# Mount Google Drive and Set Path

```
from google.colab import drive
drive.mount('/content/drive')
```

Drive already mounted at /content/drive; to attempt to forcibly remount, call drive.mount(

```
import os
import numpy as np
import matplotlib.pyplot as plt
from tensorflow.keras.preprocessing.image import ImageDataGenerator
from collections import Counter

# Define path
base_dir = '/content/drive/MyDrive/Assessment_1/dataset'
train_dir = os.path.join(base_dir, 'train')
test_dir = os.path.join(base_dir, 'test')
```

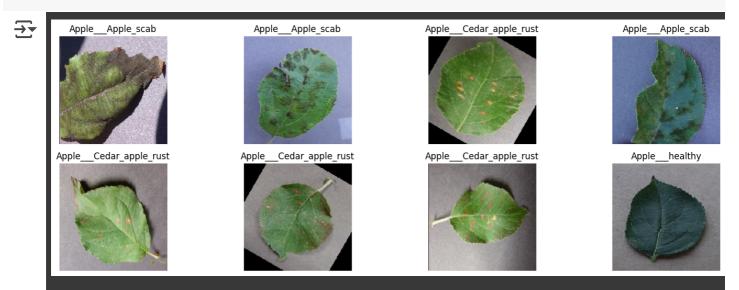
Set Parameters and Create Data Generators

```
IMG_SIZE = (224, 224)
BATCH_SIZE = 32
# No augmentation
train_datagen = ImageDataGenerator(rescale=1./255)
test_datagen = ImageDataGenerator(rescale=1./255)
# Load images
train_generator = train_datagen.flow_from_directory(
    train_dir,
    target_size=IMG_SIZE,
    batch_size=BATCH_SIZE,
    class_mode='categorical'
test_generator = test_datagen.flow_from_directory(
    test_dir,
    target_size=IMG_SIZE,
    batch_size=BATCH_SIZE,
    class_mode='categorical',
    shuffle=False
)
```

Found 6218 images belonging to 4 classes. Found 1553 images belonging to 4 classes.

### Visualize Sample Images

```
def plot_images(generator, class_indices, num_images=8):
    class_names = list(class_indices.keys())
    x_batch, y_batch = next(generator)
    plt.figure(figsize=(15, 5))
    for i in range(num_images):
        ax = plt.subplot(2, 4, i + 1)
        plt.imshow(x_batch[i])
        label_index = np.argmax(y_batch[i])
        plt.title(class_names[label_index])
        plt.axis("off")
    plt.tight_layout()
    plt.show()
```



Dataset Analysis

```
# Count total images per class
def count_images(directory):
    class_counts = {}
    for class_name in os.listdir(directory):
        class_path = os.path.join(directory, class_name)
        if os.path.isdir(class_path):
            class_counts[class_name] = len(os.listdir(class_path))
    return class_counts

train_counts = count_images(train_dir)
test_counts = count_images(test_dir)

print("Train Class Distribution:", train_counts)
print("Total training images:", sum(train_counts.values()))
print("Total testing images:", sum(test_counts.values()))
```

```
Train Class Distribution: {'Apple__healthy': 1607, 'Apple__Cedar_apple_rust': 1408, 'Apple_ Test Class Distribution: {'Apple__healthy': 401, 'Apple__Cedar_apple_rust': 352, 'Apple_ Total training images: 6218
Total testing images: 1553
```

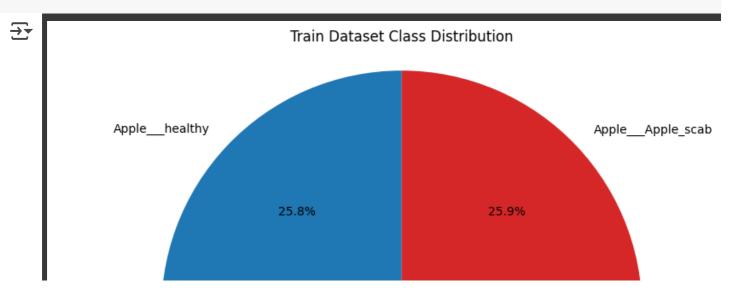
We split the dataset into training and testing sets in a folder-based format. A standard 80/20 split is often applied to ensure sufficient data for training while retaining a portion for unbiased evaluation.**bold text** 

