```
# This Python 3 environment comes with many helpful analytics
libraries installed
# It is defined by the kaggle/python Docker image:
https://github.com/kaggle/docker-python
# For example, here's several helpful packages to load
import numpy as np # linear algebra
import pandas as pd # data processing, CSV file I/O (e.g. pd.read csv)
# Input data files are available in the read-only "../input/"
directory
# For example, running this (by clicking run or pressing Shift+Enter)
will list all files under the input directory
import os
for dirname, _, filenames in os.walk('/kaggle/input'):
    for filename in filenames:
        # print(os.path.join(dirname, filename))
        pass
# You can write up to 20GB to the current directory (/kaggle/working/)
that gets preserved as output when you create a version using "Save &
Run All"
# You can also write temporary files to /kaggle/temp/, but they won't
be saved outside of the current session
```

Importing Required Modules

```
import os
import numpy as np
from sklearn.ensemble import RandomForestClassifier
from sklearn.model selection import train test split
from sklearn.metrics import classification report
from tensorflow.keras.applications import ResNet50
from tensorflow.keras.applications.resnet50 import preprocess input
from tensorflow.keras.preprocessing.image import load img,
img_to_array
2025-09-24 10:23:45.231494: E
external/local xla/xla/stream executor/cuda/cuda fft.cc:4771 Unable to
register cuFFT factory: Attempting to register factory for plugin
cuFFT when one has already been registered
WARNING: All log messages before absl::InitializeLog() is called are
written to STDERR
E0000 00:00:1758709425.587266 36 cuda dnn.cc:8310] Unable to
register cuDNN factory: Attempting to register factory for plugin
cuDNN when one has already been registered
E0000 00:00:1758709425.674907
                                   36 cuda blas.cc:1418] Unable to
register cuBLAS factory: Attempting to register factory for plugin
cuBLAS when one has already been registered
```

Loading the images

Loading the dataset

```
# [ Load dataset
dataset_path = '/kaggle/input/cassava-disease-dataset/Cassava' #
Replace with your actual path
X_images, y_labels = load_images_from_folder(dataset_path)
print("Done loading the dataset")
Done loading the dataset
```

Feature Extraction using ResNet 50

```
# □ Step 2: Feature extraction using ResNet50
resnet model = ResNet50(weights='imagenet', include top=False,
pooling='avg')
features = resnet model.predict(X images, verbose=1)
I0000 00:00:1758709481.362061
                                   36 gpu device.cc:2022] Created
device /job:localhost/replica:0/task:0/device:GPU:0 with 13942 MB
memory: -> device: 0, name: Tesla T4, pci bus id: 0000:00:04.0,
compute capability: 7.5
I0000 00:00:1758709481.362745
                                   36 gpu device.cc:20221 Created
device /job:localhost/replica:0/task:0/device:GPU:1 with 13942 MB
memory: -> device: 1, name: Tesla T4, pci bus id: 0000:00:05.0,
compute capability: 7.5
Downloading data from https://storage.googleapis.com/tensorflow/keras-
applications/resnet/
resnet50 weights tf dim ordering tf kernels notop.h5
94765736/94765736
                                     - Os Ous/step
```

```
WARNING: All log messages before absl::InitializeLog() is called are
written to STDERR
I0000 00:00:1758709498.030340
                                  98 service.cc:1481 XLA service
0x7c257414b020 initialized for platform CUDA (this does not quarantee
that XLA will be used). Devices:
I0000 00:00:1758709498.031380
                                  98 service.cc:156] StreamExecutor
device (0): Tesla T4, Compute Capability 7.5
I0000 00:00:1758709498.031402
                                  98 service.cc:156] StreamExecutor
device (1): Tesla T4, Compute Capability 7.5
I0000 00:00:1758709498.663101 98 cuda dnn.cc:529] Loaded cuDNN
version 90300
 2/235 ——
                       ——— 19s 86ms/step
                                  98 device compiler.h:188] Compiled
I0000 00:00:1758709502.930430
cluster using XLA! This line is logged at most once for the lifetime
of the process.
235/235 ———
                      ----- 31s 101ms/step
```

