Importing the Libraries

```
In [1]:
        # for manipulations
        import numpy as np
        import pandas as pd
        # for data visualizations
        import matplotlib.pyplot as plt
        import seaborn as sns
        plt.style.use('fivethirtyeight')
        # for interactivity
        import ipywidgets
        from ipywidgets import interact
        C:\Users\rbgir\anaconda3\lib\site-packages\numpy\ distributor init.py:30: Us
        erWarning: loaded more than 1 DLL from .libs:
        C:\Users\rbgir\anaconda3\lib\site-packages\numpy\.libs\libopenblas.WCDJNK7YV
        MPZQ2ME2ZZHJJRJ3JIKNDB7.gfortran-win_amd64.dll
        C:\Users\rbgir\anaconda3\lib\site-packages\numpy\.libs\libopenblas64 v0.3.2
        1-gcc_10_3_0.dll
          warnings.warn("loaded more than 1 DLL from .libs:"
        C:\Users\rbgir\anaconda3\lib\site-packages\scipy\__init__.py:138: UserWarnin
        g: A NumPy version >=1.16.5 and <1.23.0 is required for this version of SciP
        y (detected version 1.24.4)
          warnings.warn(f"A NumPy version >={np minversion} and <{np maxversion} is</pre>
        required for this version of "
        Reading the Dataset
```

```
In [2]: |# Lets read the dataset
        data = pd.read csv('Crop recommendation.csv')
        # lets check teh shape of the dataset
        print("Shape of the Dataset :", data.shape)
        Shape of the Dataset: (2200, 8)
In [3]: # Lets check the head of the dataset
        data.head()
```

Out[3]:

	N	Р	K	temperature	humidity	ph	rainfall	label
0	90	42	43	20.879744	82.002744	6.502985	202.935536	rice
1	85	58	41	21.770462	80.319644	7.038096	226.655537	rice
2	60	55	44	23.004459	82.320763	7.840207	263.964248	rice
3	74	35	40	26.491096	80.158363	6.980401	242.864034	rice
4	78	42	42	20.130175	81.604873	7.628473	262.717340	rice

```
N - ratio of Nitrogen content in soil
        P - ratio of Phosphorous content in soil
        K - ration of Potassium content in soil
        temperature - temperature in degree Celsius
        humidity - relative humidity in %
        ph - ph value of the soil
        rainfall - rainfall in mm
In [4]: | # lets check if there is any missing value present in the dataset
        data.isnull().sum()
Out[4]: N
                        0
        Ρ
                        0
        Κ
                        0
                        0
        temperature
        humidity
                        0
        ph
                        0
        rainfall
                        0
        label
                        0
        dtype: int64
In [5]: # lets check the Crops present in this Dataset
        data['label'].value_counts()
Out[5]: label
        rice
                        100
                        100
        maize
        jute
                        100
        cotton
                        100
        coconut
                        100
        papaya
                        100
                        100
        orange
        apple
                        100
        muskmelon
                        100
                        100
        watermelon
                        100
        grapes
        mango
                        100
        banana
                        100
                        100
        pomegranate
        lentil
                        100
        blackgram
                        100
        mungbean
                        100
        mothbeans
                        100
        pigeonpeas
                        100
        kidneybeans
                        100
        chickpea
                        100
        coffee
                        100
        Name: count, dtype: int64
```