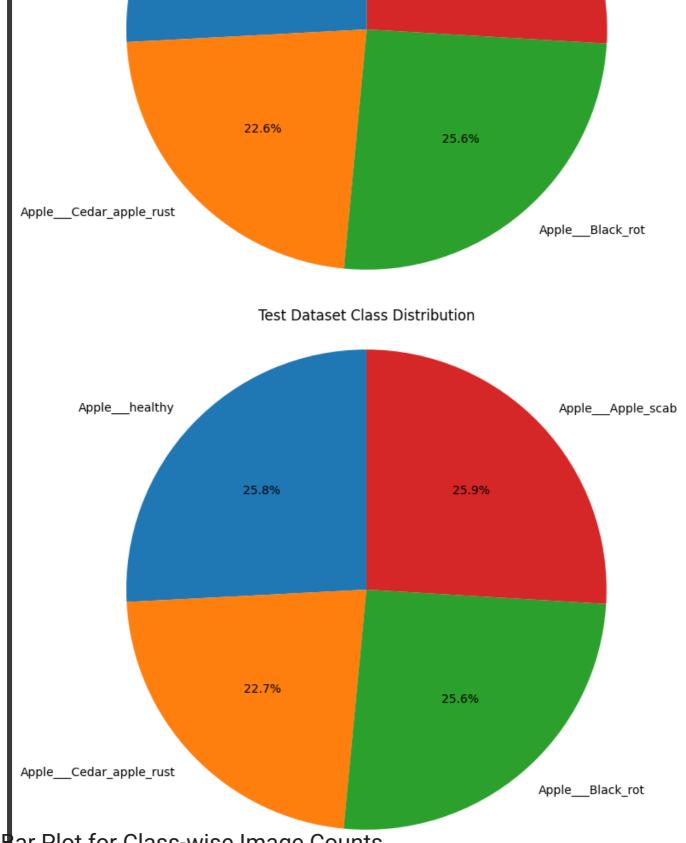
### Class Distribution Histogram

```
import matplotlib.pyplot as plt

def plot_pie_chart(counts, title):
    labels = list(counts.keys())
    sizes = list(counts.values())
    plt.figure(figsize=(8, 8))
    plt.pie(sizes, labels=labels, autopct='%1.1f%%', startangle=90)
    plt.title(title)
    plt.axis('equal')  # Equal aspect ratio ensures that pie is drawn as a circle.
    plt.show()

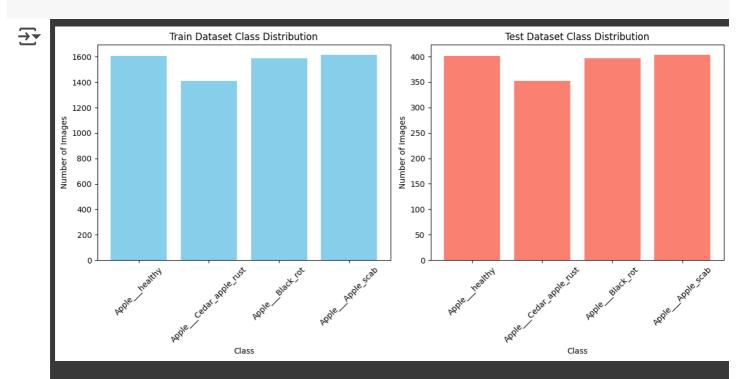
plot_pie_chart(train_counts, "Train Dataset Class Distribution")
plot_pie_chart(test_counts, "Test Dataset Class Distribution")
```







```
# Plot class distribution using bar charts
plt.figure(figsize=(12, 6))
# Train dataset bar plot
plt.subplot(1, 2, 1)
plt.bar(train_counts.keys(), train_class_counts, color='skyblue')
plt.title("Train Dataset Class Distribution")
plt.xlabel("Class")
plt.ylabel("Number of Images")
plt.xticks(rotation=45)
# Test dataset bar plot
plt.subplot(1, 2, 2)
plt.bar(test_counts.keys(), test_class_counts, color='salmon')
plt.title("Test Dataset Class Distribution")
plt.xlabel("Class")
plt.ylabel("Number of Images")
plt.xticks(rotation=45)
plt.tight_layout()
plt.show()
```



Filter A filter (or feature detector) refers to the set of values (weights) that are applied to the input data during the convolution process.