Data Split

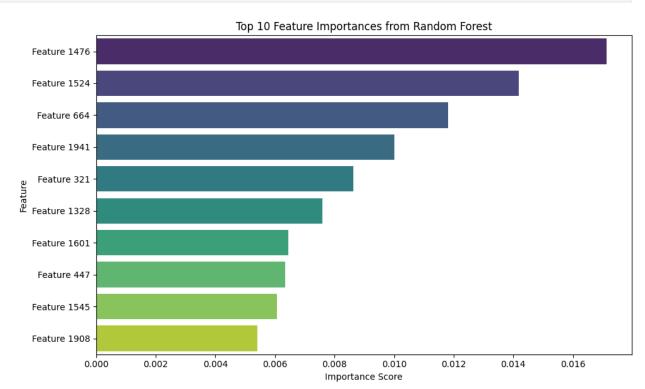
```
# Train-test split
X_train, X_test, y_train, y_test = train_test_split(features,
y_labels, test_size=0.2, random_state=42)
```

Training the random forest model

```
# □ Step 4: Train Random Forest classifier
hybrid rf model = RandomForestClassifier(n estimators=100,
random state=42)
hybrid rf_model.fit(X_train, y_train)
RandomForestClassifier(random state=42)
import matplotlib.pyplot as plt
import seaborn as sns
import pandas as pd
# Get feature importances from the trained model
importances = hybrid rf model.feature importances
# If your features are unnamed (e.g., from CNN), label them
numerically
feature names = [f'Feature {i}' for i in range(len(importances))]
# Create a DataFrame for visualization
feature df = pd.DataFrame({
    'Feature': feature names,
    'Importance': importances
```

```
}).sort_values(by='Importance', ascending=False).head(10)

# Plot the top 10 features
plt.figure(figsize=(10, 6))
sns.barplot(x='Importance', y='Feature', data=feature_df,
palette='viridis')
plt.title('Top 10 Feature Importances from Random Forest')
plt.xlabel('Importance Score')
plt.ylabel('Feature')
plt.tight_layout()
plt.show()
```



Performance Evaluation

```
# □ Step 5: Evaluate
hybrid_y_pred = hybrid_rf_model.predict(X_test)
print(classification_report(y_test, hybrid_y_pred))
                                recall f1-score
                                                    support
                   precision
                                  0.92
bacterial blight
                        0.85
                                             0.88
                                                         533
      brown spot
                        0.90
                                  0.81
                                             0.85
                                                         291
      green mite
                        0.88
                                  0.86
                                             0.87
                                                         206
                        0.93
                                  0.92
                                             0.92
                                                         226
         healthy
          mosaic
                        0.94
                                  0.91
                                             0.93
                                                         246
```

accuracy			0.89	1502
macro avg	0.90	0.88	0.89	1502
weighted avg	0.89	0.89	0.89	1502

Building a Random Forest Model without the feature extraction

```
import numpy as np
from sklearn.ensemble import RandomForestClassifier
from sklearn.model_selection import train_test_split
from sklearn.metrics import classification_report

# Flatten image data: (num_samples, height, width, channels) →
    (num_samples, features)
X_flat = X_images.reshape(X_images.shape[0], -1)

# Train-test split
rf_X_train, rf_X_test, rf_y_train, _ = train_test_split(X_flat,
y_labels, test_size=0.2, random_state=42)

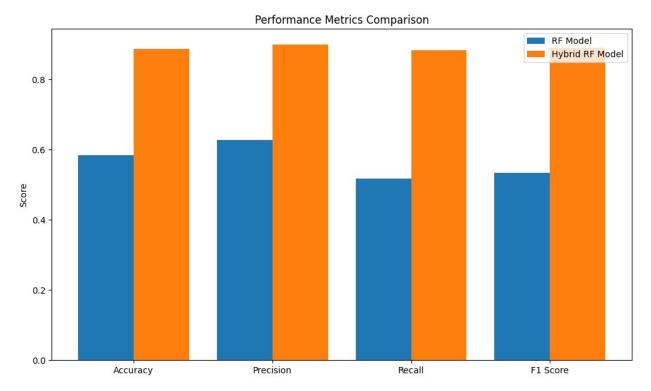
# Train Random Forest
rf_model = RandomForestClassifier(n_estimators=100, random_state=42)
rf_model.fit(rf_X_train, rf_y_train)
RandomForestClassifier(random_state=42)
```

