

# 1. Important Information

## 2. Loading The Data

Import all of the necessary libraries:

```
In [1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import plotly.express as px
```

Load the data from the '.csv' file into a 'pandas dataframe'.

```
In [2]: data = pd.read_csv('RICE.csv')
print(data.info())
data.head()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 19404 entries, 0 to 19403
Data columns (total 14 columns):
 #   Column           Non-Null Count  Dtype  
---  -- 
 0   Observation Year      19404 non-null   int64  
 1   Standard Week     19404 non-null   int64  
 2   Pest Value        19404 non-null   float64 
 3   Collection Type  19404 non-null   object  
 4   MaxT              19404 non-null   float64 
 5   MinT              19404 non-null   float64 
 6   RH1(%)            19404 non-null   float64 
 7   RH2(%)            19404 non-null   float64 
 8   RF(mm)            19404 non-null   float64 
 9   WS(kmph)          19404 non-null   float64 
 10  SSH(hrs)          19404 non-null   float64 
 11  EVP(mm)           19404 non-null   float64 
 12  PEST NAME         19404 non-null   object  
 13  Location           19404 non-null   object  
dtypes: float64(9), int64(2), object(3)
memory usage: 2.1+ MB
None
```

	Observation Year	Standard Week	Pest Value	Collection Type	MaxT	MinT	RH1(%)	RH2(%)	RF(mm)
<b>0</b>	2003	1	0.0	Number/hill	27.9	14.8	94.7	51.3	0.0
<b>1</b>	2003	2	0.0	Number/hill	27.2	15.0	93.9	53.1	0.0
<b>2</b>	2003	3	0.0	Number/hill	28.7	18.3	94.1	56.7	0.6
<b>3</b>	2003	4	0.0	Number/hill	25.3	16.4	90.9	57.4	0.3
<b>4</b>	2003	5	0.0	Number/hill	28.8	18.7	95.7	55.0	0.0

## 3. Data Exploration

### 3.1 Attribute Analysis

#### Understanding Each Feature

- **Observation Year:** The fiscal year that the observation / reading was made.
- **Standard Week:** The week of the year that the observation / reading was made (Ranging from 1 to 52).
- **Pest Value:** The numerical value of the reading that was taken. This is used to create the target attribute.
- **Collection Type:** The process / procedure used to take the reading.
- **MaxT:** The maximum temperature during the respective week.
- **MinT:** The minimum temperature during the respective week.
- **RH1(%):** The maximum relative humidity during the respective week.
- **RH2(%):** The minimum relative humidity during the respective week.
- **RF(mm):** The rainfall (in mm) during the respective week.
- **WS(kmph):** The wind speed (in kmph) during the respective week.
- **SSH(hrs):** The average sunshine hours per day during the respective week.
- **EVP(mm):** The evaporation (in mm) during the respective week.
- **Pest Name:** The recorded pest's name.
- **Location:** The location where the reading was collected / recorded.

#### Coverage

- **Temporal Coverage Start Date:** 1960/12/31
- **Temporal Coverage End Date:** 2011/12/31
- **Geospatial Coverage:** India

In [3]: `data.describe()`

	<b>Observation Year</b>	<b>Standard Week</b>	<b>Pest Value</b>	<b>MaxT</b>	<b>MinT</b>	<b>RH</b>
<b>count</b>	19404.000000	19404.000000	19404.000000	19404.000000	19404.000000	19404.00
<b>mean</b>	2000.024789	26.473717	807.944081	31.169006	20.404540	82.19
<b>std</b>	9.827306	15.016247	5290.180315	4.904610	5.388381	13.84
<b>min</b>	1959.000000	1.000000	0.000000	10.900000	0.800000	9.30
<b>25%</b>	1996.000000	13.000000	0.000000	28.800000	17.500000	79.10
<b>50%</b>	2001.000000	26.000000	3.000000	30.900000	22.000000	87.30
<b>75%</b>	2007.000000	39.000000	92.000000	33.425000	24.400000	91.00
<b>max</b>	2011.000000	52.000000	311169.000000	71.600000	30.900000	100.00

## "Collection Type"

This shows that there are 5 different 'Collection Types' used to collect the 'Pest Value'.

There are no deviations or inconsistencies with the names, therefore no normalization is needed.

```
In [4]: print(data['Collection Type'].describe())
print('')
print(data['Collection Type'].unique())
print('')
print(data['Collection Type'].value_counts())
```

```
count          19404
unique           5
top    Number/Light trap
freq            16430
Name: Collection Type, dtype: object
```

```
['Number/hill' 'Number/Light trap' 'Percent Damage'
 'Number/Pheromone trap' 'Percentage']
```

```
Collection Type
Number/Light trap      16430
Percentage            2298
Number/Pheromone trap   520
Percent Damage        104
Number/hill             52
Name: count, dtype: int64
```

## "Pest Name"

```
In [5]: print(data['PEST NAME'].describe())
print('')
print(data['PEST NAME'].unique())
print('')
print(data['PEST NAME'].value_counts())
```

```

count          19404
unique           11
top      Yellowstemborer
freq            4333
Name: PEST NAME, dtype: object

['Brownplanthopper' 'Gallmidge' 'Greenleafhopper' 'LeafFolder'
 'Yellowstemborer' 'Caseworm' 'Miridbug' 'Whitebackedplanthopper'
 'ZigZagleafhopper' 'LeafBlast' 'NeckBlast']

PEST NAME
Yellowstemborer      4333
Gallmidge             3016
Greenleafhopper       2287
LeafBlast              2090
Brownplanthopper      1958
LeafFolder             1716
Whitebackedplanthopper 1248
Miridbug               1144
Caseworm                936
ZigZagleafhopper       468
NeckBlast                 208
Name: count, dtype: int64

```

## "Location"

```
In [6]: print(data['Location'].describe())
print('')
print(data['Location'].unique())
print('')
print(data['Location'].value_counts())
```

```

count          19404
unique           6
top      Maruteru
freq            7053
Name: Location, dtype: object

['Cuttack' 'Ludhiana' 'Maruteru' 'Palampur' 'Raipur' 'Rajendranagar']

Location
Maruteru      7053
Rajendranagar 5539
Raipur        2132
Ludhiana      1976
Cuttack        1456
Palampur       1248
Name: count, dtype: int64