Kernel A kernel is the actual matrix (the set of weights) that is used to perform the convolution operation on the input image. It is typically smaller than the image (e.g., a 3x3 or 5x5 matrix)

### Build the Baseline CNN Architecture

Convolutional Layer: Extracts local features using filters. Activation Layer (ReLU): Introduces non-linearity. Pooling Layer: Reduces spatial dimensions and retains essential features. Flatten Layer: Converts 2D feature maps into 1D vectors. Fully Connected Layer: Learns high-level features for classification or regression. Dropout Layer: Prevents overfitting by randomly deactivating neurons. Batch Normalization: Normalizes layer inputs for more stable training.

```
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Conv2D, MaxPooling2D, Flatten, Dense, Dropout
from tensorflow.keras.optimizers import Adam
# Build the model
model = Sequential() #This initializes a Sequential model
# Convolutional Layer 1
model.add(Conv2D(32, (3, 3), activation='relu', input_shape=(224, 224, 3))) #32 is number of fi
model.add(MaxPooling2D(pool_size=(2, 2)))
# Convolutional Layer 2
model.add(Conv2D(64, (3, 3), activation='relu')) #(3, 3): The size of each filter (kernel)
model.add(MaxPooling2D(pool_size=(2, 2)))
# Convolutional Layer 3
model.add(Conv2D(128, (3, 3), activation='relu'))
model.add(MaxPooling2D(pool_size=(2, 2)))
# Flatten layer
model.add(Flatten())
#flatten the multi-dimensional output from the previous layer (typically the 2D feature maps a
# Fully connected layers
model.add(Dense(512, activation='relu')) #learns the feature
model.add(Dropout(0.5)) # Dropout to prevent overfitting
model.add(Dense(256, activation='relu'))
model.add(Dense(4, activation='softmax')) # 4 classes
# Model summary
model.summary()
```

### **→**

#### Model: "sequential\_1"

Layer (type)	Output Shape	Param #
conv2d_3 (Conv2D)	(None, 222, 222, 32)	896
max_pooling2d_3 (MaxPooling2D)	(None, 111, 111, 32)	0
conv2d_4 (Conv2D)	(None, 109, 109, 64)	18,496
max_pooling2d_4 (MaxPooling2D)	(None, 54, 54, 64)	0
conv2d_5 (Conv2D)	(None, 52, 52, 128)	73,856
max_pooling2d_5 (MaxPooling2D)	(None, 26, 26, 128)	0
flatten_1 (Flatten)	(None, 86528)	0
dense_3 (Dense)	(None, 512)	44,302,848
dropout_1 (Dropout)	(None, 512)	0
dense_4 (Dense)	(None, 256)	131,328
dense_5 (Dense)	(None, 4)	1,028

Total params: 44,528,452 (169.86 MB)
Trainable params: 44,528,452 (169.86 MB)
Non-trainable params: 0 (0.00 B)

# Compile the Model

```
# Compile the model
model.compile(optimizer=Adam(), loss='categorical_crossentropy', metrics=['accuracy'])
```

#### Train the Model

```
history = model.fit(
    train_generator,
    epochs=10, # Adjust as necessary
    validation_data=test_generator,
    verbose=1 #This controls how much information is shown during training.
)

# Plotting training & validation loss
plt.figure(figsize=(8, 6))
plt.plot(history.history['loss'], label='Training Loss')
plt.plot(history.history['val_loss'], label='Validation Loss')
plt.title('Training vs Validation Loss')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()
plt.show()
```



### Evaluate the Model

```
from sklearn.metrics import classification_report

# Get predictions
predictions = model.predict(test_generator)
y_pred = np.argmax(predictions, axis=1)  #picks the class with the highest probability for each
y_true = test_generator.classes

# Print classification report
print(classification_report(y_true, y_pred, target_names=test_generator.class_indices.keys()))
```

<b>→</b>	49/49 ————	<b>9/49</b> — <b>7s</b> 141ms/step			
		precision	recall	f1-score	support
	AppleApple_scab	0.98	0.90	0.94	403
	AppleBlack_rot	0.93	0.99	0.96	397
	AppleCedar_apple_rust	0.97	1.00	0.98	352
	Applehealthy	0.98	0.97	0.97	401
	2.6.0112.01			0.06	1550
	accuracy			0.96	1553
	macro avg	0.96	0.97	0.96	1553
	weighted avg	0.96	0.96	0.96	1553