Evaluating the model's performance

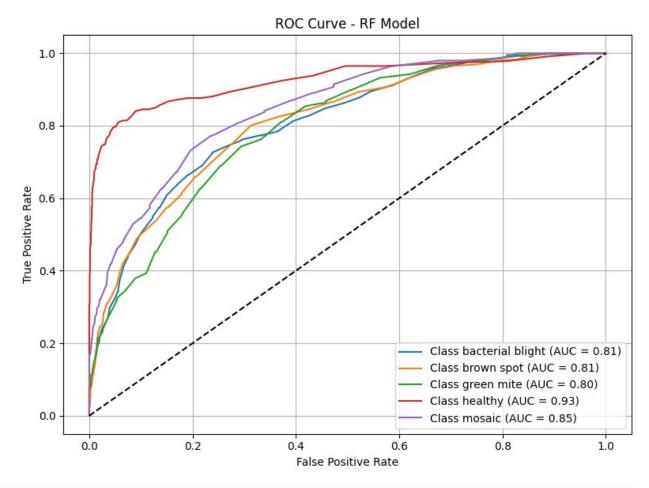
```
# Evaluate
y_pred = rf_model.predict(rf_X_test)
print(classification report(y test, y pred))
                                 recall f1-score
                   precision
                                                     support
bacterial blight
                        0.52
                                   0.85
                                             0.65
                                                         533
                        0.57
                                   0.42
                                             0.49
                                                         291
      brown spot
                        0.57
                                   0.17
                                             0.27
                                                         206
      green mite
         healthy
                        0.82
                                   0.76
                                             0.79
                                                         226
          mosaic
                        0.65
                                   0.38
                                             0.48
                                                         246
                                             0.59
                                                        1502
        accuracy
                        0.63
                                   0.52
                                              0.53
                                                        1502
       macro avg
    weighted avg
                        0.60
                                   0.59
                                             0.56
                                                        1502
```

Model Performance Comparison

```
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.metrics import (
    accuracy score, precision score, recall score, f1 score,
    roc curve, auc, confusion matrix, precision recall curve
# Evaluate metrics
def get metrics(y true, y pred):
    return {
        'Accuracy': accuracy_score(y_true, y_pred),
        'Precision': precision score(y true, y pred ,
average='macro'),
        'Recall': recall score(y true, y pred, average='macro'),
        'F1 Score': f1 score(y true, y pred, average='macro')
    }
metrics rf = get metrics(y test, y pred)
metrics_hybrid = get_metrics(y_test, hybrid_y_pred)
# □ Bar Plot of Metrics
plt.figure(figsize=(10, 6))
labels = list(metrics rf.keys())
rf values = list(metrics rf.values())
hybrid values = list(metrics hybrid.values())
x = range(len(labels))
plt.bar(x, rf values, width=0.4, label='RF Model', align='center')
plt.bar([i + 0.4 \text{ for } i \text{ in } x], hybrid values, width=0.4, label='Hybrid
RF Model', align='center')
plt.xticks([i + 0.2 for i in x], labels)
plt.ylabel('Score')
plt.title('Performance Metrics Comparison')
plt.legend()
plt.tight layout()
plt.show()
```

