

INTRODUCTION TO MACHINE LEARNING

(CSE-3008)

CROP RECOMMENDATION (WEB APPLICATION)

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Colab link :

[https://colab.research.google.com/drive/1Sti1YhGgSuOUs8PFFwwJ7DRFNWffM7Qd?
usp=sharing](https://colab.research.google.com/drive/1Sti1YhGgSuOUs8PFFwwJ7DRFNWffM7Qd?usp=sharing)

```
from google.colab import drive  
drive.mount('/content/gdrive', force_remount=True)
```

Mounted at /content/gdrive

```
import pandas as pd  
import io  
import numpy as np  
import random  
from sklearn.model_selection import cross_val_score  
  
import matplotlib.pyplot as plt  
import seaborn as sns  
  
import plotly.graph_objects as go  
import plotly.express as px  
from plotly.subplots import make_subplots  
import seaborn as sns  
from sklearn.metrics import classification_report  
from sklearn import metrics  
from sklearn import tree  
import warnings  
from sklearn.metrics import confusion_matrix  
warnings.filterwarnings('ignore')
```

```
df = pd.read_csv('/content/gdrive/MyDrive/ML Lab/Crop.csv')  
df.head()
```

	n	p	k	temperature	humidity	ph	rainfall	crop	season	area	pr
0	90	42	43	20.879744	82.002744	6.502985	202.935536	rice	winter	99	
1	85	58	41	21.770462	80.319644	7.038096	226.655537	rice	whole year	277	
2	60	55	44	23.004459	82.320763	7.840207	263.964248	rice	autumn	149	
3	74	35	40	26.491096	80.158363	6.980401	242.864034	rice	kharif	163	
4	78	42	42	20.130175	81.604873	7.628473	262.717340	rice	autumn	295	

```
df.shape
```

(2200, 11)

```
df.columns
```

```
Index(['n', 'p', 'k', 'temperature', 'humidity', 'ph', 'rainfall', 'crop',
       'season', 'area', 'production'],
      dtype='object')
```

```
df.isnull().any()
```

```
n      False
p      False
k      False
temperature  False
humidity  False
ph      False
rainfall  False
crop     False
season   False
area    False
production  False
dtype: bool
```

```
print("Number of various crops: ", len(df['crop'].unique()))
print("List of crops: ", df['crop'].unique())
```

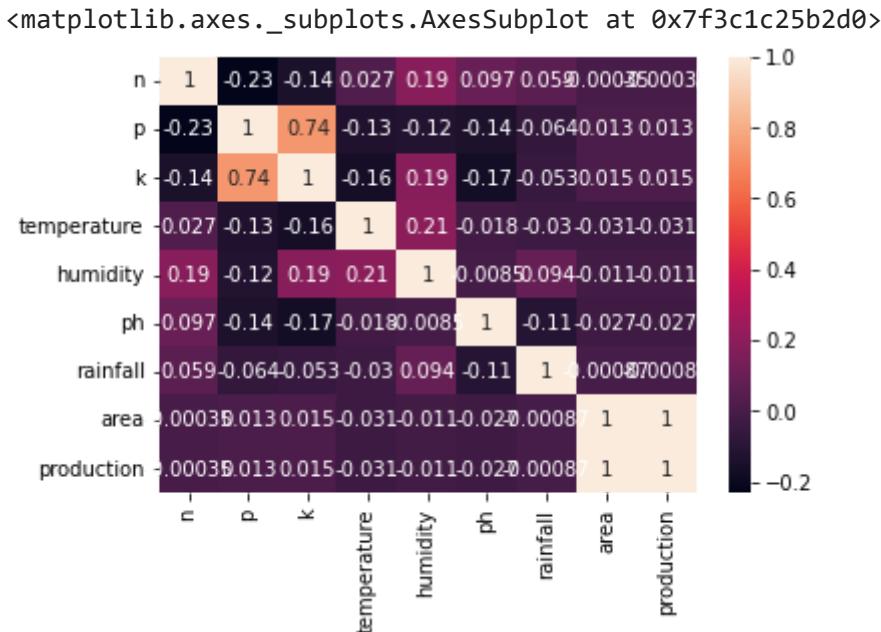
```
Number of various crops: 22
List of crops: ['rice' 'maize' 'chickpea' 'kidneybeans' 'pigeonpeas' 'mothbeans'
 'mungbean' 'blackgram' 'lentil' 'pomegranate' 'banana' 'mango' 'grapes'
 'watermelon' 'muskmelon' 'apple' 'orange' 'papaya' 'coconut' 'cotton'
 'jute' 'coffee']
```

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

```
jute      100
coconut   100
coffee    100
cotton    100
muskmelon 100
papaya   100
orange    100
pigeonpeas 100
lentil    100
grapes    100
apple     100
pomegranate 100
chickpea   100
mothbeans  100
blackgram  100
mungbean   100
banana    100
kidneybeans 100
rice      100
watermelon 100
maize     100
```

```
mango          100  
Name: crop, dtype: int64
```

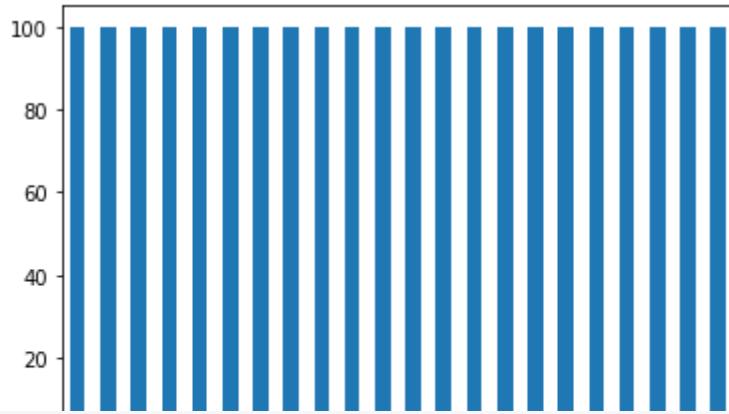
```
sns.heatmap(df.corr(), annot=True)
```