Description for each of the columns in the Dataset

Descriptive Statistics

Average Tempature in Celsius : 25.62 Average Relative Humidity in % : 71.48 Average PH Value of the soil : 6.47 Average Rainfall in mm : 103.46

```
In [6]: # Lets check the Summary for all the crops

print("Average Ratio of Nitrogen in the Soil : {0:.2f}".format(data['N'].mean(print("Average Ratio of Phosphorous in the Soil : {0:.2f}".format(data['P'].meprint("Average Ratio of Potassium in the Soil : {0:.2f}".format(data['K'].meanprint("Average Tempature in Celsius : {0:.2f}".format(data['temperature'].meanprint("Average Relative Humidity in % : {0:.2f}".format(data['humidity'].mean(print("Average PH Value of the soil : {0:.2f}".format(data['ph'].mean()))

Average Ratio of Nitrogen in the Soil : 50.55

Average Ratio of Phosphorous in the Soil : 53.36

Average Ratio of Potassium in the Soil : 48.15
```

```
In [7]: # lets check the Summary Statistics for each of the Crops
       @interact
       def summary(crops = list(data['label'].value_counts().index)):
          x = data[data['label'] == crops]
          print("----")
          print("Statistics for Nitrogen")
          print("Minimum Nitrigen required :", x['N'].min())
          print("Average Nitrogen required :", x['N'].mean())
          print("Maximum Nitrogen required :", x['N'].max())
          print("----")
          print("Statistics for Phosphorous")
          print("Minimum Phosphorous required :", x['P'].min())
          print("Average Phosphorous required :", x['P'].mean())
          print("Maximum Phosphorous required :", x['P'].max())
          print("----")
          print("Statistics for Potassium")
          print("Minimum Potassium required :", x['K'].min())
          print("Average Potassium required :", x['K'].mean())
          print("Maximum Potassium required :", x['K'].max())
          print("----")
          print("Statistics for Temperature")
          print("Minimum Temperature required : {0:.2f}".format(x['temperature'].mir
          print("Average Temperature required : {0:.2f}".format(x['temperature'].mea
          print("Maximum Temperature required : {0:.2f}".format(x['temperature'].max
          print("----")
          print("Statistics for Humidity")
          print("Minimum Humidity required : {0:.2f}".format(x['humidity'].min()))
          print("Average Humidity required : {0:.2f}".format(x['humidity'].mean()))
          print("Maximum Humidity required : {0:.2f}".format(x['humidity'].max()))
          print("----")
          print("Statistics for PH")
          print("Minimum PH required : {0:.2f}".format(x['ph'].min()))
          print("Average PH required : {0:.2f}".format(x['ph'].mean()))
          print("Maximum PH required : {0:.2f}".format(x['ph'].max()))
          print("-----")
          print("Statistics for Rainfall")
          print("Minimum Rainfall required : {0:.2f}".format(x['rainfall'].min()))
          print("Average Rainfall required : {0:.2f}".format(x['rainfall'].mean()))
          print("Maximum Rainfall required : {0:.2f}".format(x['rainfall'].max()))
```

Statistics for Nitrogen Minimum Nitrigen required: 60 Average Nitrogen required: 77.76 Maximum Nitrogen required : 100 -----Statistics for Phosphorous Minimum Phosphorous required: 35 Average Phosphorous required: 48.44 Maximum Phosphorous required : 60 -----Statistics for Potassium Minimum Potassium required: 15 Average Potassium required: 19.79 Maximum Potassium required : 25 -----Statistics for Temperature Minimum Temperature required: 18.04 Average Temperature required : 22.39 Maximum Temperature required : 26.55 -----Statistics for Humidity Minimum Humidity required: 55.28 Average Humidity required : 65.09 Maximum Humidity required : 74.83 -----Statistics for PH Minimum PH required: 5.51 Average PH required : 6.25 Maximum PH required : 7.00

Statistics for Rainfall

Minimum Rainfall required : 60.65 Average Rainfall required : 84.77 Maximum Rainfall required : 109.75