```

## 3.2 Pest Name Analysis

### "Brownplanthopper"

```
In [7]: data_BrownPlanthopper = data[data['PEST NAME'] == 'Brownplanthopper']
print(data_BrownPlanthopper['Collection Type'].value_counts())
print('')
print(data_BrownPlanthopper['Location'].value_counts())
```

```
Collection Type
Number/Light trap    1906
Number/hill          52
Name: count, dtype: int64
```

```
Location
Maruteru            918
Rajendranagar       624
Ludhiana            312
Cuttack              52
Raipur              52
Name: count, dtype: int64
```

## "Gallmidge"

```
In [8]: data_Gallmidge = data[data['PEST NAME'] == 'Gallmidge']
print(data_Gallmidge['Collection Type'].value_counts())
print('')
print(data_Gallmidge['Location'].value_counts())
```

```
Collection Type
Number/Light trap    2912
Percent Damage        104
Name: count, dtype: int64
```

```
Location
Maruteru            884
Cuttack              832
Rajendranagar       728
Raipur              520
Ludhiana             52
Name: count, dtype: int64
```

## "Greenleafhopper"

```
In [9]: data_Greenleafhopper = data[data['PEST NAME'] == 'Greenleafhopper']
print(data_Greenleafhopper['Collection Type'].value_counts())
print('')
print(data_Greenleafhopper['Location'].value_counts())
```

```
Collection Type
Number/Light trap    2287
Name: count, dtype: int64
```

```
Location
Maruteru            831
Rajendranagar       676
Raipur              416
Ludhiana            312
Cuttack              52
Name: count, dtype: int64
```

## "LeafFolder"

```
In [10]: data_LeafFolder = data[data['PEST NAME'] == 'LeafFolder']
print(data_LeafFolder['Collection Type'].value_counts())
print('')
print(data_LeafFolder['Location'].value_counts())
```

```
Collection Type
Number/Light trap    1716
Name: count, dtype: int64
```

```
Location
Maruteru        884
Rajendranagar   312
Ludhiana        260
Raipur          208
Cuttack          52
Name: count, dtype: int64
```

## "Yellowstemborer"

```
In [11]: data_Yellowstemborer = data[data['PEST NAME'] == 'Yellowstemborer']
print(data_Yellowstemborer['Collection Type'].value_counts())
print('')
print(data_Yellowstemborer['Location'].value_counts())
```

```
Collection Type
Number/Light trap    3813
Number/Pheromone trap 520
Name: count, dtype: int64
```

```
Location
Rajendranagar   1629
Maruteru        936
Ludhiana        676
Raipur          624
Cuttack          468
Name: count, dtype: int64
```

## "Caseworm"

```
In [12]: data_Caseworm = data[data['PEST NAME'] == 'Caseworm']
print(data_Caseworm['Collection Type'].value_counts())
print('')
print(data_Caseworm['Location'].value_counts())
```

```
Collection Type
Number/Light trap    936
Name: count, dtype: int64
```

```
Location
Maruteru        468
Rajendranagar   260
Raipur          156
Ludhiana        52
Name: count, dtype: int64
```

## "Miridbug"

```
In [13]: data_Miridbug = data[data['PEST NAME'] == 'Miridbug']
print(data_Miridbug['Collection Type'].value_counts())
print('')
print(data_Miridbug['Location'].value_counts())
```

```
Collection Type
Number/Light trap    1144
Name: count, dtype: int64
```

```
Location
Maruteru        780
Rajendranagar   260
Ludhiana        52
Raipur          52
Name: count, dtype: int64
```

## "Whitebackedplanthopper"

```
In [14]: data_Whitebackedplanthopper = data[data['PEST NAME'] == 'Whitebackedplanthopper']
print(data_Whitebackedplanthopper['Collection Type'].value_counts())
print('')
print(data_Whitebackedplanthopper['Location'].value_counts())
```

```
Collection Type
Number/Light trap    1248
Name: count, dtype: int64
```

```
Location
Maruteru        884
Ludhiana        260
Raipur          104
Name: count, dtype: int64
```

## "ZigZagleafhopper"

```
In [15]: data_ZigZagleafhopper = data[data['PEST NAME'] == 'ZigZagleafhopper']
print(data_ZigZagleafhopper['Collection Type'].value_counts())
print('')
print(data_ZigZagleafhopper['Location'].value_counts())
```

```
Collection Type
Number/Light trap    468
Name: count, dtype: int64
```

```
Location
Maruteru        468
Name: count, dtype: int64
```

## "LeafBlast"

```
In [16]: data_LeafBlast = data[data['PEST NAME'] == 'LeafBlast']
print(data_LeafBlast['Collection Type'].value_counts())
print('')
print(data_LeafBlast['Location'].value_counts())
```

```
Collection Type
Percentage      2090
Name: count, dtype: int64
```

```
Location
Palampur         1092
Rajendranagar    998
Name: count, dtype: int64
```

## "NeckBlast"

```
In [17]: data_NeckBlast = data[data['PEST NAME'] == 'NeckBlast']
print(data_NeckBlast['Collection Type'].value_counts())
print('')
print(data_NeckBlast['Location'].value_counts())
```