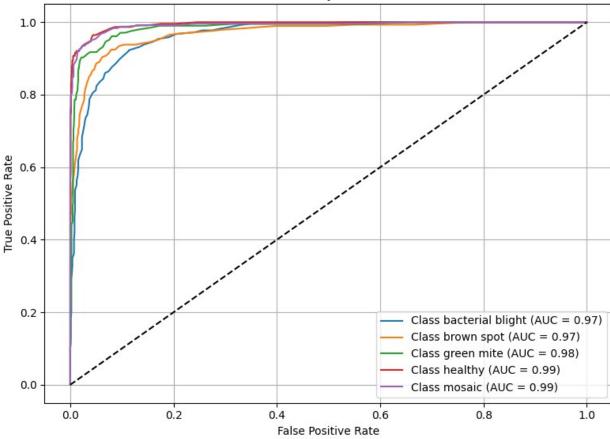


```
from sklearn.preprocessing import label binarize
# Binarize the labels for multiclass
classes = np.unique(y_test)
y_test_bin = label_binarize(y_test, classes=classes)
# □ RF Model ROC Curve
y pred proba = rf model.predict proba(rf X test)
plt.figure(figsize=(8, 6))
for i, class label in enumerate(classes):
    fpr, tpr, = roc curve(y test bin[:, i], y pred proba[:, i])
    roc auc = auc(fpr, tpr)
    plt.plot(fpr, tpr, label=f'Class {class_label} (AUC =
{roc auc:.2f})')
plt.plot([0, 1], [0, 1], 'k--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('ROC Curve - RF Model')
plt.legend()
plt.grid(True)
plt.tight_layout()
plt.show()
```



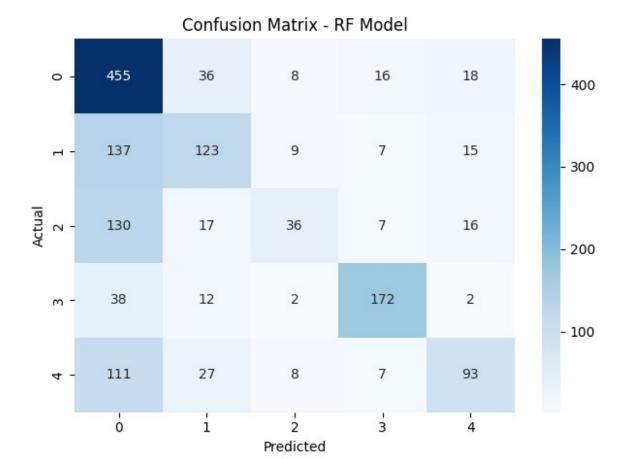
```
hybrid y pred proba = hybrid rf model.predict proba(X test)
plt.figure(figsize=(8, 6))
for i, class label in enumerate(classes):
    fpr, tpr, _ = roc_curve(y_test_bin[:, i], hybrid_y_pred_proba[:,
i])
    roc auc = auc(fpr, tpr)
    plt.plot(fpr, tpr, label=f'Class {class_label} (AUC =
{roc_auc:.2f})')
plt.plot([0, 1], [0, 1], 'k--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('ROC Curve - Hybrid RF Model')
plt.legend()
plt.grid(True)
plt.tight_layout()
plt.show()
```