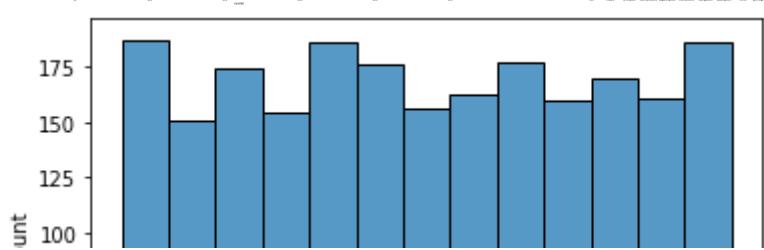
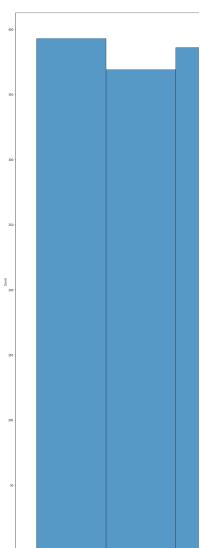
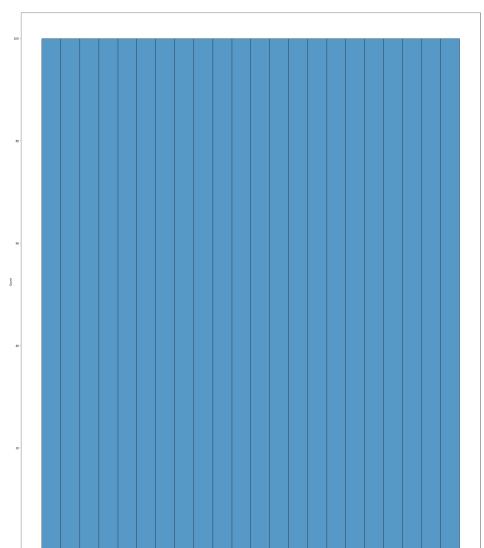
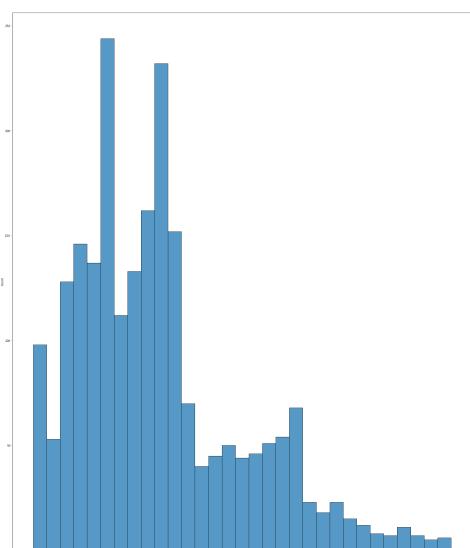
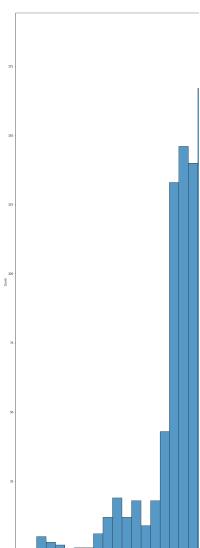
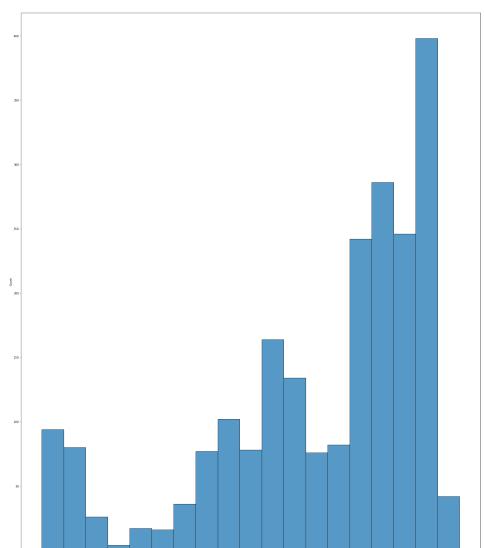
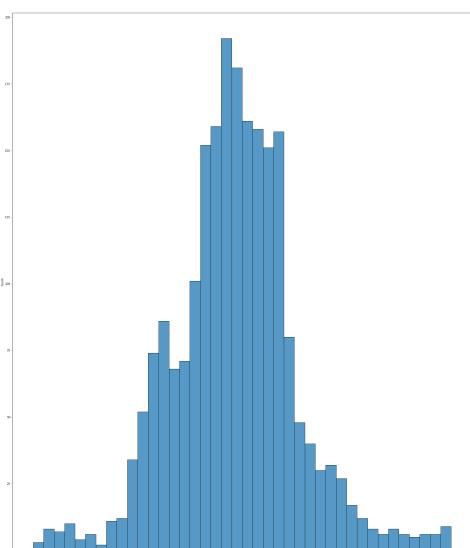
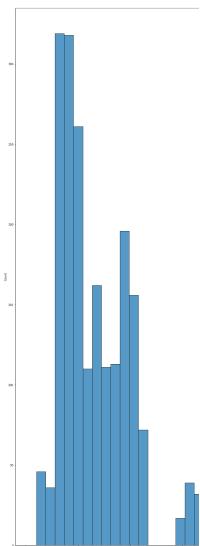
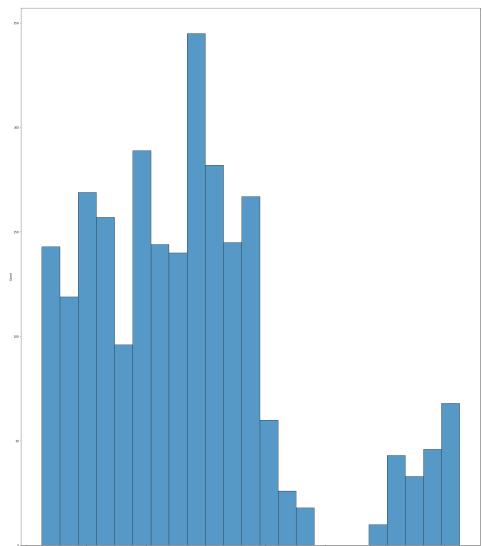
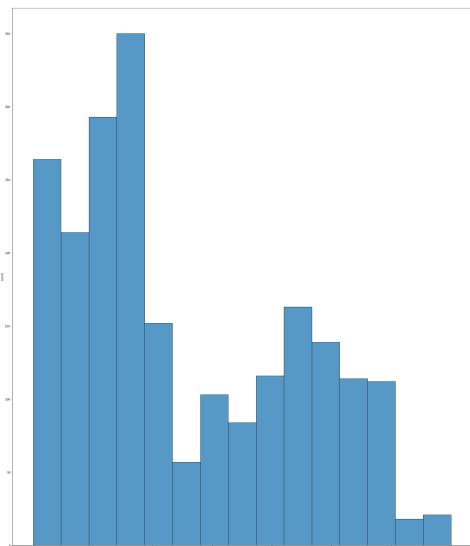
```
crop_summary = pd.pivot_table(df, index=[ 'crop' ],aggfunc='mean')  
crop_summary.head()
```

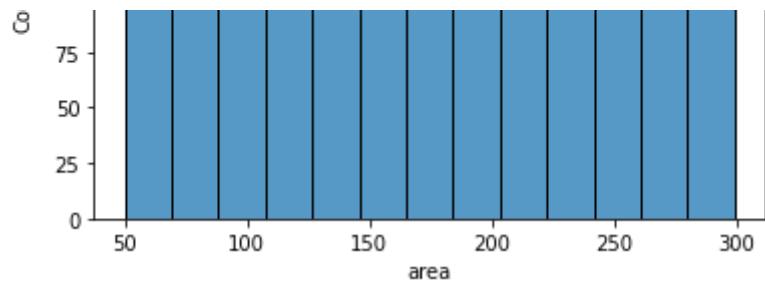
	area	humidity	k	n	p	ph	production	rainfall
crop								
apple	180.40	92.333383	199.89	20.80	134.22	5.929663	3968.80	112.654779
banana	163.85	80.358123	50.05	100.23	82.01	5.983893	3604.70	104.626980
blackgram	170.40	65.118426	19.24	40.02	67.47	7.133952	3748.80	67.884151
chickpea	177.40	16.860439	79.92	40.09	67.79	7.336957	3902.80	80.058977
coconut	173.04	94.844272	30.59	21.98	16.93	5.976562	3806.88	175.686646

```
all_columns = df.columns[:-1]  
  
labels = df["crop"].unique()  
df["crop"].value_counts().plot(kind="bar")  
  
plt.show()
```



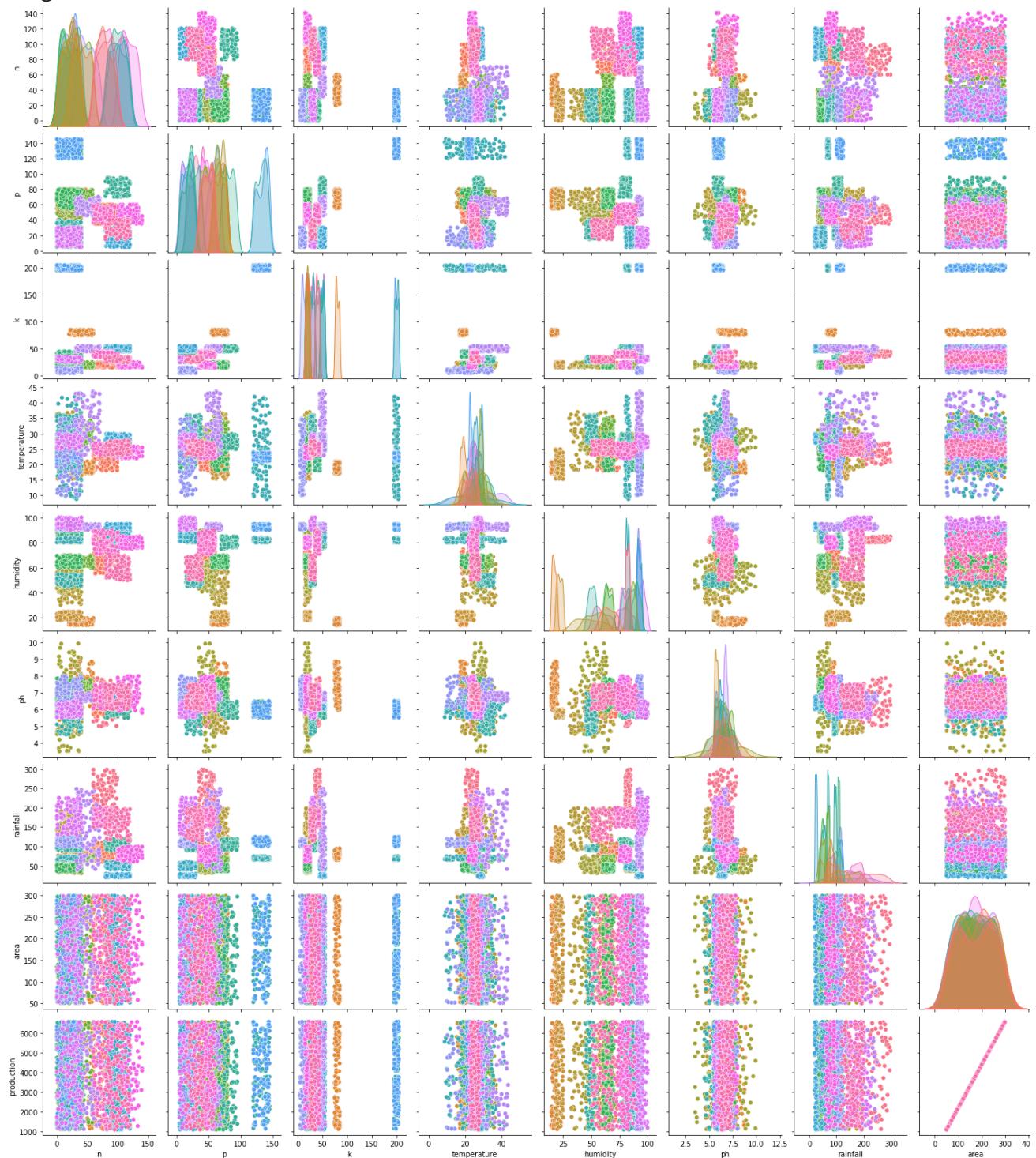
```
colorarr = ['#0592D0', '#Cd7f32', '#E97451', '#Bdb76b', '#954535', '#C2b280', '#808000', '#C  
#32cd32', '#39ff14', '#00ff7f', '#008080', '#36454f', '#F88379', '#Ff4500', '#Ff  
#Faf0e6', '#8c92ac', '#Dbd7d2', '#A7a6ba', '#B38b6d']  
= ≈ ≤ - ≈ ≥  
all_columns = df.columns[:-1]  
  
plt.figure(figsize=(100,120))  
i = 1  
for column in all_columns[:-1]:  
    plt.subplot(3,3,i)  
    sns.histplot(df[column])  
    i+=1  
plt.show()  
  
sns.histplot(df[all_columns[-1]])  
plt.show()
```





```
plt.figure(figsize=(19,17))
sns.pairplot(df, hue = "crop")
plt.show()
```