Collection Type  
Percentage 208  
Name: count, dtype: int64

Location  
Palampur 156  
Rajendranagar 52  
Name: count, dtype: int64

## 3.3 Data Visualization

### Attribute Frequency

#### Bar Charts

```
In [18]: # Select the columns for bar plots
columns = ['Observation Year', 'MaxT', 'RH1(%)', 'WS(kmph)', 'PEST NAME']

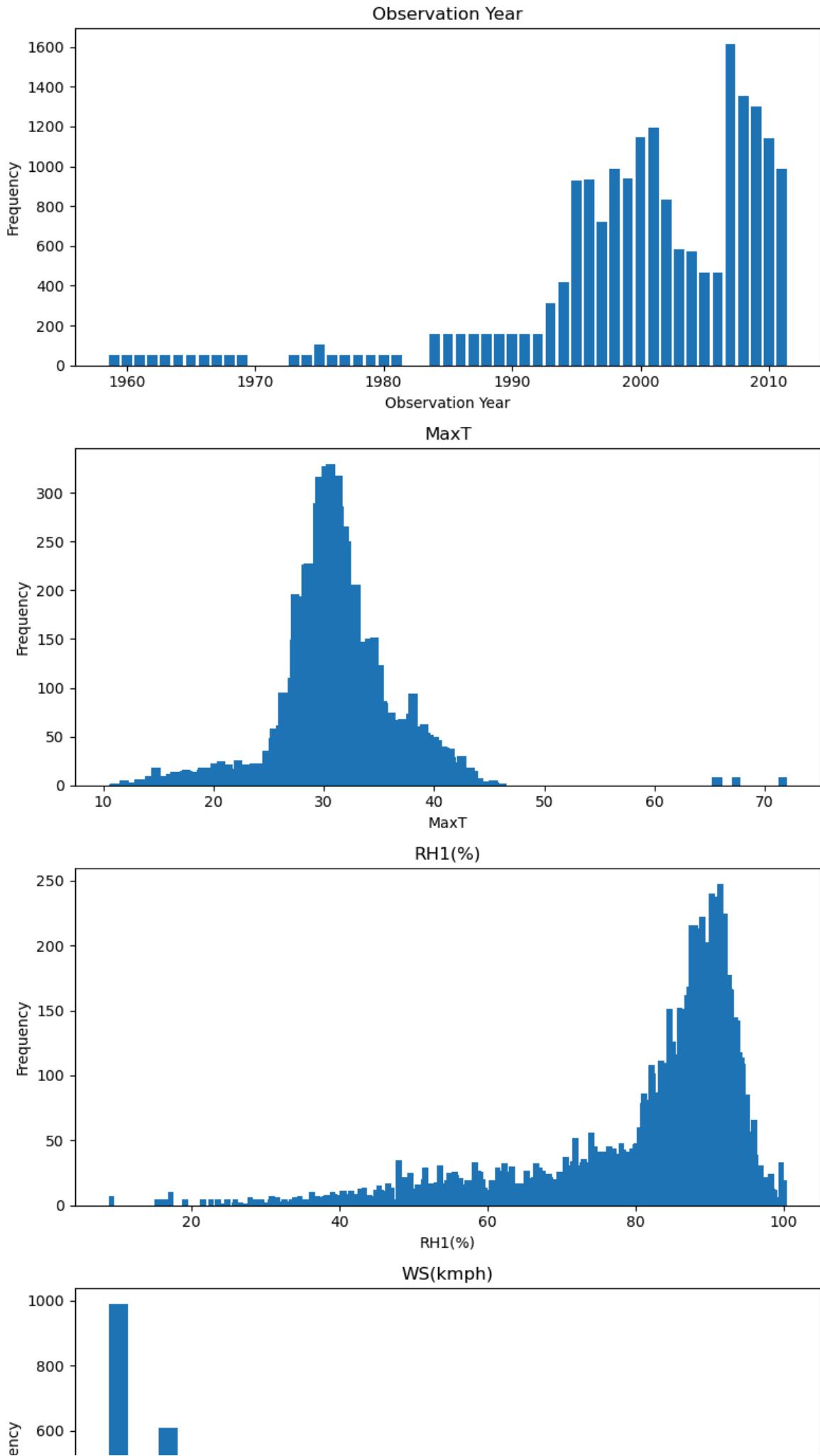
# Create subplots for bar plots
fig, axs = plt.subplots(len(columns), 1, figsize=(8, 4*len(columns)))

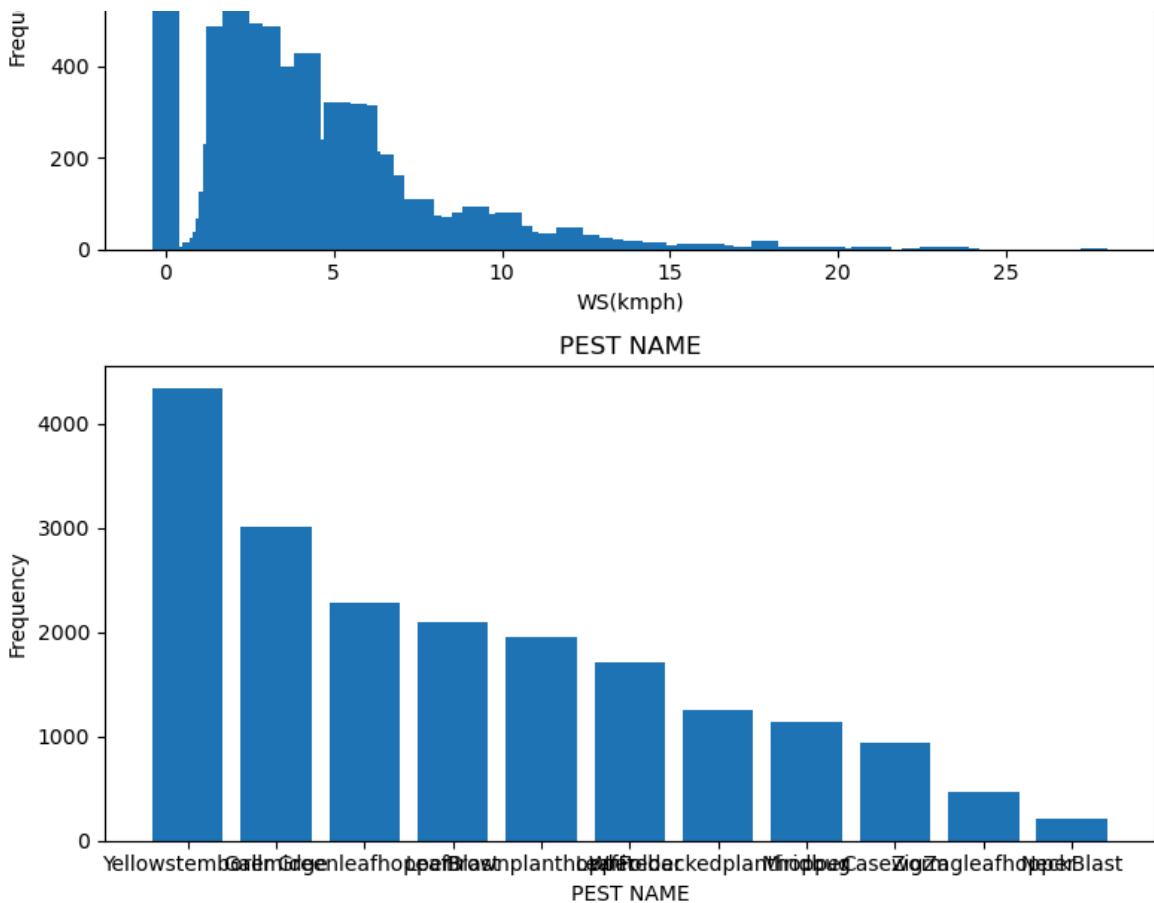
# Create bar plots for each column
for i, column in enumerate(columns):
    # Count the frequency of each unique value in the column
    value_counts = data[column].value_counts()

    # Plotting the bar graph
    axs[i].bar(value_counts.index, value_counts.values)
    axs[i].set_title(column)
    axs[i].set_xlabel(column)
    axs[i].set_ylabel("Frequency")

# Adjust the spacing between subplots
plt.tight_layout()

# Display the plots
plt.show()
```