In [8]: ## Lets compare the Average Requirement for each crops with average conditions @interact def compare(conditions = ['N','P','K','temperature','ph','humidity','rainfall' print("Average Value for", conditions,"is {0:.2f}".format(data[conditions] print("Rice : {0:.2f}".format(data[(data['label'] == 'rice')][conditions]. print("Black Grams : {0:.2f}".format(data[data['label'] == 'blackgram'][cd print("Banana : {0:.2f}".format(data[(data['label'] == 'banana')][condition print("Jute : {0:.2f}".format(data[data['label'] == 'jute'][conditions].me print("Coconut : {0:.2f}".format(data[(data['label'] == 'coconut')][condit print("Apple : {0:.2f}".format(data[data['label'] == 'apple'][conditions]. print("Papaya : {0:.2f}".format(data[(data['label'] == 'papaya')][condition print("Muskmelon : {0:.2f}".format(data[data['label'] == 'muskmelon'][cond print("Grapes : {0:.2f}".format(data[(data['label'] == 'grapes')][condition print("Watermelon : {0:.2f}".format(data[data['label'] == 'watermelon'][cd print("Kidney Beans: {0:.2f}".format(data[(data['label'] == 'kidneybeans') print("Mung Beans : {0:.2f}".format(data[data['label'] == 'mungbean'][cond print("Oranges : {0:.2f}".format(data[(data['label'] == 'orange')][conditi print("Chick Peas : {0:.2f}".format(data[data['label'] == 'chickpea'][cond print("Lentils : {0:.2f}".format(data[(data['label'] == 'lentil')][conditj print("Cotton : {0:.2f}".format(data[data['label'] == 'cotton'][conditions print("Maize : {0:.2f}".format(data[(data['label'] == 'maize')][conditions print("Moth Beans : {0:.2f}".format(data[data['label'] == 'mothbeans'][cor print("Pigeon Peas : {0:.2f}".format(data[(data['label'] == 'pigeonpeas')] print("Mango : {0:.2f}".format(data[data['label'] == 'mango'][conditions]. print("Pomegranate : {0:.2f}".format(data[(data['label'] == 'pomegranate') print("Coffee : {0:.2f}".format(data[data['label'] == 'coffee'][conditions

conditions N

Average Value for N is 50.55

Rice: 79.89

Black Grams : 40.02 Banana : 100.23 Jute: 78.40 Coconut : 21.98 Apple : 20.80 Papaya : 49.88 Muskmelon: 100.32 Grapes : 23.18 Watermelon: 99.42 Kidney Beans: 20.75 Mung Beans : 20.99 Oranges : 19.58 Chick Peas : 40.09 Lentils : 18.77 Cotton : 117.77 Maize : 77.76

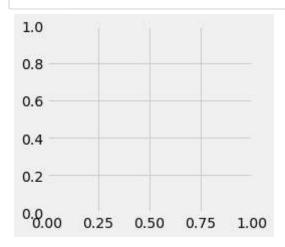
Moth Beans : 21.44 Pigeon Peas : 20.73 Mango : 20.07

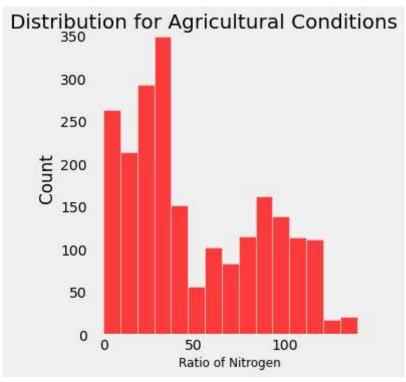
Pomegranate : 18.87 Coffee : 101.20

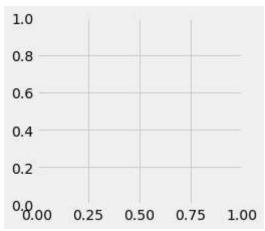
```
In [9]: # Lets make this funtion more Intuitive
       @interact
        def compare(conditions = ['N','P','K','temperature','ph','humidity','rainfall'
            print("Crops which require greater than average", conditions,'\n')
           print(data[data[conditions] > data[conditions].mean()]['label'].unique())
           print("----")
           print("Crops which require less than average", conditions,'\n')
           print(data[data[conditions] <= data[conditions].mean()]['label'].unique())</pre>
           conditions
                    Ν
        Crops which require greater than average N
         ['rice' 'maize' 'chickpea' 'blackgram' 'banana' 'watermelon' 'muskmelon'
          'papaya' 'cotton' 'jute' 'coffee']
        Crops which require less than average N
         ['chickpea' 'kidneybeans' 'pigeonpeas' 'mothbeans' 'mungbean' 'blackgram'
          'lentil' 'pomegranate' 'mango' 'grapes' 'apple' 'orange' 'papaya'
          'coconut']
```

