=True)
```

```
df.isnull().sum()
```

```
AnimalName      0
symptoms1       0
symptoms2       0
symptoms3       0
symptoms4       0
symptoms5       0
Dangerous        0
dtype: int64
```

|                     |  |
|---------------------|--|
| Data Transformation | <pre> from sklearn.preprocessing import LabelEncoder  le = LabelEncoder() df['AnimalName'] = le.fit_transform(df['AnimalName']) df['symptoms1'] = le.fit_transform(df['symptoms1']) df['symptoms2'] = le.fit_transform(df['symptoms2']) df['symptoms3'] = le.fit_transform(df['symptoms3']) df['symptoms4'] = le.fit_transform(df['symptoms4']) df['symptoms5'] = le.fit_transform(df['symptoms5']) df['Dangerous'] = le.fit_transform(df['Dangerous']) </pre> |
| Feature Engineering | Attached the codes in final submission.  |
| Save Processed Data | --   |