<Figure size 1368x1224 with 0 Axes>



```
from sklearn import preprocessing
le=preprocessing.LabelEncoder()
le.fit(['df.season'])
df['season']=le.fit_transform(df['season'])
```

```
f= plt.figure(figsize=(20,5))
ax=f.add_subplot(121)
sns.distplot(df['n'] , color ='red',ax=ax)

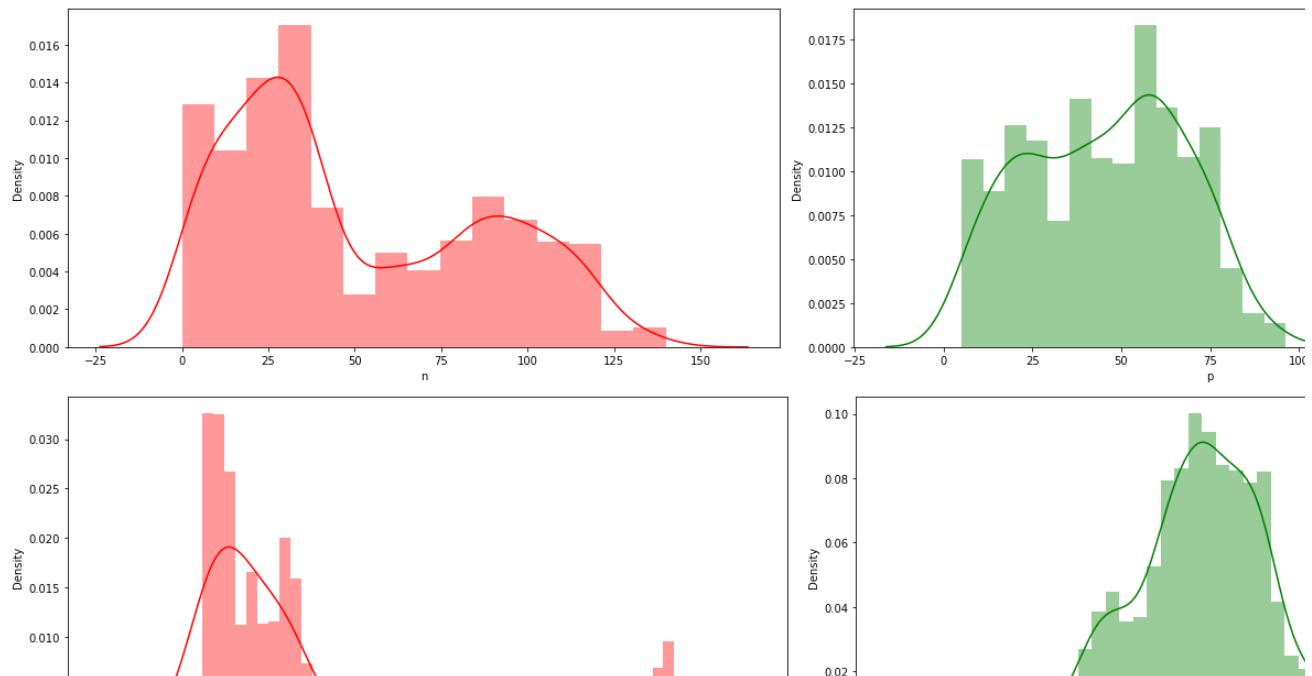
ax=f.add_subplot(122)
sns.distplot(df['p'] , color ='green' , ax = ax)
```

```
plt.tight_layout()
f= plt.figure(figsize=(20,5))
ax=f.add_subplot(121)
sns.distplot(df['k'] , color ='red',ax=ax)

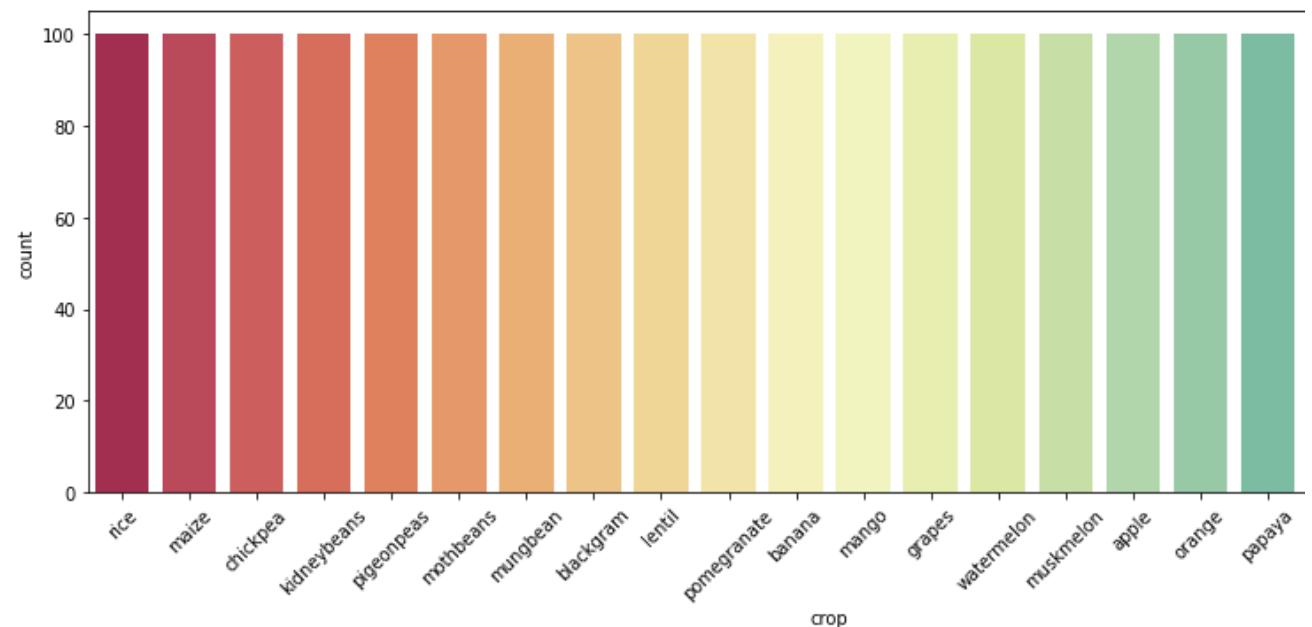
ax=f.add_subplot(122)
sns.distplot(df['temperature'] , color ='green' , ax = ax)
plt.tight_layout()
f= plt.figure(figsize=(20,5))
ax=f.add_subplot(121)
sns.distplot(df['humidity'] , color ='red',ax=ax)

ax=f.add_subplot(122)
sns.distplot(df['ph'] , color ='green' , ax = ax)
plt.tight_layout()
sns.distplot(df['rainfall'],color ='red')
sns.distplot(df['season'],color ='red')
```

```
<matplotlib.axes._subplots.AxesSubplot at 0x7f3c1bc254d0>
```



```
f= plt.figure(figsize=(15,5))
sns.countplot(df['crop'] , palette = 'Spectral')
plt.xticks(rotation=45)
plt.show()
```



```
crop_summary_N = crop_summary.sort_values(by='n', ascending=False)

fig = make_subplots(rows=1, cols=2)

top = {
    'y' : crop_summary_N['n'][0:10].sort_values().index,
    'x' : crop_summary_N['n'][0:10].sort_values()
}
```

```

last = {
    'y' : crop_summary_N['n'][ -10: ].index,
    'x' : crop_summary_N['n'][ -10: ]
}

fig.add_trace(
    go.Bar(top,
        name="Most nitrogen required",
        marker_color=random.choice(colorarr),
        orientation='h',
        text=top['x']),
    row=1, col=1
)

fig.add_trace(
    go.Bar(last,
        name="Least nitrogen required",
        marker_color=random.choice(colorarr),
        orientation='h',
        text=last['x']),
    row=1, col=2
)
fig.update_traces(texttemplate='%{text}', textposition='inside')
fig.update_layout(title_text="Nitrogen (N)",
                  plot_bgcolor='white',
                  font_size=12,
                  font_color='black',
                  height=500)
fig.update_xaxes(showgrid=False)
fig.update_yaxes(showgrid=False)
fig.show()