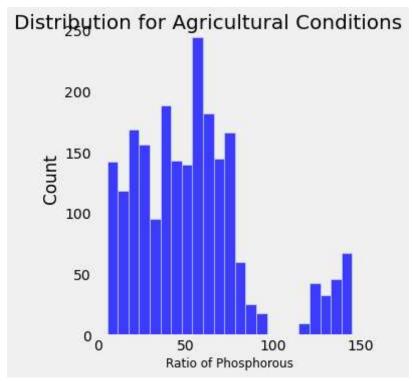
Analyzing Agricultural Conditions

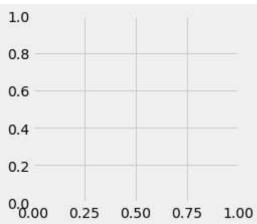
```
In [10]: import numpy as np
         import seaborn as sns
         import matplotlib.pyplot as plt
         plt.rcParams['figure.figsize'] = (15, 7)
         plt.subplot(2, 4, 1)
         sns.displot(data['N'].values, color='red') # Convert to numpy array with .val
         plt.xlabel('Ratio of Nitrogen', fontsize=12)
         plt.grid()
         plt.suptitle('Distribution for Agricultural Conditions', fontsize=20)
         plt.show()
         plt.subplot(2, 4, 2)
         sns.displot(data['P'].values, color='blue')
         plt.xlabel('Ratio of Phosphorous', fontsize=12)
         plt.grid()
         plt.suptitle('Distribution for Agricultural Conditions', fontsize=20)
         plt.show()
         plt.subplot(2, 4, 3)
         sns.displot(data['K'].values, color='darkblue')
         plt.xlabel('Ratio of Potassium', fontsize=12)
         plt.grid()
         plt.suptitle('Distribution for Agricultural Conditions', fontsize=20)
         plt.show()
         plt.subplot(2, 4, 4)
         sns.displot(data['temperature'].values, color='black')
         plt.xlabel('Temperature', fontsize=12)
         plt.grid()
         plt.suptitle('Distribution for Agricultural Conditions', fontsize=20)
         plt.show()
         plt.subplot(2, 4, 5)
         sns.displot(data['rainfall'].values, color='purple')
         plt.xlabel('Rainfall', fontsize=12)
         plt.suptitle('Distribution for Agricultural Conditions', fontsize=20)
         plt.show()
         plt.subplot(2, 4, 6)
         sns.displot(data['humidity'].values, color='lightgreen')
         plt.xlabel('Humidity', fontsize=12)
         plt.grid()
         plt.suptitle('Distribution for Agricultural Conditions', fontsize=20)
         plt.show()
         plt.subplot(2, 4, 7)
         sns.displot(data['ph'].values, color='darkgreen')
         plt.xlabel('pH Level', fontsize=12)
         plt.grid()
         plt.suptitle('Distribution for Agricultural Conditions', fontsize=20)
         plt.show()
```