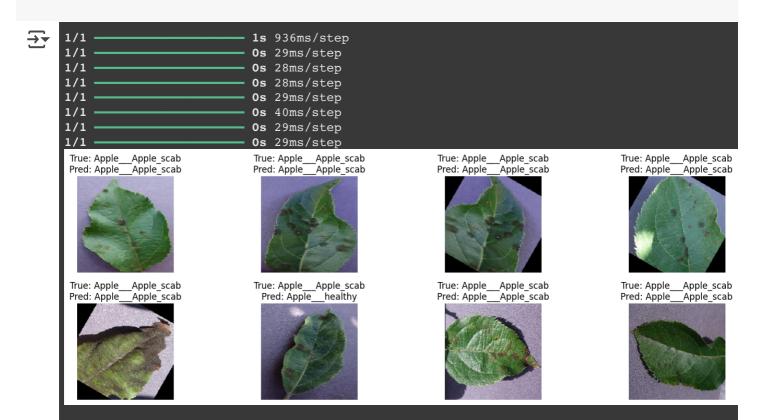
Precision measures the accuracy of positive predictions. Recall measures how well the model identifies actual positive instances. F1-Score is the harmonic mean of precision and recall.

### Visualize Predictions

```
def plot_predictions(generator, model, num_images=8):
    class_names = list(generator.class_indices.keys())
    x_batch, y_batch = next(generator)
    plt.figure(figsize=(15, 5))

for i in range(num_images):
        ax = plt.subplot(2, 4, i + 1)
        plt.imshow(x_batch[i])
        predicted_class = class_names[np.argmax(model.predict(x_batch[i:i+1]))]
        true_class = class_names[np.argmax(y_batch[i])]
        plt.title(f"True: {true_class}\nPred: {predicted_class}")
        plt.axis('off')

plt.tight_layout()
    plt.show()
```



batch normalization is normalizes the inputs to each layer of the network. This means it adjusts the data so that each input has a mean of 0 and a variance of 1. This makes the training process smoother and faster.

### Build the Deeper Model

```
from tensorflow.keras.layers import BatchNormalization

# Build deeper model
model_deeper = Sequential()
```

```
# Convolutional Layer 1
model_deeper.add(Conv2D(64, (3, 3), activation='relu', input_shape=(224, 224, 3)))
model_deeper.add(BatchNormalization()) # Batch normalization
model_deeper.add(MaxPooling2D(pool_size=(2, 2)))
# Convolutional Layer 2
model_deeper.add(Conv2D(128, (3, 3), activation='relu'))
model_deeper.add(BatchNormalization())
model_deeper.add(MaxPooling2D(pool_size=(2, 2)))
# Convolutional Layer 3
model_deeper.add(Conv2D(256, (3, 3), activation='relu'))
model_deeper.add(BatchNormalization())
model_deeper.add(MaxPooling2D(pool_size=(2, 2)))
# Convolutional Layer 4 (Added layer)
model_deeper.add(Conv2D(512, (3, 3), activation='relu'))
model_deeper.add(BatchNormalization())
model_deeper.add(MaxPooling2D(pool_size=(2, 2)))
# Convolutional Layer 5 (Added layer)
model_deeper.add(Conv2D(512, (3, 3), activation='relu'))
model_deeper.add(BatchNormalization())
model_deeper.add(MaxPooling2D(pool_size=(2, 2)))
# Flatten layer
model_deeper.add(Flatten())
# Fully connected layers
model_deeper.add(Dense(1024, activation='relu'))
model_deeper.add(Dropout(0.5)) # Dropout for regularization
model_deeper.add(Dense(512, activation='relu'))
model_deeper.add(Dense(4, activation='softmax')) # 4 classes
#decreasing will neurons to 64, you are decreasing the model's capacity.
# Model summary
model_deeper.summary()
```

/usr/local/lib/python3.11/dist-packages/keras/src/layers/convolutional/base\_conv.py:107: U
 super().\_\_init\_\_(activity\_regularizer=activity\_regularizer, \*\*kwargs)
Model: "sequential\_2"