## Pie Charts

```
In [19]: # Select the columns for pie charts
columns = ['Collection Type', 'PEST NAME', 'Location']

# Create subplots for pie charts (stacked vertically)
fig, axs = plt.subplots(len(columns), 1, figsize=(8, 4*len(columns)))

# Ensure axs is iterable for single subplot case
if len(columns) == 1:
    axs = [axs]

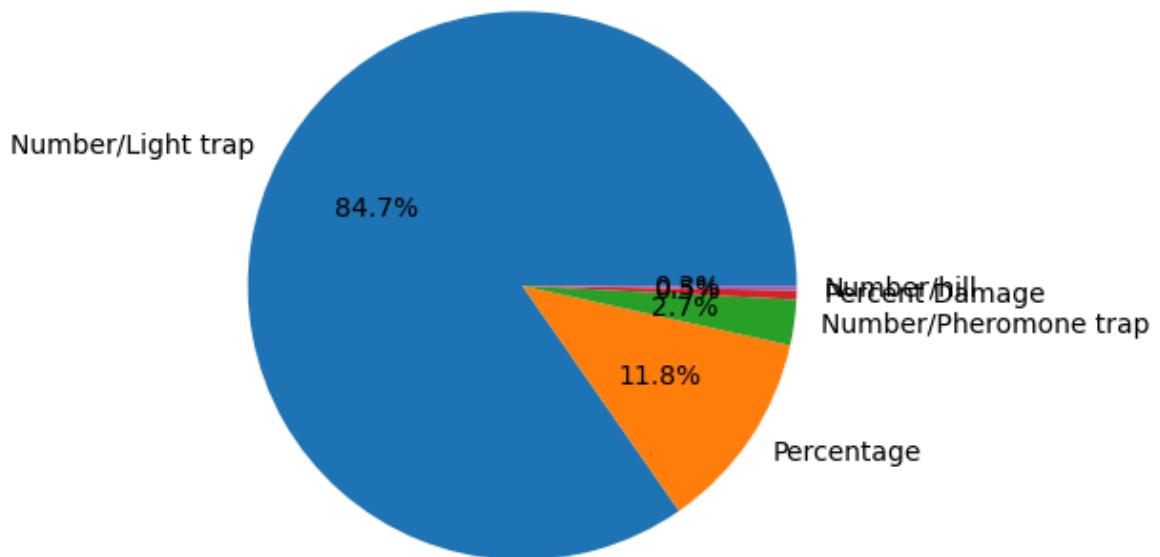
# Create pie charts for each column
for i, column in enumerate(columns):
    # Count the frequency of each unique value in the column
    value_counts = data[column].value_counts()

    # Plotting the pie chart
    axs[i].pie(value_counts, labels=value_counts.index, autopct='%1.1f%%')
    axs[i].set_title(column)

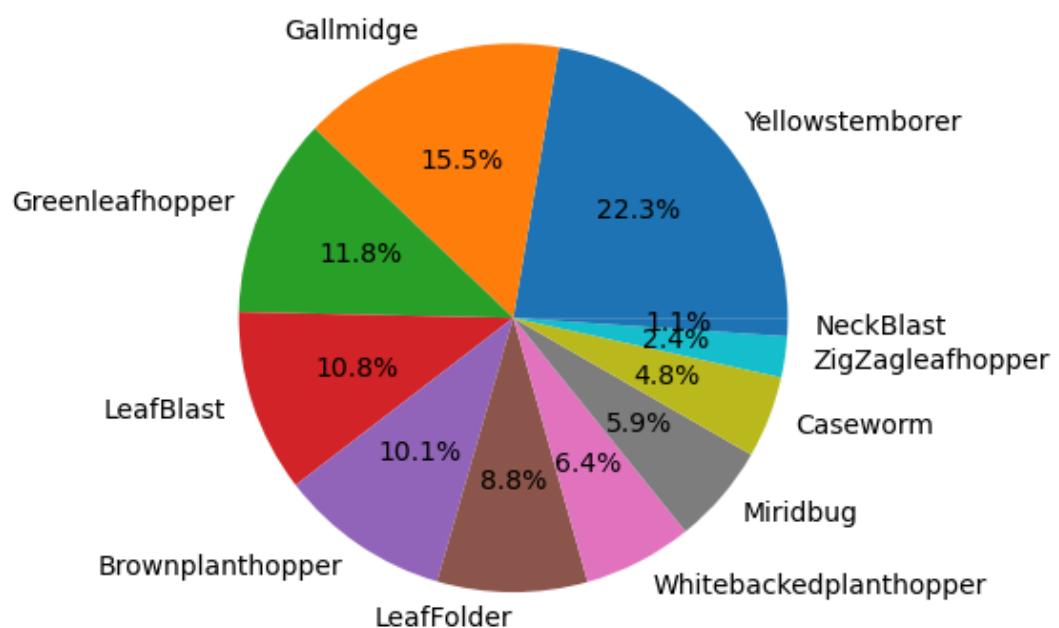
# Adjust the spacing between subplots
plt.tight_layout()

# Display the plots
plt.show()
```

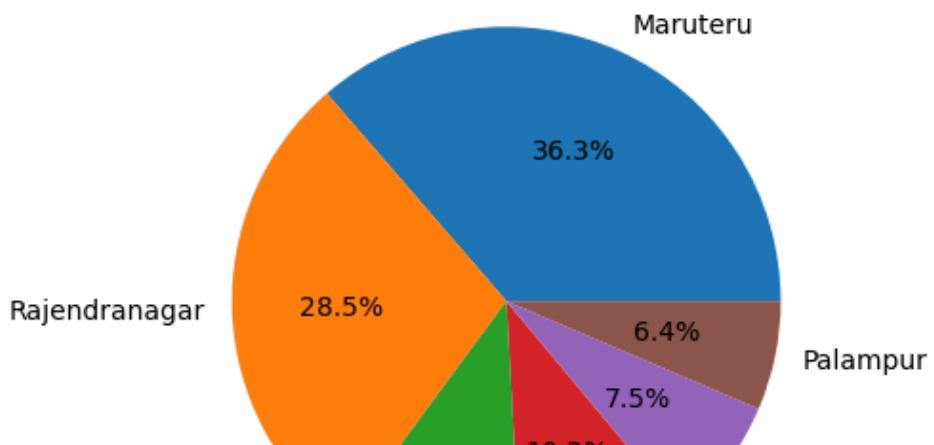
## Collection Type

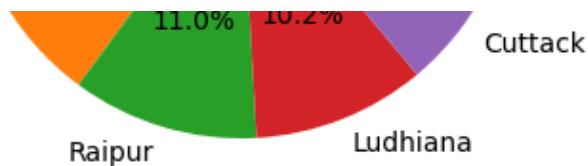


## PEST NAME



## Location





## 4. Data Preprocessing

### 4.1 Creating The Target Attribute

The "Outbreak" attribute is used as the target attribute for the Machine Learning Models for the "pest outbreak prediction", which is a typical Decision Support System (DSS). It is created with the following logic:

"If the 'Pest Value' is anything above '0', it is seen as an outbreak."

```
In [20]: data['Outbreak'] = (data['Pest Value'] > 0).astype(int)
data.head()
```

```
Out[20]:
```

	Observation Year	Standard Week	Pest Value	Collection Type	MaxT	MinT	RH1(%)	RH2(%)	RF(mm)
0	2003	1	0.0	Number/hill	27.9	14.8	94.7	51.3	0.0
1	2003	2	0.0	Number/hill	27.2	15.0	93.9	53.1	0.0
2	2003	3	0.0	Number/hill	28.7	18.3	94.1	56.7	0.6
3	2003	4	0.0	Number/hill	25.3	16.4	90.9	57.4	0.3
4	2003	5	0.0	Number/hill	28.8	18.7	95.7	55.0	0.0

### Total Outbreak Percentage

```
In [21]: print(data['Outbreak'].value_counts())
print('')
count_ones = data['Outbreak'].sum()
total = data['Outbreak'].count()
pct = count_ones / total * 100
print(f"Outbreak == 1: {count_ones}/{total} ({pct:.2f}%)")
```

```
Outbreak
1    10734
0     8670
Name: count, dtype: int64
```

```
Outbreak == 1: 10734/19404 (55.32%)
```

This represents a balanced dataset. Allowing the model to be trained equally with records that contain an outbreak and records that don't.

### Removing The 'Pest Value' Attribute

The 'Pest Value' attribute was used to create the target attribute called 'Outbreak', therefore it is no longer needed. Now it will be removed from the dataset.

```
In [22]: data.drop(columns=['Pest Value'], inplace=True)
data.head()
```

Out[22]:

	Observation Year	Standard Week	Collection Type	MaxT	MinT	RH1(%)	RH2(%)	RF(mm)	WS(km)
0	2003	1	Number/hill	27.9	14.8	94.7	51.3	0.0	
1	2003	2	Number/hill	27.2	15.0	93.9	53.1	0.0	
2	2003	3	Number/hill	28.7	18.3	94.1	56.7	0.6	
3	2003	4	Number/hill	25.3	16.4	90.9	57.4	0.3	
4	2003	5	Number/hill	28.8	18.7	95.7	55.0	0.0	

## Empty Values

This is to check if there are any empty values in the dataset. This shows that all of the columns and records are populated.

```
In [23]: data.isnull().sum()
```

Out[23]:

Observation	Year	0
Standard	Week	0
Collection	Type	0
MaxT		0
MinT		0
RH1(%)		0
RH2(%)		0
RF(mm)		0
WS(kmph)		0
SSH(hrs)		0
EVP(mm)		0
PEST NAME		0
Location		0
Outbreak		0

dtype: int64

## 4.2 Duplicate Records

This is to check if there are any duplicate records found in the dataset. They are then removed. This dataset doesn't contain any duplicates.

```
In [24]: print(data.duplicated().value_counts())
data = data.drop_duplicates()
```

```
False      19404
Name: count, dtype: int64
```

## 4.3 Handling Outliers

```
In [25]: # Plot / Visualize the outliers of the numerical features
for col in data[['MaxT', 'MinT', 'RH1(%)', 'RH2(%)', 'RF(mm)', 'WS(kmph)', 'SSH(%)']]:
    fig = px.box(
        data_frame=data,
        x=col,
        orientation='h',
        title=f'Boxplot of the Target ({col}) - With Outliers')
    )
fig.show()
```

```
In [26]: # Create a mask to filter out the outliers for 'LoanAmount'
# mask_MaxT = (data['MaxT'] >= 23.3) & (data['MaxT'] <= 38.7)

# # print(data[mask_MaxT].head())
# # print(data[mask_MaxT].info())

# fig = px.box(
#     data_frame=data[mask_MaxT],
#     x='MaxT',
#     orientation='h',
#     title='Boxplot of the Target (MaxT) - Without Outliers')

# fig.update_layout(xaxis_title='Target')
# fig.show()
```

## 4.4 Encode Categorical Variables

Use One-Hot encoding for 'PEST NAME', 'Location', and 'Collection Type'.

```
In [27]: # 1. Define categorical and numeric columns
cat_cols = ['Collection Type', 'PEST NAME', 'Location']
num_cols = [c for c in data.columns if c not in cat_cols + ['Outbreak']]

# 2. One-hot encode categorical features
data = pd.get_dummies(data, columns=cat_cols, drop_first=False)
```

# 5. Modelling

## 5.1 Splitting The Data

The data needs to be split into the training set and the testing set. 80% of the dataset will be assigned to the training set and 20% will be assigned to the testing set. The sets are assigned randomly to ensure integrity.

```
In [28]: # Splitting the dataset into the Training set and Test set
from sklearn.model_selection import train_test_split

X = data.drop(columns=['Outbreak'], inplace=False)
y = data['Outbreak']

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
```

## 5.2 Random Forest Model

A Random Forest Model is trained using the training set from above.

```
In [29]: from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import classification_report, roc_auc_score

# Random Forest model
rf_clf = RandomForestClassifier(
    n_estimators=300,
    max_depth=None,
    random_state=42,
    n_jobs=-1,
    class_weight=None # or 'balanced' if you use a stricter Outbreak threshold
)

rf_clf.fit(X_train, y_train)

y_pred = rf_clf.predict(X_test)
y_prob = rf_clf.predict_proba(X_test)[:, 1]

print("Random Forest classification report:")
print(classification_report(y_test, y_pred))
print("Random Forest ROC AUC:", roc_auc_score(y_test, y_prob))
```

	precision	recall	f1-score	support
0	0.88	0.81	0.84	1734
1	0.85	0.91	0.88	2147
accuracy			0.87	3881
macro avg	0.87	0.86	0.86	3881
weighted avg	0.87	0.87	0.86	3881