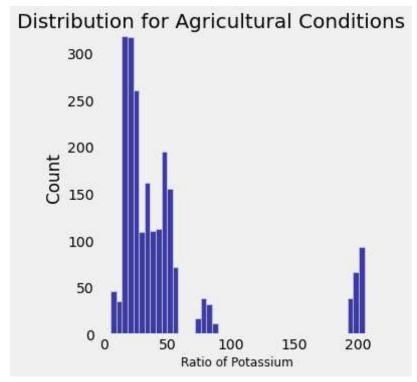


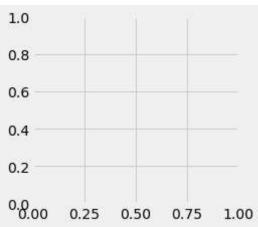


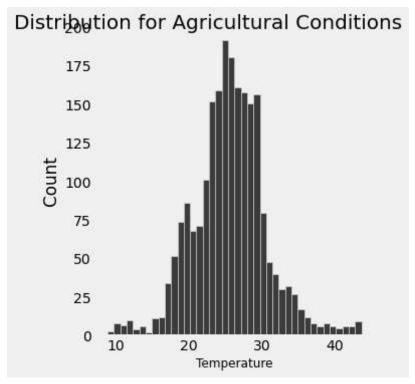


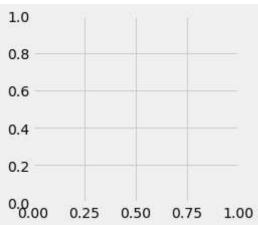


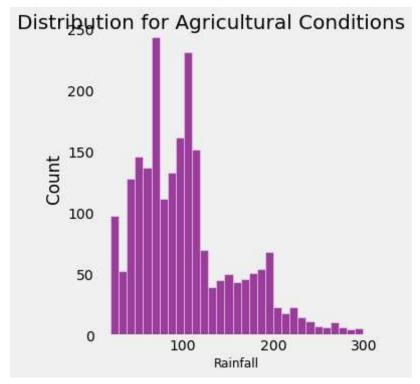


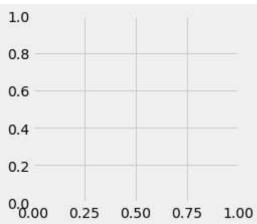


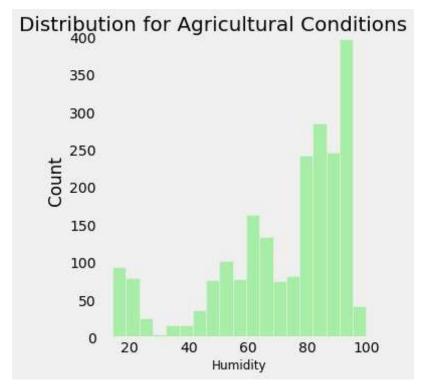


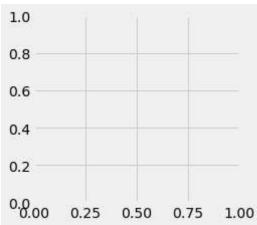


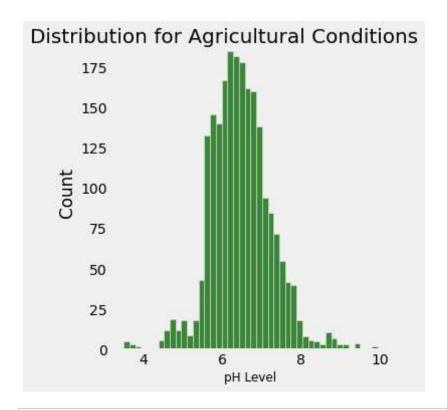












Clustering Similar Crops

```
In [13]: ### Lets try to Cluster these Crops

# Lets import the warnings library so that we can avoid warnings
import warnings
warnings.filterwarnings('ignore')

# Lets select the Spending score, and Annual Income Columns from the Data
x = data.loc[:, ['N','P','K','temperature','ph','humidity','rainfall']].values

# Let's check the shape of x
print(x.shape)

# Lets convert this data into a dataframe
x_data = pd.DataFrame(x)
x_data.head()
```

Out[13]:

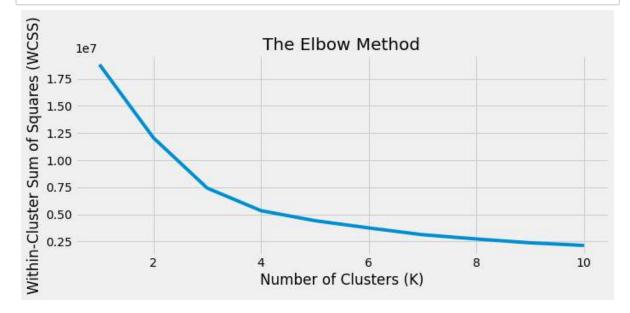
(2200, 7)