```

Nitrogen (N)

```
crop_summary_P = crop_summary.sort_values(by='p', ascending=False)

fig = make_subplots(rows=1, cols=2)

top = {
    'y' : crop_summary_P['p'][0:10].sort_values().index,
    'x' : crop_summary_P['p'][0:10].sort_values()
}

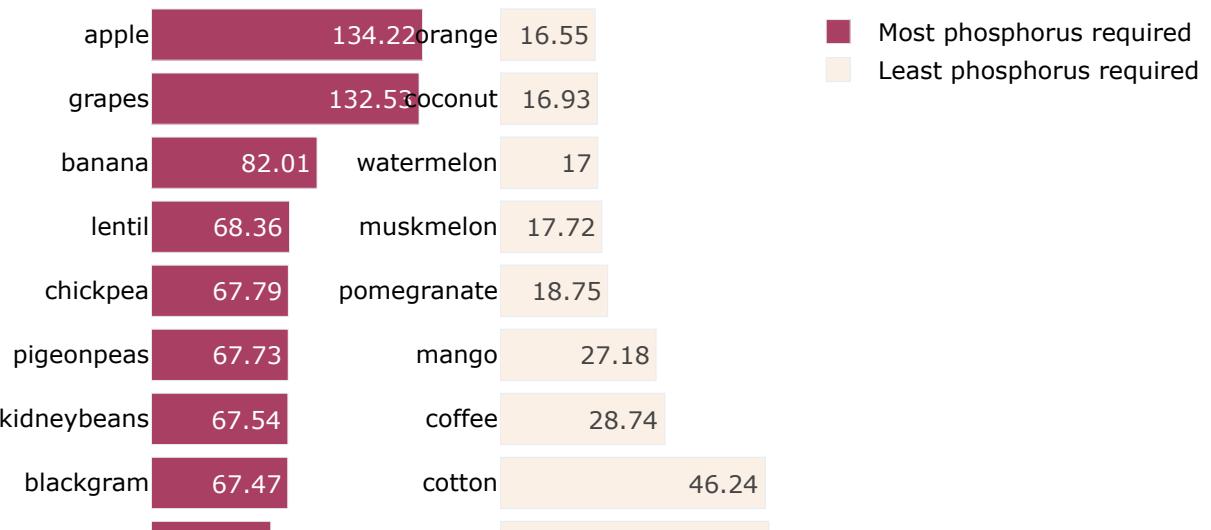
last = {
    'y' : crop_summary_P['p'][-10:].index,
    'x' : crop_summary_P['p'][-10:]
}

fig.add_trace(
    go.Bar(top,
        name="Most phosphorus required",
        marker_color=random.choice(colorarr),
        orientation='h',
        text=top['x']),
    row=1, col=1
)

fig.add_trace(
    go.Bar(last,
        name="Least phosphorus required",
        marker_color=random.choice(colorarr),
        orientation='h',
        text=last['x']),
    row=1, col=2
)
fig.update_traces(texttemplate='%{text}', textposition='inside')
fig.update_layout(title_text="Phosphorus (P)",
                  plot_bgcolor='white',
                  font_size=12,
                  font_color='black',
                  height=500)

fig.update_xaxes(showgrid=False)
fig.update_yaxes(showgrid=False)
fig.show()
```

Phosphorus (P)



```
crop_summary_K = crop_summary.sort_values(by='k', ascending=False)
```

```
fig = make_subplots(rows=1, cols=2)
```

```
top = {  
    'y' : crop_summary_K['k'][0:10].sort_values().index,  
    'x' : crop_summary_K['k'][0:10].sort_values()  
}
```

```
last = {  
    'y' : crop_summary_K['k'][-10:].index,  
    'x' : crop_summary_K['k'][-10:]  
}
```

```
fig.add_trace(  
    go.Bar(top,  
        name="Most potassium required",  
        marker_color=random.choice(colorarr),  
        orientation='h',  
        text=top['x']),  
    row=1, col=1  
)
```

```
fig.add_trace(  
    go.Bar(last,  
        name="Least potassium required",  
        marker_color=random.choice(colorarr),  
        orientation='h',  
        text=last['x']),  
    row=1, col=2  
)  
fig.update_traces(texttemplate='%{text}', textposition='inside')  
fig.update_layout(title_text="Potassium (K)",  
                  plot_bgcolor='white',
```

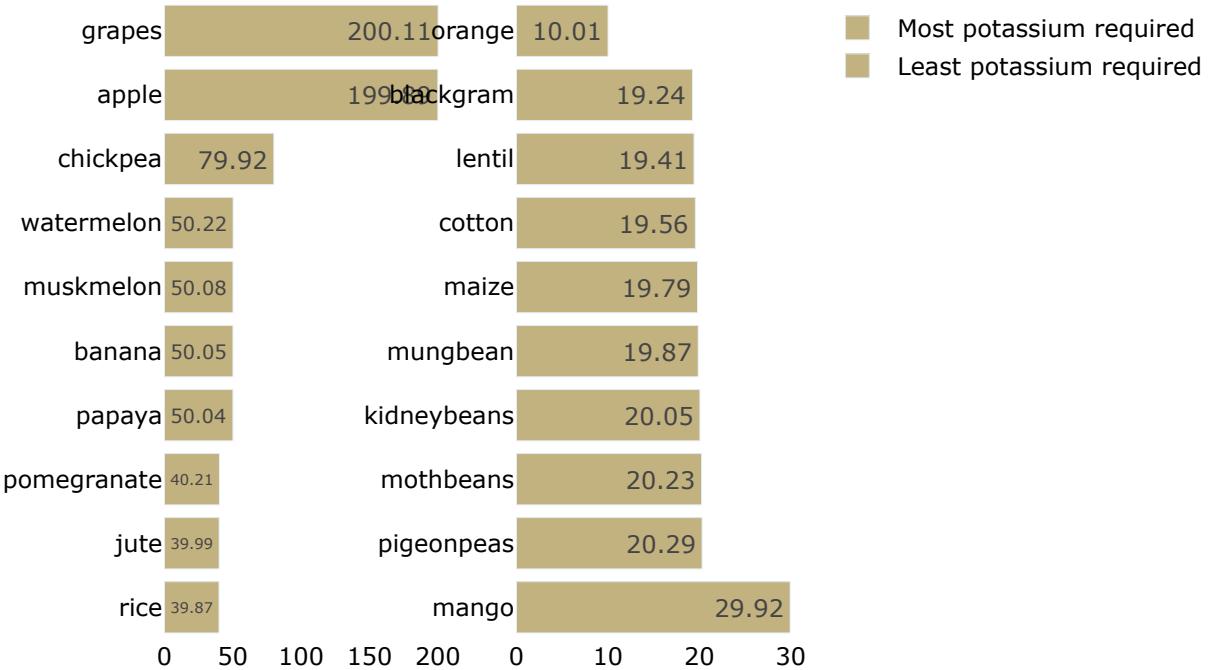
```

        font_size=12,
        font_color='black',
        height=500)

fig.update_xaxes(showgrid=False)
fig.update_yaxes(showgrid=False)
fig.show()

```

Potassium (K)



```

fig = go.Figure()
fig.add_trace(go.Bar(
    x=crop_summary.index,
    y=crop_summary['n'],
    name='Nitrogen',
    marker_color='blue'
))
fig.add_trace(go.Bar(
    x=crop_summary.index,
    y=crop_summary['p'],
    name='Phosphorous',
    marker_color='yellow'
))
fig.add_trace(go.Bar(
    x=crop_summary.index,
    y=crop_summary['k'],
    name='Potash',
    marker_color='crimson'
))

```

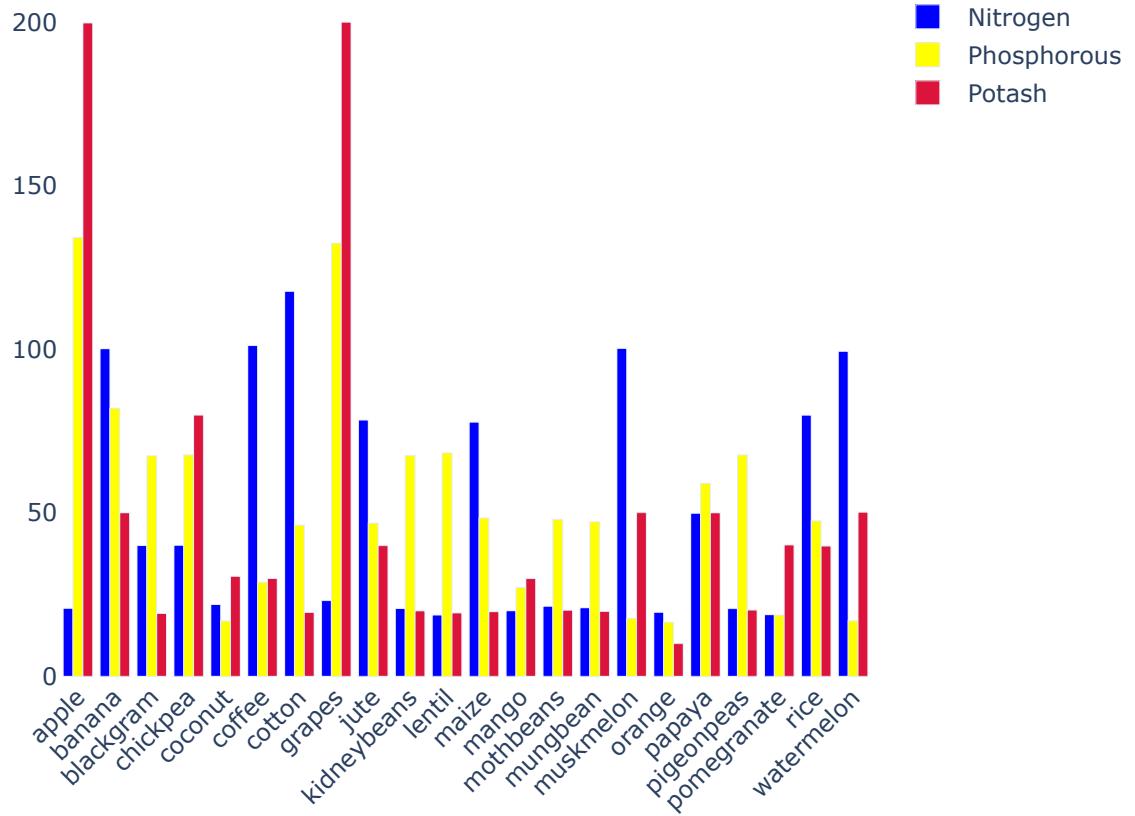
```

fig.update_layout(title="N, P, K values comparision between crops",
                  plot_bgcolor='white',
                  barmode='group',
                  xaxis_tickangle=-45)

fig.show()