Random Forest ROC AUC: 0.9414335283964266

## 5.3 Gradient Boosting Model

A Gradient Boosting Model is trained using the training set from above.

```
In [30]: from xgboost import XGBClassifier

xgb_clf = XGBClassifier(
    n_estimators=400,
    learning_rate=0.05,
    max_depth=5,
    subsample=0.8,
    colsample_bytree=0.8,
    random_state=42,
    n_jobs=-1,
    eval_metric='logloss' # suppresses warning
    # scale_pos_weight can be tuned if classes become imbalanced
)

xgb_clf.fit(X_train, y_train)
```

```

y_pred_xgb = xgb_clf.predict(X_test)
y_prob_xgb = xgb_clf.predict_proba(X_test)[:, 1]

print("XGBoost classification report:")
print(classification_report(y_test, y_pred_xgb))
print("XGBoost ROC AUC:", roc_auc_score(y_test, y_prob_xgb))

```

	precision	recall	f1-score	support
0	0.89	0.81	0.85	1734
1	0.86	0.92	0.89	2147
accuracy			0.87	3881
macro avg	0.87	0.87	0.87	3881
weighted avg	0.87	0.87	0.87	3881

XGBoost ROC AUC: 0.9433640137333874

## 5.4 Making Predictions With The Models

This is to use the trained models on new raw records. When you want to predict for new raw rows (with original columns).

```

In [31]: # 1. Example new raw records (same columns as original data before encoding)
df_new_raw = pd.DataFrame([
    {
        "Observation Year": 2010,
        "Standard Week": 30,
        "Pest Value": 50.0, # this will be dropped, same as when we
        "Collection Type": "Number/Light trap",
        "MaxT": 32.5,
        "MinT": 24.0,
        "RH1(%)": 90.0,
        "RH2(%)": 60.0,
        "RF(mm)": 25.0,
        "WS(kmph)": 4.0,
        "SSH(hrs)": 7.0,
        "EVP(mm)": 4.5,
        "PEST NAME": "Brownplanthopper",
        "Location": "Cuttack"
    },
    {
        "Observation Year": 2005,
        "Standard Week": 10,
        "Pest Value": 0.0,
        "Collection Type": "Number/Light trap",
        "MaxT": 28.0,
        "MinT": 18.0,
        "RH1(%)": 80.0,
        "RH2(%)": 45.0,
        "RF(mm)": 0.0,
        "WS(kmph)": 3.0,
        "SSH(hrs)": 8.5,
        "EVP(mm)": 3.0,
        "PEST NAME": "Yellowstemborer",
        "Location": "Maruterau"
    }
])

```

```

    },
    {
        "Observation Year": 2008,
        "Standard Week": 38,
        "Pest Value": 200.0,
        "Collection Type": "Number/Light trap",
        "MaxT": 33.0,
        "MinT": 23.0,
        "RH1(%)": 92.0,
        "RH2(%)": 65.0,
        "RF(mm)": 40.0,
        "WS(kmph)": 6.0,
        "SSH(hrs)": 5.5,
        "EVP(mm)": 5.0,
        "PEST NAME": "LeafBlast",
        "Location": "Rajendranagar"
    }
])
# 2. Drop the "Pest Value" attribute
df_new_raw = df_new_raw.drop(columns=["Pest Value"])

# 3. Function from previous answer, adjusted to use global cat_cols
cat_cols = ['Collection Type', 'PEST NAME', 'Location'] # must match training
trained_feature_cols = X_train.columns # from the training step

def prepare_new_records(df_raw, trained_columns, cat_cols):
    # One-hot encode categorical as in training
    df_encoded = pd.get_dummies(df_raw, columns=cat_cols, drop_first=False)

    # Add missing dummy columns
    for col in trained_columns:
        if col not in df_encoded.columns:
            df_encoded[col] = 0

    # Drop extra columns not used in training
    df_encoded = df_encoded[trained_columns]

    return df_encoded

# 4. Prepare new data
X_new = prepare_new_records(df_new_raw, trained_feature_cols, cat_cols)

# 5. Predict with trained models
# Random Forest predictions
rf_preds_new = rf_clf.predict(X_new)
rf_probs_new = rf_clf.predict_proba(X_new)[:, 1]

print("New raw records:")
print(df_new_raw)
print("\nRandom Forest predicted Outbreak (0/1):", rf_preds_new)
print("Random Forest predicted probability of Outbreak:", rf_probs_new)

print("\n")

# XGBoost predictions
xgb_preds_new = xgb_clf.predict(X_new)
xgb_probs_new = xgb_clf.predict_proba(X_new)[:, 1]

```

```
print("XGBoost predicted Outbreak (0/1):", xgb_preds_new)
print("XGBoost predicted probability of Outbreak:", xgb_probs_new)
```

New raw records:

	Observation Year	Standard Week	Collection Type	MaxT	MinT	RH1(%)	\
0	2010	30	Number/Light trap	32.5	24.0	90.0	
1	2005	10	Number/Light trap	28.0	18.0	80.0	
2	2008	38	Number/Light trap	33.0	23.0	92.0	

	RH2(%)	RF(mm)	WS(kmph)	SSH(hrs)	EVP(mm)	PEST NAME	\
0	60.0	25.0	4.0	7.0	4.5	Brownplanthopper	
1	45.0	0.0	3.0	8.5	3.0	Yellowstemborer	
2	65.0	40.0	6.0	5.5	5.0	LeafBlast	

	Location
0	Cuttack
1	Maruteru
2	Rajendranagar

```
Random Forest predicted Outbreak (0/1): [0 1 1]
Random Forest predicted probability of Outbreak: [0.48      0.91333333 0.79
]
```

```
XGBoost predicted Outbreak (0/1): [0 1 1]
XGBoost predicted probability of Outbreak: [0.258903  0.96805286 0.91739213]
XGBoost predicted Outbreak (0/1): [0 1 1]
XGBoost predicted probability of Outbreak: [0.258903  0.96805286 0.91739213]
```

## 5.5 Long Short-Term Memory Model - Specific

The LSTM Model will be trained on one pest and one location. The following code focuses on Yellowstemborer in Rajendranagar. The pest (Yellowstemborer) was chosen, seeing as it is the pest with the most amount of records. The location (Rajendranagar) was chosen, because it was the location with the most amount of records for this pest.