	0	1	2	3	4	5	6
0	90.0	42.0	43.0	20.879744	6.502985	82.002744	202.935536
1	85.0	58.0	41.0	21.770462	7.038096	80.319644	226.655537
2	60.0	55.0	44.0	23.004459	7.840207	82.320763	263.964248
3	74.0	35.0	40.0	26.491096	6.980401	80.158363	242.864034
4	78.0	42.0	42.0	20.130175	7.628473	81.604873	262.717340

```
In [14]: !pip install --upgrade threadpoolctl
```

Requirement already satisfied: threadpoolctl in c:\users\rbgir\anaconda3\lib\site-packages (3.2.0)

```
In [16]:
         from sklearn.cluster import KMeans
         import matplotlib.pyplot as plt
         plt.rcParams['figure.figsize'] = (10, 4)
         wcss = [] # List to store the within-cluster sum of squares (WCSS) for differ
         for i in range(1, 11):
             # Create a KMeans instance with i clusters, 'k-means++' initialization, an
             km = KMeans(n_clusters=i, init='k-means++', max_iter=300, n_init=10, rando
             # Fit the KMeans model to the data and calculate the WCSS
             km.fit(x) # Assuming 'x' is your dataset
             wcss.append(km.inertia_)
         # Plot the results
         plt.plot(range(1, 11), wcss)
         plt.title('The Elbow Method', fontsize=20)
         plt.xlabel('Number of Clusters (K)')
         plt.ylabel('Within-Cluster Sum of Squares (WCSS)')
         plt.show()
```



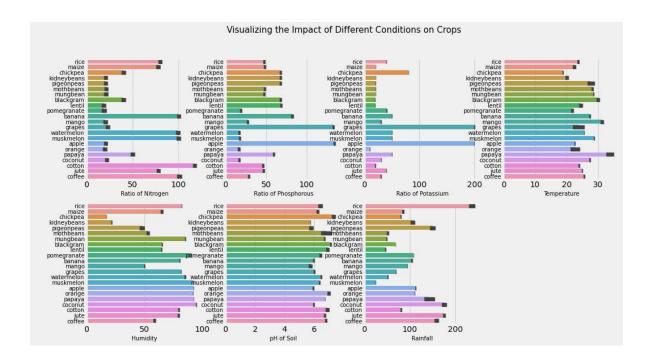
```
In [17]: # lets implement the K Means algorithm to perform Clustering analysis
       km = KMeans(n clusters = 4, init = 'k-means++', max iter = 300, n init = 10, r
       y_means = km.fit_predict(x)
       # Lets find out the Results
       a = data['label']
       y_means = pd.DataFrame(y_means)
       z = pd.concat([y means, a], axis = 1)
       z = z.rename(columns = {0: 'cluster'})
       # lets check the Clusters of each Crops
       print("Lets check the Results After Applying the K Means Clustering Analysis
       print("Crops in First Cluster:", z[z['cluster'] == 0]['label'].unique())
       print("-----")
       print("Crops in Second Cluster:", z[z['cluster'] == 1]['label'].unique())
       print("----")
       print("Crops in Third Cluster:", z[z['cluster'] == 2]['label'].unique())
       print("----")
       print("Crops in Forth Cluster:", z[z['cluster'] == 3]['label'].unique())
       Lets check the Results After Applying the K Means Clustering Analysis
       Crops in First Cluster: ['maize' 'chickpea' 'kidneybeans' 'pigeonpeas' 'moth
       beans' 'mungbean'
        'blackgram' 'lentil' 'pomegranate' 'mango' 'orange' 'papaya' 'coconut']
       _____
       Crops in Second Cluster: ['maize' 'banana' 'watermelon' 'muskmelon' 'papaya'
       'cotton' 'coffee']
       Crops in Third Cluster: ['grapes' 'apple']
       -----
       Crops in Forth Cluster: ['rice' 'pigeonpeas' 'papaya' 'coconut' 'jute' 'coff
```