```

N, P, K values comparision between crops



```

labels = ['Nitrogen(N)', 'Phosphorous(P)', 'Potash(K)']
fig = make_subplots(rows=1, cols=5, specs=[[{'type':'domain'}, {'type':'domain'},
                                              {'type':'domain'}, {'type':'domain'},
                                              {'type':'domain'}]])

rice_np = crop_summary[crop_summary.index=='rice']
values = [rice_np['n'][0], rice_np['p'][0], rice_np['k'][0]]
fig.add_trace(go.Pie(labels=labels, values=values, name="Rice"), 1, 1)

cotton_np = crop_summary[crop_summary.index=='cotton']
values = [cotton_np['n'][0], cotton_np['p'][0], cotton_np['k'][0]]
fig.add_trace(go.Pie(labels=labels, values=values, name="Cotton"), 1, 2)

jute_np = crop_summary[crop_summary.index=='jute']
values = [jute_np['n'][0], jute_np['p'][0], jute_np['k'][0]]
fig.add_trace(go.Pie(labels=labels, values=values, name="Jute"), 1, 3)

maize_np = crop_summary[crop_summary.index=='maize']
values = [maize_np['n'][0], maize_np['p'][0], maize_np['k'][0]]
fig.add_trace(go.Pie(labels=labels, values=values, name="Maize"), 1, 4)

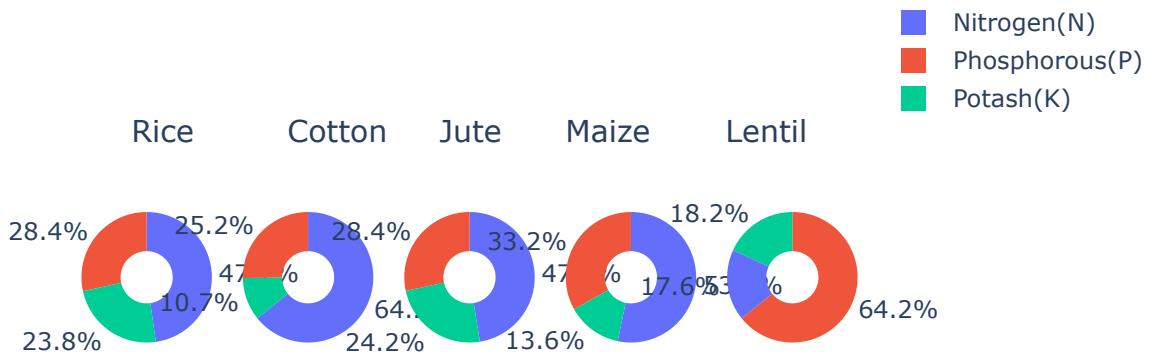
```

```

lentil_npk = crop_summary[crop_summary.index=='lentil']
values = [lentil_npk['n'][0], lentil_npk['p'][0], lentil_npk['k'][0]]
fig.add_trace(go.Pie(labels=labels, values=values, name="Lentil"), 1, 5)
fig.update_traces(hole=.4, hoverinfo="label+percent+name")
fig.update_layout(
    title_text="NPK ratio for rice, cotton, jute, maize, lentil",
    annotations=[dict(text='Rice', x=0.06, y=0.8, font_size=15, showarrow=False),
                 dict(text='Cotton', x=0.26, y=0.8, font_size=15, showarrow=False),
                 dict(text='Jute', x=0.5, y=0.8, font_size=15, showarrow=False),
                 dict(text='Maize', x=0.74, y=0.8, font_size=15, showarrow=False),
                 dict(text='Lentil', x=0.94, y=0.8, font_size=15, showarrow=False)])
fig.show()

```

NPK ratio for rice, cotton, jute, maize, lentil



```

labels = ['Nitrogen(N)', 'Phosphorous(P)', 'Potash(K)']
specs = [[{'type': 'domain'}, {'type': 'domain'}, {'type': 'domain'}, {'type': 'domain'}, {'type': 'domain'},
          {'type': 'domain'}, {'type': 'domain'}, {"type": "domain"}, {"type": "domain"}, {"type": "domain"}],
fig = make_subplots(rows=2, cols=5, specs=specs)
cafe_colors = ['rgb(255, 128, 0)', 'rgb(0, 153, 204)', 'rgb(173, 173, 133)']

apple_npk = crop_summary[crop_summary.index=='apple']
values = [apple_npk['n'][0], apple_npk['p'][0], apple_npk['k'][0]]
fig.add_trace(go.Pie(labels=labels, values=values, name="Apple", marker_colors=cafe_colors))

banana_npk = crop_summary[crop_summary.index=='banana']
values = [banana_npk['n'][0], banana_npk['p'][0], banana_npk['k'][0]]

```

```

fig.add_trace(go.Pie(labels=labels, values=values, name="Banana", marker_colors=cafe_colors)

grapes_npk = crop_summary[crop_summary.index=='grapes']
values = [grapes_npk['n'][0], grapes_npk['p'][0], grapes_npk['k'][0]]
fig.add_trace(go.Pie(labels=labels, values=values, name="Grapes", marker_colors=cafe_colors)

orange_npk = crop_summary[crop_summary.index=='orange']
values = [orange_npk['n'][0], orange_npk['p'][0], orange_npk['k'][0]]
fig.add_trace(go.Pie(labels=labels, values=values, name="Orange", marker_colors=cafe_colors)

mango_npk = crop_summary[crop_summary.index=='mango']
values = [mango_npk['n'][0], mango_npk['p'][0], mango_npk['k'][0]]
fig.add_trace(go.Pie(labels=labels, values=values, name="Mango", marker_colors=cafe_colors)

coconut_npk = crop_summary[crop_summary.index=='coconut']
values = [coconut_npk['n'][0], coconut_npk['p'][0], coconut_npk['k'][0]]
fig.add_trace(go.Pie(labels=labels, values=values, name="Coconut", marker_colors=cafe_color)

papaya_npk = crop_summary[crop_summary.index=='papaya']
values = [papaya_npk['n'][0], papaya_npk['p'][0], papaya_npk['k'][0]]
fig.add_trace(go.Pie(labels=labels, values=values, name="Papaya", marker_colors=cafe_colors)

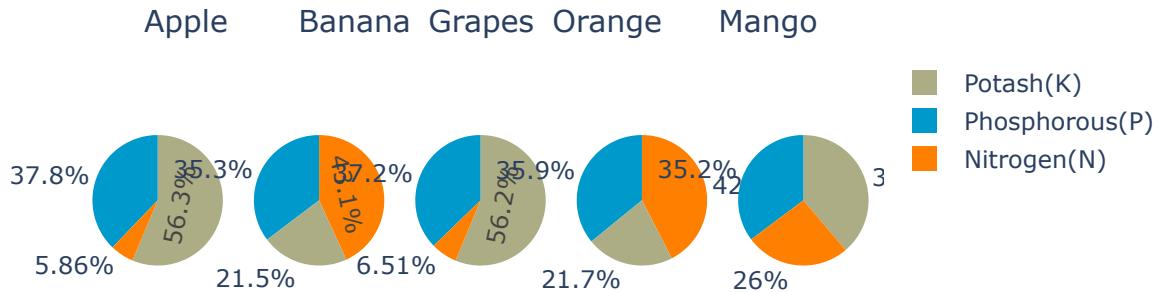
pomegranate_npk = crop_summary[crop_summary.index=='pomegranate']
values = [pomegranate_npk['n'][0], pomegranate_npk['p'][0], pomegranate_npk['k'][0]]
fig.add_trace(go.Pie(labels=labels, values=values, name="Pomegranate", marker_colors=cafe_c

watermelon_npk = crop_summary[crop_summary.index=='watermelon']
values = [watermelon_npk['n'][0], watermelon_npk['p'][0], watermelon_npk['k'][0]]
fig.add_trace(go.Pie(labels=labels, values=values, name="Watermelon", marker_colors=cafe_co

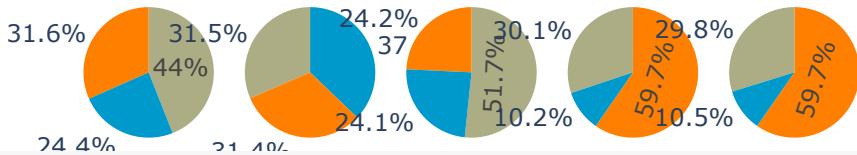
muskmelon_npk = crop_summary[crop_summary.index=='muskmelon']
values = [muskmelon_npk['n'][0], muskmelon_npk['p'][0], muskmelon_npk['k'][0]]
fig.add_trace(go.Pie(labels=labels, values=values, name="Muskmelon", marker_colors=cafe_col
fig.update_layout(
    title_text="NPK ratio for fruits",
    annotations=[dict(text='Apple', x=0.06, y=1.08, font_size=15, showarrow=False),
                 dict(text='Banana', x=0.26, y=1.08, font_size=15, showarrow=False),
                 dict(text='Grapes', x=0.50, y=1.08, font_size=15, showarrow=False),
                 dict(text='Orange', x=0.74, y=1.08, font_size=15, showarrow=False),
                 dict(text='Mango', x=0.94, y=1.08, font_size=15, showarrow=False),
                 dict(text='Coconut', x=0.06, y=0.46, font_size=15, showarrow=False),
                 dict(text='Papaya', x=0.26, y=0.46, font_size=15, showarrow=False),
                 dict(text='Pomegranate', x=0.50, y=0.46, font_size=15, showarrow=False),
                 dict(text='Watermelon', x=0.74, y=0.46, font_size=15, showarrow=False),
                 dict(text='Muskmelon', x=0.94, y=0.46, font_size=15, showarrow=False)])
fig.show()