### 1. Choose Features And Sort

```
In [39]: data = pd.read_csv('RICE.csv')
data['Outbreak'] = (data['Pest Value'] > 0).astype(int)

# Start from your cleaned 'data' where Pest Value is dropped and Outbreak is def
# data columns: Observation Year, Standard Week, Collection Type, MaxT, ..., PES

# 1. Filter to one pest and one location (simplest case)
pest = "Yellowstemborer"
loc = "Rajendranagar"

df = data[(data["PEST NAME"] == pest) & (data["Location"] == loc)].copy()
print(df.head())
print(df.info())

# 2. Sort by time
df = df.sort_values(by=["Observation Year", "Standard Week"]).reset_index(drop=True)

# 3. (Optional) Drop columns you don't want as inputs
# For a first LSTM, you might use only numeric weather + week + year as inputs
```

```

input_cols = [
    "Observation Year",
    "Standard Week",
    "MaxT",
    "MinT",
    "RH1(%)",
    "RH2(%)",
    "RF(mm)",
    "WS(kmph)",
    "SSH(hrs)",
    "EVP(mm)",
]
target_col = "Outbreak"

df_model = df[input_cols + [target_col]].copy()

```

	Observation Year	Standard Week	Pest Value	Collection Type	MaxT	\
17775	1975	1	11.0	Number/Light trap	31.4	
17776	1975	2	27.0	Number/Light trap	31.3	
17777	1975	3	52.0	Number/Light trap	30.4	
17778	1975	4	63.0	Number/Light trap	31.0	
17779	1975	5	0.0	Number/Light trap	30.2	

	MinT	RH1(%)	RH2(%)	RF(mm)	WS(kmph)	SSH(hrs)	EVP(mm)	\
17775	17.5	79.6	43.7	0.8	1.8	8.0	3.4	
17776	16.2	83.7	38.0	0.0	2.1	9.4	4.1	
17777	13.7	79.0	37.3	0.0	2.2	10.0	3.8	
17778	14.4	80.4	34.9	0.0	1.6	9.1	3.5	
17779	16.3	81.1	38.2	1.2	2.7	7.2	3.5	

	PEST NAME	Location	Outbreak
17775	Yellowstemborer	Rajendranagar	1
17776	Yellowstemborer	Rajendranagar	1
17777	Yellowstemborer	Rajendranagar	1
17778	Yellowstemborer	Rajendranagar	1
17779	Yellowstemborer	Rajendranagar	0

<class 'pandas.core.frame.DataFrame'>  
Index: 1629 entries, 17775 to 19403  
Data columns (total 15 columns):

#	Column	Non-Null Count	Dtype
0	Observation Year	1629	non-null int64
1	Standard Week	1629	non-null int64
2	Pest Value	1629	non-null float64
3	Collection Type	1629	non-null object
4	MaxT	1629	non-null float64
5	MinT	1629	non-null float64
6	RH1(%)	1629	non-null float64
7	RH2(%)	1629	non-null float64
8	RF(mm)	1629	non-null float64
9	WS(kmph)	1629	non-null float64
10	SSH(hrs)	1629	non-null float64
11	EVP(mm)	1629	non-null float64
12	PEST NAME	1629	non-null object
13	Location	1629	non-null object
14	Outbreak	1629	non-null int32

dtypes: float64(9), int32(1), int64(2), object(3)

memory usage: 197.3+ KB

None

## 2. Scale numeric features

Use MinMax or standard scaling. Here is MinMax to using scikit-learn. Save "scaler" for later to transform any future data identically.

```
In [40]: from sklearn.preprocessing import MinMaxScaler

scaler = MinMaxScaler()
df_model_scaled = df_model.copy()
df_model_scaled[input_cols] = scaler.fit_transform(df_model[input_cols])
df_model_scaled.head()
```

Out[40]:

	<b>Observation Year</b>	<b>Standard Week</b>	<b>MaxT</b>	<b>MinT</b>	<b>RH1(%)</b>	<b>RH2(%)</b>	<b>RF(mm)</b>	<b>WS(kmph)</b>
<b>0</b>	0.0	0.000000	0.396739	0.492958	0.744648	0.405759	0.003810	0.065217
<b>1</b>	0.0	0.019608	0.391304	0.431925	0.807339	0.331152	0.000000	0.076087
<b>2</b>	0.0	0.039216	0.342391	0.314554	0.735474	0.321990	0.000000	0.079710
<b>3</b>	0.0	0.058824	0.375000	0.347418	0.756881	0.290576	0.000000	0.057971
<b>4</b>	0.0	0.078431	0.331522	0.436620	0.767584	0.333770	0.005714	0.097826

## 3. Build supervised sequences (sliding windows)

Define a helper to turn a univariate time series with features into sequences of length T predicting the next week's Outbreak.

```
In [41]: def make_sequences(df_scaled, input_cols, target_col, window_size=4):
    """
    df_scaled: DataFrame with scaled inputs and target
    Returns X (num_samples, window_size, num_features), y (num_samples,)
    """
    X_list, y_list = [], []
    values = df_scaled[input_cols + [target_col]].values
    n_total = len(values)

    for i in range(n_total - window_size):
        window = values[i : i + window_size]
        target = values[i + window_size, -1] # Outbreak after the window
        X_list.append(window[:, :-1]) # all input features over the window
        y_list.append(target)

    X = np.array(X_list)
    y = np.array(y_list).astype(int)
    return X, y

window_size = 4 # e.g., use 4 weeks history
X, y = make_sequences(df_model_scaled, input_cols, target_col, window_size=windo
print("X shape:", X.shape) # (samples, time_steps=window_size, features=len(inp
print("y shape:", y.shape)
```

```
X shape: (1625, 4, 10)
y shape: (1625,)
```

## 4. Train/test split for sequences

Use a chronological split to respect time ordering (no shuffling).

```
In [42]: # Simple temporal split: first 80% for train, last 20% for test
n_samples = X.shape[0]
split_index = int(n_samples * 0.8)

X_train, X_test = X[:split_index], X[split_index:]
y_train, y_test = y[:split_index], y[split_index:]

print("Train samples:", X_train.shape[0], "Test samples:", X_test.shape[0])
```