ee']

```
In [18]: # Hard Clustering
        print("Results for Hard Clustering\n")
        counts = z[z['cluster'] == 0]['label'].value_counts()
        d = z.loc[z['label'].isin(counts.index[counts >= 50])]
        d = d['label'].value counts()
        print("Crops in Cluster 1:", list(d.index))
        print("-----")
        counts = z[z['cluster'] == 1]['label'].value_counts()
        d = z.loc[z['label'].isin(counts.index[counts >= 50])]
        d = d['label'].value counts()
        print("Crops in Cluster 2:", list(d.index))
        print("-----")
        counts = z[z['cluster'] == 2]['label'].value_counts()
        d = z.loc[z['label'].isin(counts.index[counts >= 50])]
        d = d['label'].value_counts()
        print("Crops in Cluster 3:", list(d.index))
        print("-----")
        counts = z[z['cluster'] == 3]['label'].value_counts()
        d = z.loc[z['label'].isin(counts.index[counts >= 50])]
        d = d['label'].value counts()
        print("Crops in Cluster 4:", list(d.index))
        Results for Hard Clustering
        Crops in Cluster 1: ['chickpea', 'kidneybeans', 'mothbeans', 'mungbean', 'bl
```

visualizing the Hidden Patterns

```
In [19]: ### Data Visualizations
         plt.rcParams['figure.figsize'] = (15, 8)
         plt.subplot(2, 4, 1)
         sns.barplot(data['N'], data['label'])
         plt.ylabel(' ')
         plt.xlabel('Ratio of Nitrogen', fontsize = 10)
         plt.yticks(fontsize = 10)
         plt.subplot(2, 4, 2)
         sns.barplot(data['P'], data['label'])
         plt.ylabel(' ')
         plt.xlabel('Ratio of Phosphorous', fontsize = 10)
         plt.yticks(fontsize = 10)
         plt.subplot(2, 4, 3)
         sns.barplot(data['K'], data['label'])
         plt.ylabel(' ')
         plt.xlabel('Ratio of Potassium', fontsize = 10)
         plt.yticks(fontsize = 10)
         plt.subplot(2, 4, 4)
         sns.barplot(data['temperature'], data['label'])
         plt.ylabel(' ')
         plt.xlabel('Temperature', fontsize = 10)
         plt.yticks(fontsize = 10)
         plt.subplot(2, 4, 5)
         sns.barplot(data['humidity'], data['label'])
         plt.ylabel(' ')
         plt.xlabel('Humidity', fontsize = 10)
         plt.yticks(fontsize = 10)
         plt.subplot(2, 4, 6)
         sns.barplot(data['ph'], data['label'])
         plt.ylabel(' ')
         plt.xlabel('pH of Soil', fontsize = 10)
         plt.yticks(fontsize = 10)
         plt.subplot(2, 4, 7)
         sns.barplot(data['rainfall'], data['label'])
         plt.ylabel(' ')
         plt.xlabel('Rainfall', fontsize = 10)
         plt.yticks(fontsize = 10)
         plt.suptitle('Visualizing the Impact of Different Conditions on Crops', fontsi
         plt.show()
```



Predictive Modelling

```
In [21]: # Lets split the Dataset for Predictive Modelling
         y = data['label']
         x = data.drop(['label'], axis = 1)
         print("Shape of x:", x.shape)
         print("Shape of y:", y.shape)
         Shape of x: (2200, 7)
         Shape of y: (2200,)
In [22]: # lets create Training and Testing Sets for Validation of Results
         from sklearn.model_selection import train_test_split
         x_train, x_test, y_train, y_test = train_test_split(x, y, test_size = 0.2, rar
         print("The Shape of x train:", x_train.shape)
         print("The Shape of x test:", x_test.shape)
         print("The Shape of y train:", y_train.shape)
         print("The Shape of y test:", y_test.shape)
         The Shape of x train: (1760, 7)
         The Shape of x test: (440, 7)
         The Shape of y train: (1760,)
         The Shape of y test: (440,)
```