```

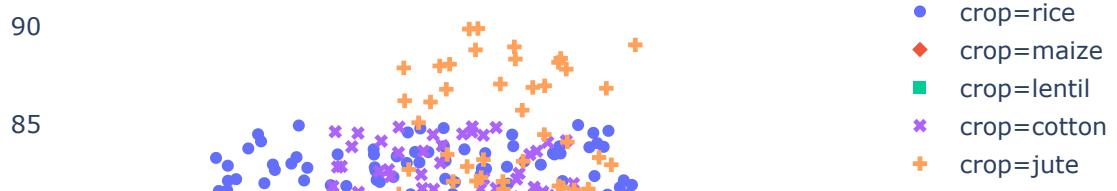
NPK ratio for fruits



Coconut Papaya Pomegranate Watermelon Muskmelon



```
crop_scatter = df[(df['crop']=='rice') |  
                   (df['crop']=='jute') |  
                   (df['crop']=='cotton') |  
                   (df['crop']=='maize') |  
                   (df['crop']=='lentil')]  
  
fig = px.scatter(crop_scatter, x="temperature", y="humidity", color="crop", symbol="crop")  
fig.update_layout(plot_bgcolor='white')  
fig.update_xaxes(showgrid=False)  
fig.update_yaxes(showgrid=False)  
  
fig.show()
```

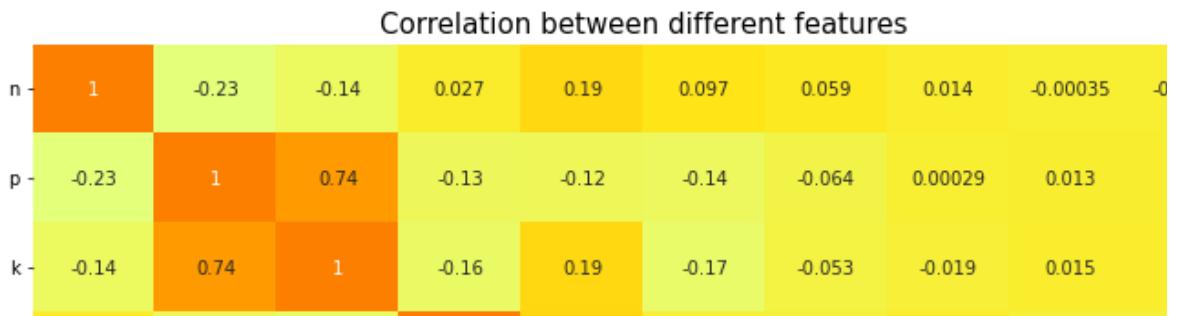


```
df['season']
```

```
0      5
1      4
2      0
3      1
4      0
..
2195    4
2196    0
2197    2
2198    4
2199    5
Name: season, Length: 2200, dtype: int64
```

```
fig, ax = plt.subplots(1, 1, figsize=(15, 9))
sns.heatmap(df.corr(), annot=True, cmap='Wistia' )
ax.set(xlabel='features')
ax.set(ylabel='features')

plt.title('Correlation between different features', fontsize = 15, c='black')
plt.show()
```



```
target=['crop']
features=['n','p','k','temperature','humidity','ph','rainfall','season']
```

```
humidity -0.0085 0.094 -0.001 -0.011
```

```
X=df[features]
y=df[target]
```

```
from sklearn.model_selection import train_test_split
Xtrain, Xtest, Ytrain,Ytest = train_test_split(X, y, test_size = 0.3,
                                              shuffle = True, random_state = 0)
```

```
# Initializing empty lists to append all model's name and corresponding name
acc = []
model = []
```

ID3 DECISION TREE

```
from sklearn.tree import DecisionTreeClassifier

DecisionTree = DecisionTreeClassifier(criterion="entropy",random_state=2,max_depth=4)

DecisionTree.fit(Xtrain,Ytrain)

predicted_values = DecisionTree.predict(Xtest)
x = metrics.accuracy_score(Ytest, predicted_values)
acc.append(x)
model.append('Decision Tree')
print("DecisionTrees's Accuracy is: ", x*100)

print(classification_report(Ytest,predicted_values))
plt.figure(figsize = (15,9))
sns.heatmap(confusion_matrix(Ytest, predicted_values), annot = True)
plt.title("Confusion Matrix for Test Data")
plt.show()
```

DecisionTrees's Accuracy is: 65.30303030303031

	precision	recall	f1-score	support
apple	1.00	1.00	1.00	28
banana	0.50	1.00	0.67	30
blackgram	0.49	1.00	0.66	31
chickpea	1.00	1.00	1.00	34
coconut	1.00	1.00	1.00	26
coffee	1.00	0.90	0.95	29
cotton	0.47	1.00	0.64	28
grapes	0.00	0.00	0.00	30
jute	1.00	0.23	0.37	31
kidneybeans	1.00	0.81	0.89	26
lentil	0.51	1.00	0.68	22
maize	0.00	0.00	0.00	27
mango	1.00	0.93	0.96	28
mothbeans	0.00	0.00	0.00	36
mungbean	0.00	0.00	0.00	29
muskmelon	1.00	1.00	1.00	30
orange	0.00	0.00	0.00	34
papaya	0.00	0.00	0.00	39
pigeonpeas	0.61	1.00	0.76	28
pomegranate	0.48	1.00	0.65	32
rice	0.37	1.00	0.54	37
watermelon	1.00	1.00	1.00	25
accuracy			0.65	660
macro avg	0.57	0.68	0.58	660
weighted avg	0.54	0.65	0.56	660

Confusion Matrix for Test Data

NAIVE BAYES ALGORITHM

```
from sklearn.naive_bayes import GaussianNB
```

```
NaiveBayes = GaussianNB()

NaiveBayes.fit(Xtrain,Ytrain)

predicted_values = NaiveBayes.predict(Xtest)
x = metrics.accuracy_score(Ytest, predicted_values)
acc.append(x)
model.append('Naive Bayes')
print("Naive Bayes's Accuracy is: ", x*100)

print(classification_report(Ytest,predicted_values))
plt.figure(figsize = (15,9))
sns.heatmap(confusion_matrix(Ytest,predicted_values), annot = True)
plt.title("Confusion Matrix for Test Data")
plt.show()
```

Naive Bayes's Accuracy is: 99.39393939393939

	precision	recall	f1-score	support
apple	1.00	1.00	1.00	28
banana	1.00	1.00	1.00	30
blackgram	1.00	1.00	1.00	31
chickpea	1.00	1.00	1.00	34
coconut	1.00	1.00	1.00	26
coffee	1.00	1.00	1.00	29
cotton	1.00	1.00	1.00	28
grapes	1.00	1.00	1.00	30
jute	0.91	0.97	0.94	31
kidneybeans	1.00	1.00	1.00	26
lentil	1.00	1.00	1.00	22
maize	1.00	1.00	1.00	27
mango	1.00	1.00	1.00	28
mothbeans	1.00	1.00	1.00	36
mungbean	1.00	1.00	1.00	29
muskmelon	1.00	1.00	1.00	30

KNN Classifier

```
प्रगतिशीलता १.०० १.०० १.०० २०
```

```
from sklearn.neighbors import KNeighborsClassifier
clf_knn = KNeighborsClassifier(n_neighbors=4)
clf_knn.fit(Xtrain,Ytrain)
y_pred1 = clf_knn.predict(Xtest)
accu=metrics.accuracy_score(Ytest,y_pred1)

acc.append(accu)
model.append('KNN Classifier')

print("Accuracy Score of KNN:",accu*100)

print(classification_report(Ytest,y_pred1))
plt.figure(figsize = (15,9))
sns.heatmap(confusion_matrix(Ytest,y_pred1), annot = True)
plt.title("Confusion Matrix for Test Data")
plt.show()
```

Accuracy Score of KNN: 97.72727272727273

	precision	recall	f1-score	support
apple	1.00	1.00	1.00	28
banana	1.00	1.00	1.00	30
blackgram	0.97	1.00	0.98	31
chickpea	1.00	1.00	1.00	34
coconut	1.00	1.00	1.00	26
coffee	1.00	1.00	1.00	29
cotton	0.97	1.00	0.98	28
grapes	1.00	1.00	1.00	30
jute	0.79	1.00	0.89	31
kidneybeans	0.87	1.00	0.93	26
lentil	0.96	1.00	0.98	22
maize	1.00	0.96	0.98	27
mango	1.00	1.00	1.00	28
mothbeans	1.00	0.94	0.97	36
mungbean	1.00	1.00	1.00	29
muskmelon	1.00	1.00	1.00	30
orange	1.00	1.00	1.00	34
papaya	1.00	0.97	0.99	39
pigeonpeas	1.00	0.86	0.92	28
pomegranate	1.00	1.00	1.00	32
rice	1.00	0.81	0.90	37
watermelon	1.00	1.00	1.00	25
accuracy			0.98	660
macro avg	0.98	0.98	0.98	660
weighted avg	0.98	0.98	0.98	660