Layer (type)	Output Shape	Param #
conv2d_6 (Conv2D)	(None, 222, 222, 64)	1,792
batch_normalization (BatchNormalization)	(None, 222, 222, 64)	256
<pre>max_pooling2d_6 (MaxPooling2D)</pre>	(None, 111, 111, 64)	0
conv2d_7 (Conv2D)	(None, 109, 109, 128)	73,856
<pre>batch_normalization_1 (BatchNormalization)</pre>	(None, 109, 109, 128)	512
<pre>max_pooling2d_7 (MaxPooling2D)</pre>	(None, 54, 54, 128)	0
conv2d_8 (Conv2D)	(None, 52, 52, 256)	295,168
<pre>batch_normalization_2 (BatchNormalization)</pre>	(None, 52, 52, 256)	1,024
max_pooling2d_8 (MaxPooling2D)	(None, 26, 26, 256)	Ø
conv2d_9 (Conv2D)	(None, 24, 24, 512)	1,180,160
<pre>batch_normalization_3 (BatchNormalization)</pre>	(None, 24, 24, 512)	2,048
<pre>max_pooling2d_9 (MaxPooling2D)</pre>	(None, 12, 12, 512)	0
conv2d_10 (Conv2D)	(None, 10, 10, 512)	2,359,808
batch_normalization_4 (BatchNormalization)	(None, 10, 10, 512)	2,048
max_pooling2d_10 (MaxPooling2D)	(None, 5, 5, 512)	0
flatten_2 (Flatten)	(None, 12800)	0
dense_6 (Dense)	(None, 1024)	13,108,224
dropout_2 (Dropout)	(None, 1024)	0
dense_7 (Dense)	(None, 512)	524,800
dense_8 (Dense)	(None, 4)	2,052

Total params: 17,551,748 (66.95 MB)
Trainable params: 17,548,804 (66.94 MB)
Non-trainable params: 2,944 (11.50 KB)

# Compile the Model

```
# Compile the deeper model
model_deeper.compile(optimizer=Adam(), loss='categorical_crossentropy', metrics=['accuracy'])
```

# Train the Deeper Model

```
history_deeper = model_deeper.fit(
    train_generator,
    epochs=10, # Adjust as necessary
    validation_data=test_generator,
    verbose=1 #This controls how much information is shown during training
)

# Plotting training & validation loss for deeper model
plt.figure(figsize=(8, 6))
plt.plot(history_deeper.history['loss'], label='Training Loss')
plt.plot(history_deeper.history['val_loss'], label='Validation Loss')
plt.title('Training vs Validation Loss (Deeper Model)')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()
plt.show()
```



### Evaluate the Deeper Model

```
# Get predictions from deeper model
predictions_deeper = model_deeper.predict(test_generator)
y_pred_deeper = np.argmax(predictions_deeper, axis=1)

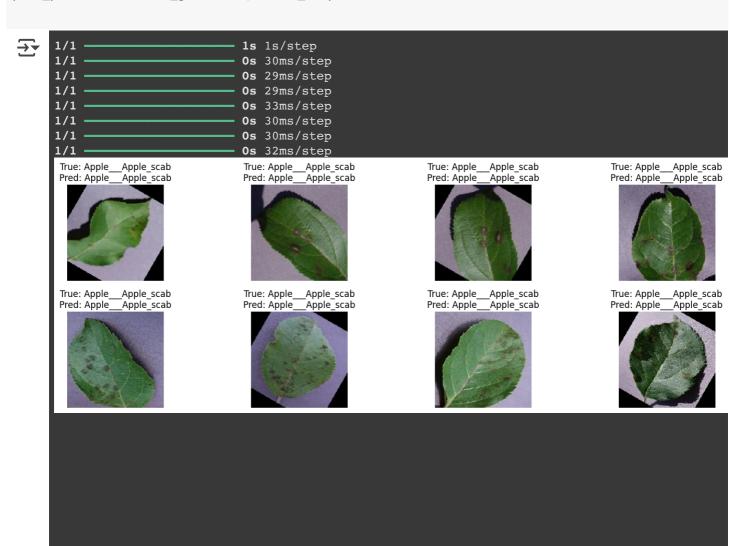
# Print classification report
from sklearn.metrics import classification_report

print(classification_report(y_true, y_pred_deeper, target_names=test_generator.class_indices.kg
```

<b>→</b>	<b>49/49</b> — <b>8s</b> 144ms/step				
_		precision	recall	f1-score	support
	AppleApple_scab	0.75	0.95	0.84	403
	AppleBlack_rot	0.96	0.99	0.98	397
	AppleCedar_apple_rust	0.84	0.82	0.83	352
	Applehealthy	0.97	0.70	0.81	401
	accuracy			0.87	1553
	macro avg	0.88	0.87	0.87	1553
	weighted avg	0.88	0.87	0.87	1553