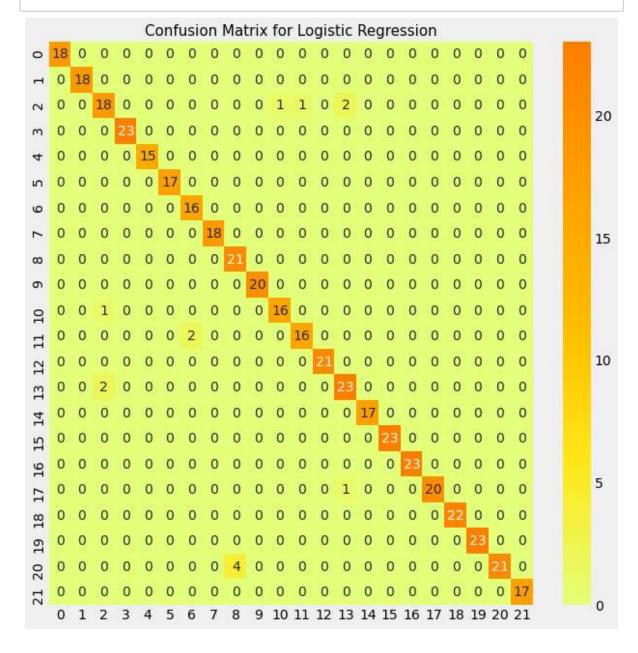
```
In [23]: # Lets create a Predictive Model
from sklearn.linear_model import LogisticRegression

model = LogisticRegression()
model.fit(x_train, y_train)
y_pred = model.predict(x_test)
```

```
In [24]: # lets evaluate the Model Performance
    from sklearn.metrics import classification_report, confusion_matrix

# lets print the Confusion matrix first
    plt.rcParams['figure.figsize'] = (10, 10)
    cm = confusion_matrix(y_test, y_pred)
    sns.heatmap(cm, annot = True, cmap = 'Wistia')
    plt.title('Confusion Matrix for Logistic Regression', fontsize = 15)
    plt.show()

# lets print the Classification Report also
    cr = classification_report(y_test, y_pred)
    print(cr)
```



precision	recall	f1-score	support
1.00	1.00	1.00	18
1.00	1.00	1.00	18
0.86	0.82	0.84	22
1.00	1.00	1.00	23
1.00	1.00	1.00	15
1.00	1.00	1.00	17
0.89	1.00	0.94	16
1.00	1.00	1.00	18
0.84	1.00	0.91	21
1.00	1.00	1.00	20
0.94	0.94	0.94	17
0.94	0.89	0.91	18
1.00	1.00	1.00	21
0.88	0.92	0.90	25
1.00	1.00	1.00	17
1.00	1.00	1.00	23
1.00	1.00	1.00	23
1.00	0.95	0.98	21
1.00	1.00	1.00	22
1.00	1.00	1.00	23
1.00	0.84	0.91	25
1.00	1.00	1.00	17
		0.97	440
0.97	0.97	0.97	440
0.97	0.97	0.97	440
	1.00 1.00 0.86 1.00 1.00 1.00 0.89 1.00 0.84 1.00 0.94 0.94 1.00 1.00 1.00 1.00 1.00	1.00	1.00

Calculate Accuracy for Logistic Regression in this model:

```
In [25]: from sklearn.metrics import accuracy_score
    from sklearn.linear_model import LogisticRegression # Assuming you're using I

# Create and train your model
    model = LogisticRegression() # Replace with your actual model initialization
    model.fit(x_train, y_train)

# Make predictions on the test set
    y_pred = model.predict(x_test)

# Calculate accuracy
    accuracy = accuracy_score(y_test, y_pred)

print(f'Accuracy: {accuracy:.2f}')
```

Accuracy: 0.97

It has 97 % Accuracy in Crop prediction.

Real time Predictions