Confusion Matrix for Test Data

LOGISTIC REGRESSION

```
- 0 0 1 0 0 0 0 0 0 0 1  
from sklearn.linear_model import LogisticRegression  
  
LogReg = LogisticRegression(random_state=2)  
  
LogReg.fit(Xtrain,Ytrain)  
  
predicted_values = LogReg.predict(Xtest)  
  
x = metrics.accuracy_score(Ytest, predicted_values)
```

```
acc.append(x)
model.append('Logistic Regression')
print("Logistic Regression's Accuracy is: ", x*100, '\n')

print(classification_report(Ytest,predicted_values))
plt.figure(figsize = (15,9))
sns.heatmap(confusion_matrix(Ytest,predicted_values), annot = True)
plt.title("Confusion Matrix for Test Data")
plt.show()
```

Logistic Regression's Accuracy is: 95.90909090909090

	precision	recall	f1-score	support
apple	1.00	1.00	1.00	28
banana	1.00	1.00	1.00	30
blackgram	0.84	0.87	0.86	31
chickpea	1.00	1.00	1.00	34
coconut	1.00	1.00	1.00	26
coffee	1.00	1.00	1.00	29
cotton	0.82	0.96	0.89	28
grapes	1.00	1.00	1.00	30
jute	0.88	0.97	0.92	31
...

RANDOM FOREST

```
mango      1.00      1.00      1.00      28
from sklearn.ensemble import RandomForestClassifier

RF = RandomForestClassifier(n_estimators=20, random_state=0)
RF.fit(Xtrain,Ytrain)

predicted_values = RF.predict(Xtest)

x = metrics.accuracy_score(Ytest, predicted_values)
acc.append(x)
model.append('Random Forest')
print("RF's Accuracy is: ", x*100)

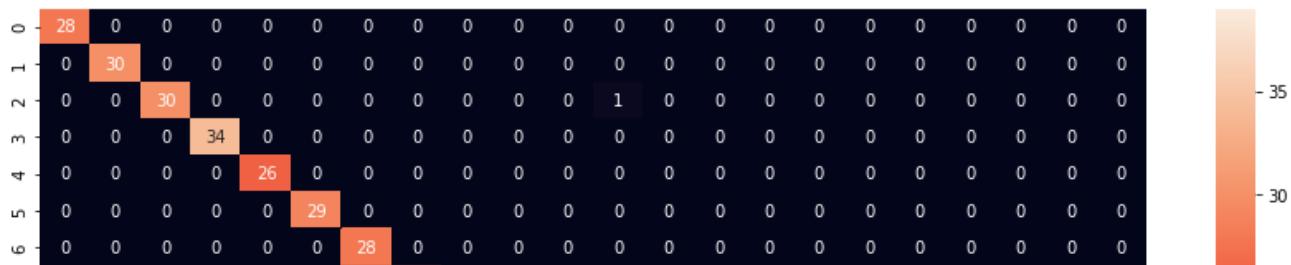
print(classification_report(Ytest,predicted_values))
plt.figure(figsize = (15,9))
sns.heatmap(confusion_matrix(Ytest,predicted_values), annot = True)
plt.title("Confusion Matrix for Test Data")
plt.show()
```



RF's Accuracy is: 99.69696969696969

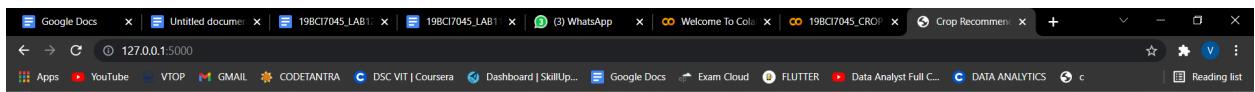
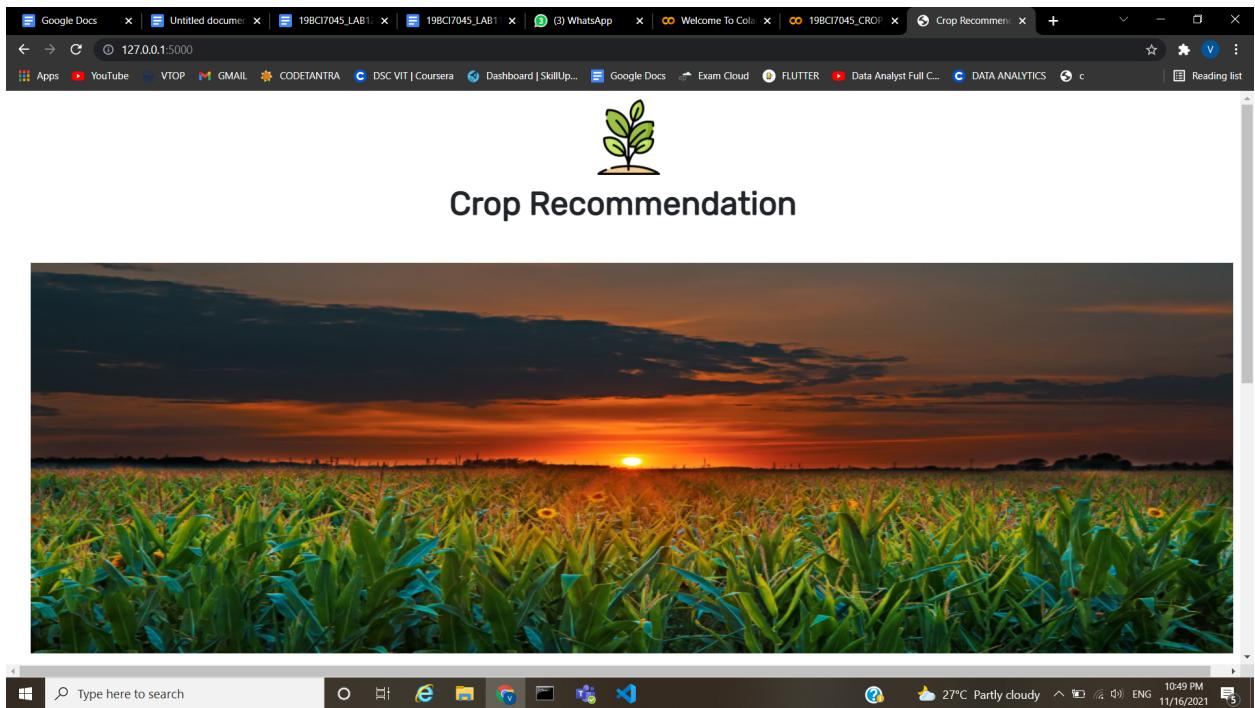
	precision	recall	f1-score	support
apple	1.00	1.00	1.00	28
banana	1.00	1.00	1.00	30
blackgram	1.00	0.97	0.98	31
chickpea	1.00	1.00	1.00	34
coconut	1.00	1.00	1.00	26
coffee	1.00	1.00	1.00	29
cotton	1.00	1.00	1.00	28
grapes	1.00	1.00	1.00	30
jute	0.97	1.00	0.98	31
kidneybeans	1.00	1.00	1.00	26
lentil	1.00	1.00	1.00	22
maize	0.96	1.00	0.98	27
mango	1.00	1.00	1.00	28
mothbeans	1.00	1.00	1.00	36
mungbean	1.00	1.00	1.00	29
muskmelon	1.00	1.00	1.00	30
orange	1.00	1.00	1.00	34
papaya	1.00	1.00	1.00	39
pigeonpeas	1.00	1.00	1.00	28
pomegranate	1.00	1.00	1.00	32
rice	1.00	0.97	0.99	37
watermelon	1.00	1.00	1.00	25
accuracy			1.00	660
macro avg	1.00	1.00	1.00	660
weighted avg	1.00	1.00	1.00	660

Confusion Matrix for Test Data



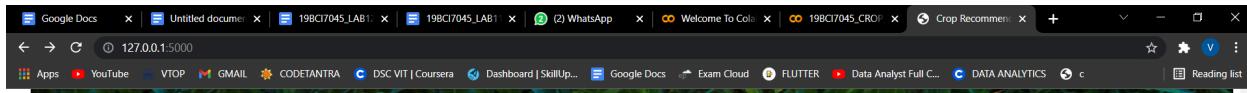
```
plt.figure(figsize=[10,5],dpi = 100)
plt.title('Accuracy Comparison')
plt.xlabel('Accuracy')
plt.ylabel('Algorithm')
sns.barplot(x = acc,y = model,palette='dark')
```

DISPLAYING WEBPAGE :



EXAMPLES :

1.