### Visualize Predictions from the Deeper Model

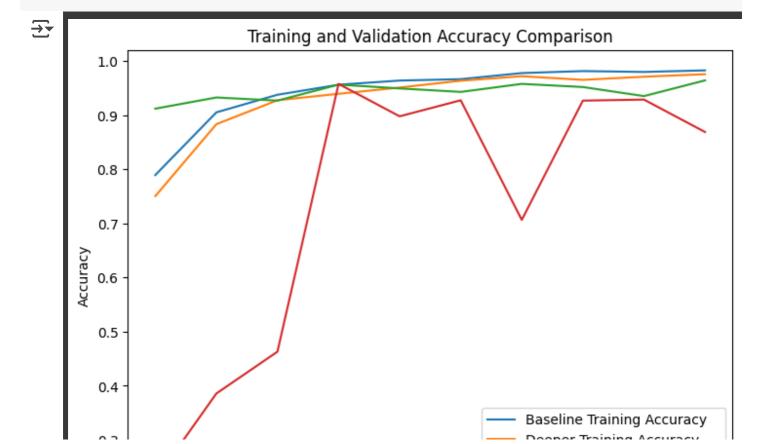
plot\_predictions(test\_generator, model\_deeper)

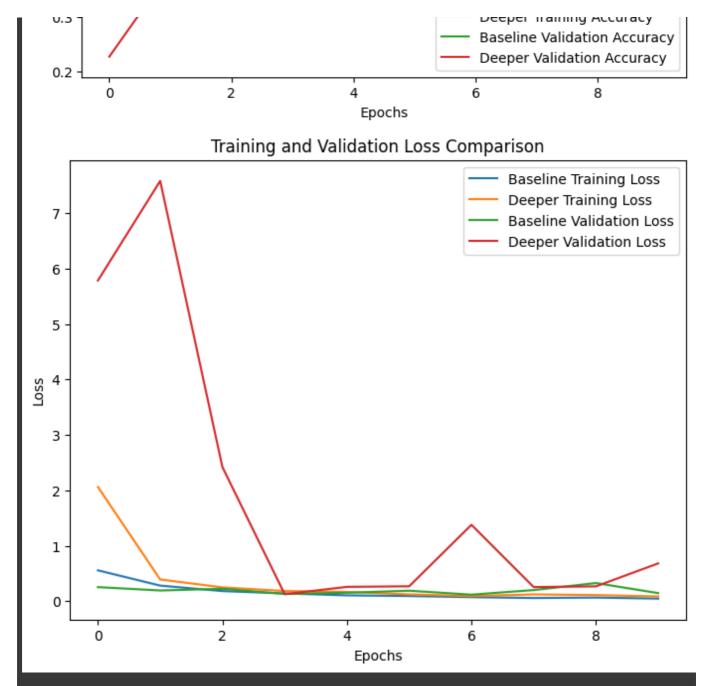


# Comparative Analysis of Baseline vs. Deeper Model

### Model Performance Comparison

```
# Compare accuracy and loss for baseline and deeper models
baseline_history = history.history
deeper_history = history_deeper.history
# Plot accuracy comparison
plt.figure(figsize=(8, 6))
plt.plot(baseline_history['accuracy'], label='Baseline Training Accuracy')
plt.plot(deeper_history['accuracy'], label='Deeper Training Accuracy')
plt.plot(baseline_history['val_accuracy'], label='Baseline Validation Accuracy')
plt.plot(deeper_history['val_accuracy'], label='Deeper Validation Accuracy')
plt.title('Training and Validation Accuracy Comparison')
plt.xlabel('Epochs')
plt.ylabel('Accuracy')
plt.legend()
plt.show()
# Plot loss comparison
plt.figure(figsize=(8, 6))
plt.plot(baseline_history['loss'], label='Baseline Training Loss')
plt.plot(deeper_history['loss'], label='Deeper Training Loss')
plt.plot(baseline_history['val_loss'], label='Baseline Validation Loss')
plt.plot(deeper_history['val_loss'], label='Deeper Validation Loss')
plt.title('Training and Validation Loss Comparison')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()
plt.show()
```





### **Computational Efficiency Comparison**

### Optimizer Comparison (SGD vs Adam)

```
# Compile model with SGD optimizer
model_deeper.compile(optimizer='sgd', loss='categorical_crossentropy', metrics=['accuracy'])
# Train with SGD learning rate is 0.01 meanwhile adam is 0.0005
history_sgd = model_deeper.fit(
    train_generator,
    epochs=10, #one complete pass through the entire training dataset during the training procevalidation_data=test_generator,
    verbose=1
)
# Plot SGD vs Adam Loss
plt.figure(figsize=(8, 6))
plt.plot(history_sgd.history['loss'], label='SGD Training Loss')
```

```
plt.plot(deeper_history['loss'], label='Adam Training Loss')
plt.plot(history_sgd.history['val_loss'], label='SGD Validation Loss')
plt.plot(deeper_history['val_loss'], label='Adam Validation Loss')
plt.title('SGD vs Adam Loss Comparison')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()
plt.show()
```