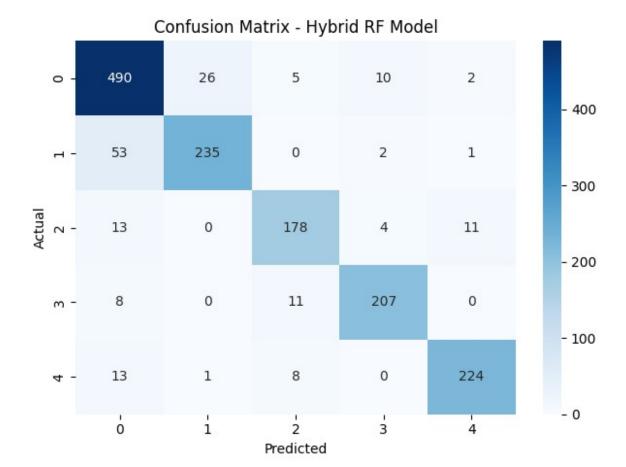




```
# Confusion Matrices
def plot_conf_matrix(y_true, y_pred, title):
    cm = confusion_matrix(y_true, y_pred)
    sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')
    plt.title(title)
    plt.xlabel('Predicted')
    plt.ylabel('Actual')
    plt.tight_layout()
    plt.show()

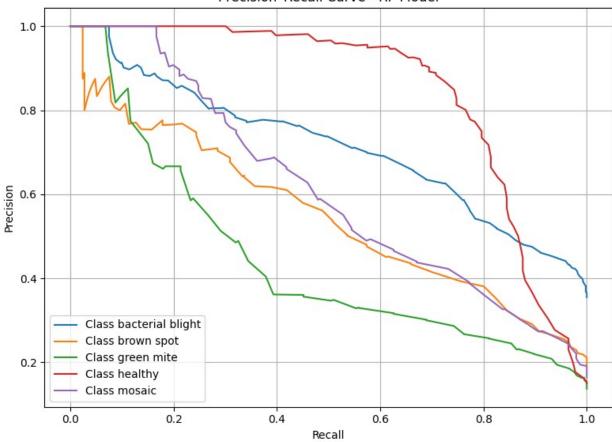
plot_conf_matrix(y_test, y_pred, 'Confusion Matrix - RF Model')
plot_conf_matrix(y_test, hybrid_y_pred, 'Confusion Matrix - Hybrid RF Model')
```

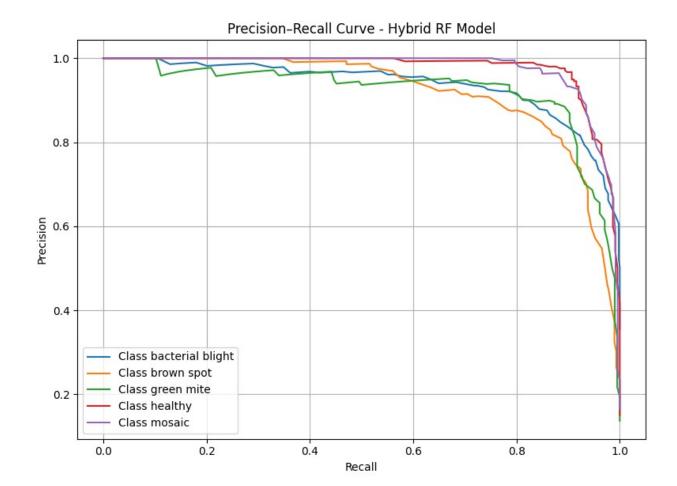




```
from sklearn.preprocessing import label binarize
# Binarize true labels for multiclass
classes = np.unique(y test)
y test bin = label binarize(y test, classes=classes)
y pred proba = rf model.predict proba(rf X test)
plt.figure(figsize=(8, 6))
for i, class_label in enumerate(classes):
    precision, recall, _ = precision_recall_curve(y_test_bin[:, i],
y_pred_proba[:, i])
    plt.plot(recall, precision, label=f'Class {class label}')
plt.xlabel('Recall')
plt.ylabel('Precision')
plt.title('Precision-Recall Curve - RF Model')
plt.legend()
plt.grid(True)
plt.tight layout()
plt.show()
```

Precision-Recall Curve - RF Model





Saving Model for Development

```
import joblib

joblib.dump(hybrid_rf_model, "hybrid_model.pkl")
['hybrid_model.pkl']
```