This web application is created in the wake of helping the farmers get the best suited crop depending on various parameters like Nitrogen, Phosphorous, Potassium, Temperature, Rainfall, PH, Season and Humidity. Please enter the parameters below and our ML model will predict crop with accuracy of 99%

Nitrogen :

Phosphorous :

Potassium :

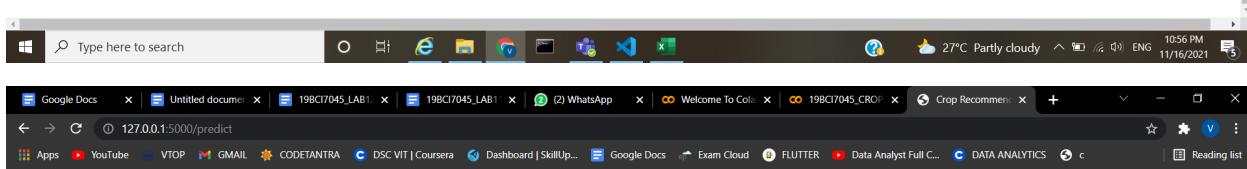
Temperature :

Humidity :

PH :

Rainfall :

Season :



This web application is created in the wake of helping the farmers get the best suited crop depending on various parameters like Nitrogen, Phosphorous, Potassium, Temperature, Rainfall, PH, Season and Humidity. Please enter the parameters below and our ML model will predict crop with accuracy of 99%

Nitrogen :

Phosphorous :

Potassium :

Temperature :

Humidity :

PH :

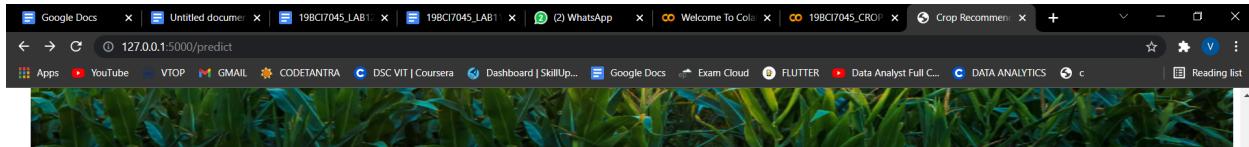
Rainfall :

Season :

Recommended Crop is rice

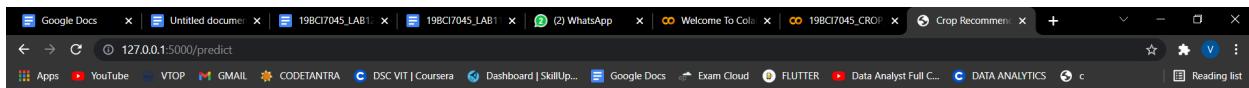


2.



This web application is created in the wake of helping the farmers get the best suited crop depending on various parameters like Nitrogen, Phosphorous, Potassium, Temperature, Rainfall, PH, Season and Humidity. Please enter the parameters below and our ML model will predict crop with accuracy of 99%

Nitrogen :
Phosphorous :
Potassium :
Temperature :
Humidity :
PH :
Rainfall :
Season :



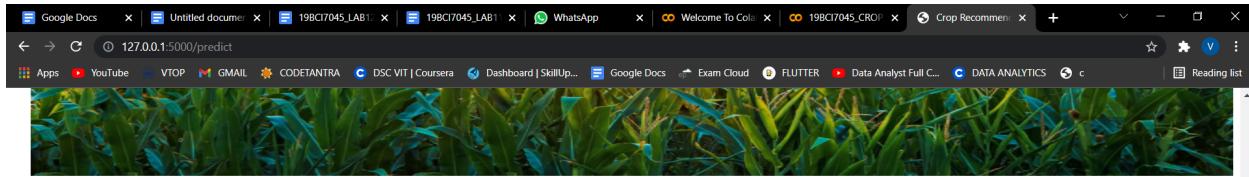
This web application is created in the wake of helping the farmers get the best suited crop depending on various parameters like Nitrogen, Phosphorous, Potassium, Temperature, Rainfall, PH, Season and Humidity. Please enter the parameters below and our ML model will predict crop with accuracy of 99%

Nitrogen :
Phosphorous :
Potassium :
Temperature :
Humidity :
PH :
Rainfall :
Season :

Recommended Crop is pomegranate



3.



This web application is created in the wake of helping the farmers get the best suited crop depending on various parameters like Nitrogen, Phosphorous, Potassium, Temperature, Rainfall, PH, Season and Humidity. Please enter the parameters below and our ML model will predict crop with accuracy of 99%

Nitrogen :

Phosphorous :

Potassium :

Temperature :

Humidity :

PH :

Rainfall :

Season :



This web application is created in the wake of helping the farmers get the best suited crop depending on various parameters like Nitrogen, Phosphorous, Potassium, Temperature, Rainfall, PH, Season and Humidity. Please enter the parameters below and our ML model will predict crop with accuracy of 99%

Nitrogen :

Phosphorous :

Potassium :

Temperature :

Humidity :

PH :

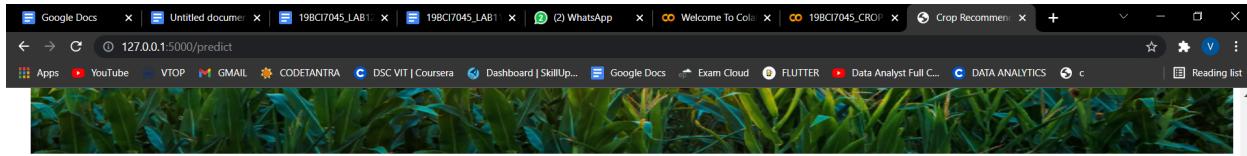
Rainfall :

Season :

Recommended Crop is mango



4.



This web application is created in the wake of helping the farmers get the best suited crop depending on various parameters like Nitrogen, Phosphorous, Potassium, Temperature, Rainfall, PH, Season and Humidity. Please enter the parameters below and our ML model will predict crop with accuracy of 99%

Nitrogen :

Phosphorous :

Potassium :

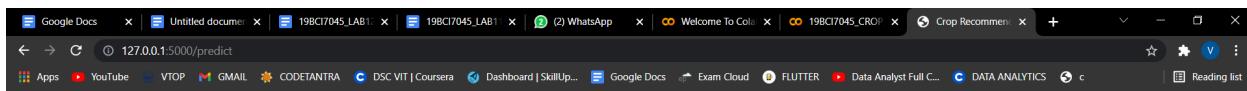
Temperature :

Humidity :

PH :

Rainfall :

Season :



This web application is created in the wake of helping the farmers get the best suited crop depending on various parameters like Nitrogen, Phosphorous, Potassium, Temperature, Rainfall, PH, Season and Humidity. Please enter the parameters below and our ML model will predict crop with accuracy of 99%

Nitrogen :

Phosphorous :

Potassium :

Temperature :

Humidity :

PH :

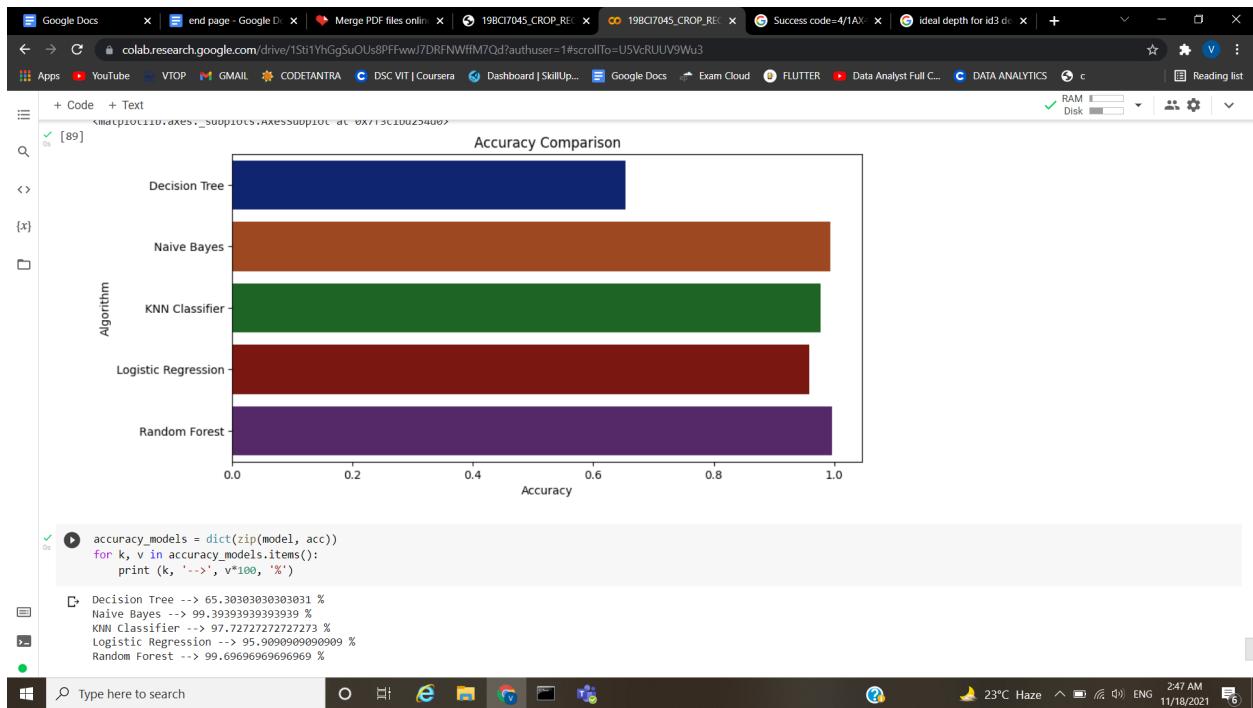
Rainfall :

Season :

Recommended Crop is coffee



ACCURACY :



Highest Accuracy : 99.69% (Random Forest)

Least Accuracy : 65.3 % (Decision Tree)

Colab link :

<https://colab.research.google.com/drive/1Sti1YhGgSuOUs8PFFwwJ7DRFNWffM7Qd?usp=sharing>