### Loading and Adapting a Pre-Trained Model

```
from tensorflow.keras.models import Model
from tensorflow.keras.optimizers import Adam
# Load VGG16 with pre-trained ImageNet weights, without the top fully connected layers
base_model = VGG16(weights='imagenet', include_top=False, input_shape=(224, 224, 3))
#will only retain the convolutional layers and pooling layers, which are responsible for featu
# Freeze the base model (so the weights won't be updated during training)
for layer in base_model.layers:
    layer.trainable = False
# Add a custom top layer for your classification task
x = base_model.output
x = GlobalAveragePooling2D()(x) # Pooling layer to reduce dimensionality
x = Dense(1024, activation='relu')(x)
x = Dropout(0.5)(x) # Regularization
x = Dense(4, activation='softmax')(x) # 4 classes for classification
# Create the final model
model_transfer = Model(inputs=base_model.input, outputs=x)
# Compile the model
model_transfer.compile(optimizer=Adam(), loss='categorical_crossentropy', metrics=['accuracy']
model_transfer.summary()
#Adam(learning_rate=0.0005)
```

58889256/58889256 - 4s Ous/step

Model: "functional\_42"

	1	- "
Layer (type)	Output Shape	Param #
input_layer_3 (InputLayer)	(None, 224, 224, 3)	0
block1_conv1 (Conv2D)	(None, 224, 224, 64)	1,792
block1_conv2 (Conv2D)	(None, 224, 224, 64)	36,928
block1_pool (MaxPooling2D)	(None, 112, 112, 64)	0
block2_conv1 (Conv2D)	(None, 112, 112, 128)	73,856
block2_conv2 (Conv2D)	(None, 112, 112, 128)	147,584
block2_pool (MaxPooling2D)	(None, 56, 56, 128)	0
block3_conv1 (Conv2D)	(None, 56, 56, 256)	295,168
block3_conv2 (Conv2D)	(None, 56, 56, 256)	590,080
block3_conv3 (Conv2D)	(None, 56, 56, 256)	590,080
block3_pool (MaxPooling2D)	(None, 28, 28, 256)	Ø
block4_conv1 (Conv2D)	(None, 28, 28, 512)	1,180,160
block4_conv2 (Conv2D)	(None, 28, 28, 512)	2,359,808
block4_conv3 (Conv2D)	(None, 28, 28, 512)	2,359,808
block4_pool (MaxPooling2D)	(None, 14, 14, 512)	Ø
block5_conv1 (Conv2D)	(None, 14, 14, 512)	2,359,808
block5_conv2 (Conv2D)	(None, 14, 14, 512)	2,359,808
block5_conv3 (Conv2D)	(None, 14, 14, 512)	2,359,808
block5_pool (MaxPooling2D)	(None, 7, 7, 512)	0
global_average_pooling2d (GlobalAveragePooling2D)	(None, 512)	0
dense_9 (Dense)	(None, 1024)	525,312
dropout_3 (Dropout)	(None, 1024)	0
dense_10 (Dense)	(None, 4)	4,100

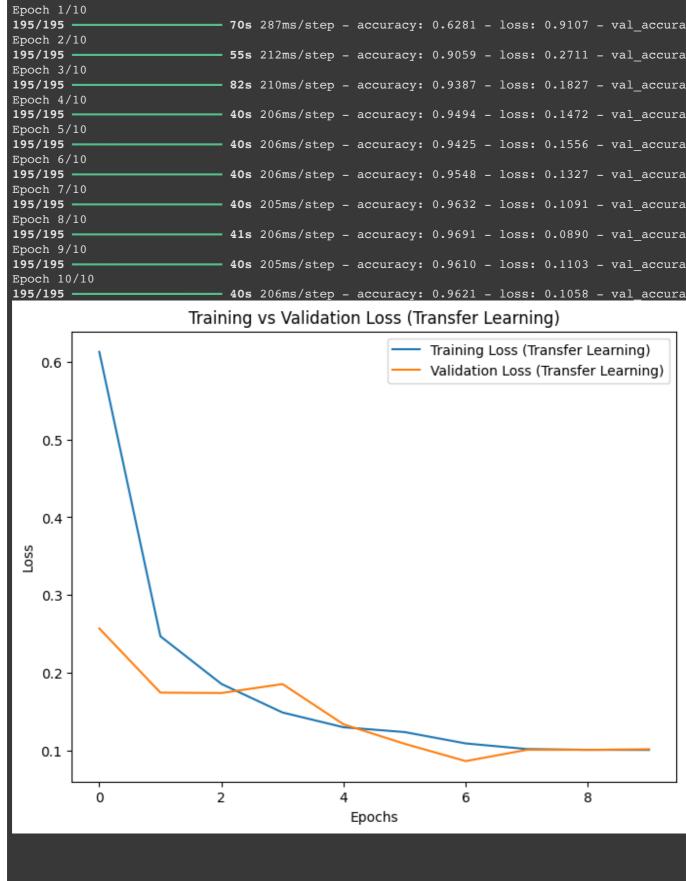
Total params: 15,244,100 (58.15 MB)
Trainable params: 529,412 (2.02 MB)
Non-trainable params: 14,714,688 (56.13 MB)