Train samples: 1300 Test samples: 325

## 5. Define and train an LSTM model

Example with Keras (TensorFlow backend):

```
In [43]: import tensorflow as tf
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import LSTM, Dense, Dropout

n_timesteps = X_train.shape[1]
n_features = X_train.shape[2]

model = Sequential()
model.add(LSTM(64, input_shape=(n_timesteps, n_features), return_sequences=False))
model.add(Dropout(0.3))
model.add(Dense(32, activation="relu"))
model.add(Dense(1, activation="sigmoid")) # binary classification

model.compile(
    loss="binary_crossentropy",
    optimizer="adam",
    metrics=["accuracy"]
)

history = model.fit(
    X_train, y_train,
    epochs=30,
    batch_size=32,
    validation_split=0.2,
    shuffle=False # keep temporal order
)
```

Epoch 1/30

C:\Users\hroux\AppData\Roaming\Python\Python312\site-packages\keras\src\layers\rnn\rnn.py:199: UserWarning:

Do not pass an `input\_shape`/`input\_dim` argument to a layer. When using Sequential models, prefer using an `Input(shape)` object as the first layer in the model instead.

```
33/33 ━━━━━━━━ 1s 6ms/step - accuracy: 0.5144 - loss: 0.6918 - val_accuracy: 0.7192 - val_loss: 0.6615
Epoch 2/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.5683 - loss: 0.6829 - val_accuracy: 0.7192 - val_loss: 0.6380
Epoch 3/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.5673 - loss: 0.6812 - val_accuracy: 0.7192 - val_loss: 0.6248
Epoch 4/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.6087 - loss: 0.6711 - val_accuracy: 0.7192 - val_loss: 0.6035
Epoch 5/30
33/33 ━━━━━━━━ 0s 2ms/step - accuracy: 0.6231 - loss: 0.6622 - val_accuracy: 0.7231 - val_loss: 0.5798
Epoch 6/30
33/33 ━━━━━━━━ 0s 2ms/step - accuracy: 0.6442 - loss: 0.6459 - val_accuracy: 0.7308 - val_loss: 0.5549
Epoch 7/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.6500 - loss: 0.6331 - val_accuracy: 0.7423 - val_loss: 0.5268
Epoch 8/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.6740 - loss: 0.6132 - val_accuracy: 0.7692 - val_loss: 0.4993
Epoch 9/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.6981 - loss: 0.6025 - val_accuracy: 0.7769 - val_loss: 0.4741
Epoch 10/30
33/33 ━━━━━━━━ 0s 2ms/step - accuracy: 0.6827 - loss: 0.5999 - val_accuracy: 0.7769 - val_loss: 0.4728
Epoch 11/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.6981 - loss: 0.5886 - val_accuracy: 0.7885 - val_loss: 0.4595
Epoch 12/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.7010 - loss: 0.5847 - val_accuracy: 0.7923 - val_loss: 0.4581
Epoch 13/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.7067 - loss: 0.5777 - val_accuracy: 0.7923 - val_loss: 0.4526
Epoch 14/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.7106 - loss: 0.5749 - val_accuracy: 0.7923 - val_loss: 0.4530
Epoch 15/30
33/33 ━━━━━━━━ 0s 2ms/step - accuracy: 0.7125 - loss: 0.5654 - val_accuracy: 0.7923 - val_loss: 0.4570
Epoch 16/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.7106 - loss: 0.5642 - val_accuracy: 0.7923 - val_loss: 0.4532
Epoch 17/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.7154 - loss: 0.5617 - val_accuracy: 0.7923 - val_loss: 0.4539
Epoch 18/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.7115 - loss: 0.5605 - val_accuracy: 0.7885 - val_loss: 0.4563
Epoch 19/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.7154 - loss: 0.5570 - val_accuracy: 0.7962 - val_loss: 0.4614
Epoch 20/30
33/33 ━━━━━━━━ 0s 3ms/step - accuracy: 0.7260 - loss: 0.5518 - val_accuracy: 0.7885 - val_loss: 0.4549
Epoch 21/30
```

```
33/33 ━━━━━━━━ 0s 2ms/step - accuracy: 0.7240 - loss: 0.5548 - val_accuracy: 0.7885 - val_loss: 0.4634
Epoch 22/30
33/33 ━━━━━━━━ 0s 2ms/step - accuracy: 0.7202 - loss: 0.5520 - val_accuracy: 0.7885 - val_loss: 0.4587
Epoch 23/30
33/33 ━━━━━━ 0s 3ms/step - accuracy: 0.7308 - loss: 0.5499 - val_accuracy: 0.7962 - val_loss: 0.4534
Epoch 24/30
33/33 ━━━━━━ 0s 2ms/step - accuracy: 0.7288 - loss: 0.5390 - val_accuracy: 0.7885 - val_loss: 0.4627
Epoch 25/30
33/33 ━━━━━━ 0s 3ms/step - accuracy: 0.7231 - loss: 0.5499 - val_accuracy: 0.7885 - val_loss: 0.4554
Epoch 26/30
33/33 ━━━━━━ 0s 3ms/step - accuracy: 0.7356 - loss: 0.5399 - val_accuracy: 0.8000 - val_loss: 0.4507
Epoch 27/30
33/33 ━━━━━━ 0s 3ms/step - accuracy: 0.7413 - loss: 0.5400 - val_accuracy: 0.8000 - val_loss: 0.4423
Epoch 28/30
33/33 ━━━━━━ 0s 2ms/step - accuracy: 0.7327 - loss: 0.5414 - val_accuracy: 0.8000 - val_loss: 0.4487
Epoch 29/30
33/33 ━━━━━━ 0s 3ms/step - accuracy: 0.7413 - loss: 0.5387 - val_accuracy: 0.8038 - val_loss: 0.4503
Epoch 30/30
33/33 ━━━━━━ 0s 2ms/step - accuracy: 0.7442 - loss: 0.5283 - val_accuracy: 0.8038 - val_loss: 0.4489
```

## 6. Evaluate the LSTM

```
In [44]: from sklearn.metrics import classification_report, roc_auc_score

y_prob = model.predict(X_test).ravel()
y_pred = (y_prob >= 0.5).astype(int)

print("LSTM classification report:")
print(classification_report(y_test, y_pred))
print("LSTM ROC AUC:", roc_auc_score(y_test, y_prob))
```

```
11/11 ━━━━━━ 0s 8ms/step
LSTM classification report:
precision    recall    f1-score   support
          0       0.48      0.29      0.36       45
          1       0.89      0.95      0.92      280

accuracy                           0.86      325
macro avg       0.69      0.62      0.64      325
weighted avg     0.84      0.86      0.84      325
```

LSTM ROC AUC: 0.6388888888888888