# **Training Strategies**

```
# Train only the custom top layers
history_transfer = model_transfer.fit(
    train_generator,
    epochs=10, # You can adjust this based on training time
```

```
validation_data=test_generator,
   verbose=1 #This controls the level of logging output during training
)

# Plot training vs validation loss for transfer learning
plt.figure(figsize=(8, 6))
plt.plot(history_transfer.history['loss'], label='Training Loss (Transfer Learning)')
plt.plot(history_transfer.history['val_loss'], label='Validation Loss (Transfer Learning)')
plt.title('Training vs Validation Loss (Transfer Learning)')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()
plt.show()
```



### Model Evaluation and Prediction

```
# Get predictions from the transfer learning model
predictions_transfer = model_transfer.predict(test_generator)
y_pred_transfer = np.argmax(predictions_transfer, axis=1)

# Print classification report for transfer learning model
print(classification_report(y_true, y_pred_transfer, target_names=test_generator.class_indices)
```

<b>→</b>	49/49 ————	<b>– 10s</b> 184ms/	step		
<u> </u>		precision	recall	f1-score	support
	AppleApple_scab	0.99	0.87	0.92	403
	AppleBlack_rot	0.97	0.99	0.98	397
	AppleCedar_apple_rust	0.93	0.99	0.96	352
	Applehealthy	0.94	0.99	0.96	401
	accuracy			0.96	1553
	macro avg	0.96	0.96	0.96	1553
	weighted avg	0.96	0.96	0.96	1553

### ROC curve (Receiver Operating Characteristic curve)

```
from sklearn.metrics import roc curve, auc
from sklearn.preprocessing import label_binarize
# Binarize the true labels (multi-class one-hot encoding)
y_true_binarized = label_binarize(y_true, classes=[0, 1, 2, 3])
# Function to plot the ROC curve for each class
def plot_roc_curve(y_true, y_pred_prob, model_name):
    n_classes = y_true.shape[1] # Number of classes
    fpr, tpr, auc_scores = {}, {}, {}
    # Plot ROC curve for each class
    plt.figure(figsize=(10, 8))
    for i in range(n classes):
        fpr[i], tpr[i], _ = roc_curve(y_true[:, i], y_pred_prob[:, i]) # Get FPR, TPR for clas
        auc_scores[i] = auc(fpr[i], tpr[i]) # Calculate AUC for each class
        plt.plot(fpr[i], tpr[i], label=f'Class {i} (AUC = {auc_scores[i]:.2f})')
    # Plot a random classifier (diagonal line)
    plt.plot([0, 1], [0, 1], 'k--', label='Random Classifier')
    plt.title(f'ROC Curve for {model_name}')
    plt.xlabel('False Positive Rate')
    plt.ylabel('True Positive Rate')
    plt.legend(loc='best')
    plt.show()
# Get predicted probabilities for each model (probabilities for each class)
y_pred_baseline_prob = model.predict(test_generator)
y_pred_deeper_prob = model_deeper.predict(test_generator)
y_pred_transfer_prob = model_transfer.predict(test_generator)
# Plot ROC curve for each model
plot_roc_curve(y_true_binarized, y_pred_baseline_prob, 'Baseline Model')
plot_roc_curve(y_true_binarized, y_pred_deeper_prob, 'Deeper Model')
plot_roc_curve(y_true_binarized, y_pred_transfer_prob, 'Transfer Learning Model')
```

