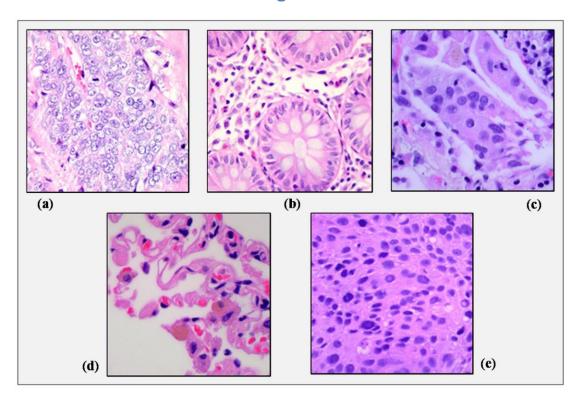
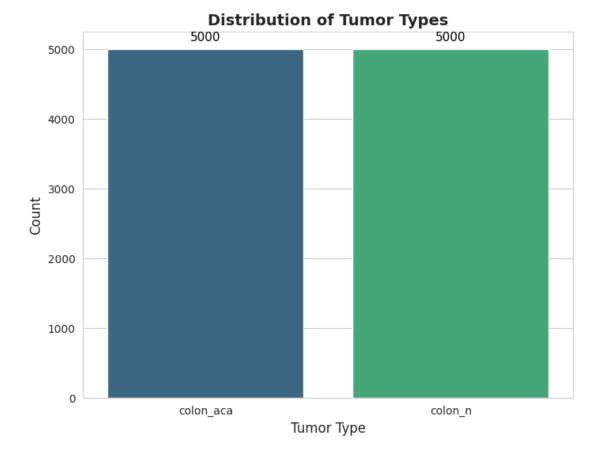
MRANet with SE Block for Lung Colon Cancer Classification



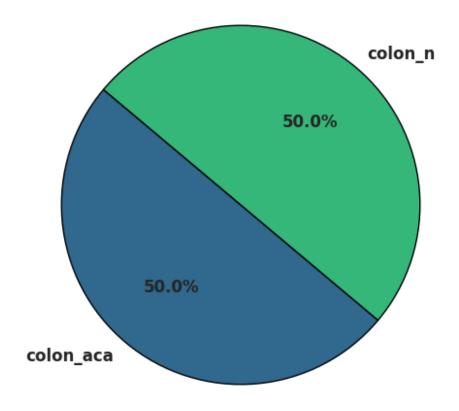
```
import numpy as np
import pandas as pd
base_path = "/kaggle/input/lung-and-colon-cancer-histopathological-
images/lung_colon_image_set/colon_image_sets"
categories = ["colon aca", "colon n"]
image_paths = []
labels = []
for category in categories:
    category_path = os.path.join(base_path, category)
    for image_name in os.listdir(category_path):
        image_path = os.path.join(category_path, image_name)
        image_paths.append(image_path)
        labels.append(category)
df = pd.DataFrame({
    "image_path": image_paths,
    "label": labels
})
df.head()
```

```
image path
                                                          label
0 /kaggle/input/lung-and-colon-cancer-histopatho...
                                                      colon aca
1 /kaggle/input/lung-and-colon-cancer-histopatho...
                                                      colon_aca
2 /kaggle/input/lung-and-colon-cancer-histopatho...
                                                      colon aca
3 /kaggle/input/lung-and-colon-cancer-histopatho...
                                                      colon_aca
4 /kaggle/input/lung-and-colon-cancer-histopatho...
                                                      colon aca
df.tail()
                                             image path
                                                           label
      /kaggle/input/lung-and-colon-cancer-histopatho...
9995
                                                         colon n
9996
      /kaggle/input/lung-and-colon-cancer-histopatho...
                                                         colon_n
9997
      /kaggle/input/lung-and-colon-cancer-histopatho...
                                                         colon n
      /kaggle/input/lung-and-colon-cancer-histopatho...
9998
                                                         colon_n
     /kaggle/input/lung-and-colon-cancer-histopatho...
9999
                                                         colon n
df.shape
(10000, 2)
df.columns
Index(['image_path', 'label'], dtype='object')
df.duplicated().sum()
0
df.isnull().sum()
image path
              0
label
              0
dtype: int64
df.info()
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 10000 entries, 0 to 9999
Data columns (total 2 columns):
                Non-Null Count Dtype
     Column
                 -----
 0
     image path 10000 non-null object
 1
     label
                 10000 non-null object
dtypes: object(2)
memory usage: 156.4+ KB
df['label'].unique()
array(['colon aca', 'colon n'], dtype=object)
df['label'].value_counts()
```

```
label
colon aca
            5000
colon n
            5000
Name: count, dtype: int64
import seaborn as sns
import matplotlib.pyplot as plt
sns.set_style("whitegrid")
fig, ax = plt.subplots(figsize=(8, 6))
sns.countplot(data=df, x="label", palette="viridis", ax=ax)
ax.set_title("Distribution of Tumor Types", fontsize=14, fontweight='bold')
ax.set_xlabel("Tumor Type", fontsize=12)
ax.set_ylabel("Count", fontsize=12)
for p in ax.patches:
    ax.annotate(f'{int(p.get_height())}',
                (p.get_x() + p.get_width() / 2., p.get_height()),
                ha='center', va='bottom', fontsize=11, color='black',
                xytext=(0, 5), textcoords='offset points')
plt.show()
label_counts = df["label"].value_counts()
fig, ax = plt.subplots(figsize=(8, 6))
colors = sns.color_palette("viridis", len(label_counts))
ax.pie(label counts, labels=label counts.index, autopct='%1.1f%%',
       startangle=140, colors=colors, textprops={'fontsize': 12, 'weight':
'bold'},
       wedgeprops={'edgecolor': 'black', 'linewidth': 1})
ax.set_title("Distribution of Tumor Types - Pie Chart", fontsize=14,
fontweight='bold')
plt.show()
```



Distribution of Tumor Types - Pie Chart



```
import cv2
num_images = 5
plt.figure(figsize=(15, 12))
for i, category in enumerate(categories):
    category_images = df[df['label'] ==
category]['image_path'].iloc[:num_images]

for j, img_path in enumerate(category_images):
    img = cv2.imread(img_path)
    img = cv2.cvtColor(img, cv2.COLOR_BGR2RGB)

    plt.subplot(len(categories), num_images, i * num_images + j + 1)
    plt.imshow(img)
    plt.axis('off')
    plt.title(category)
```

```
plt.tight_layout()
plt.show()
from sklearn.preprocessing import LabelEncoder
label_encoder = LabelEncoder()
df['category_encoded'] = label_encoder.fit_transform(df['label'])
df = df[['image_path', 'category_encoded']]
from imblearn.over_sampling import RandomOverSampler
ros = RandomOverSampler(random_state=42)
X_resampled, y_resampled = ros.fit_resample(df[['image_path']],
df['category_encoded'])
df_resampled = pd.DataFrame(X_resampled, columns=['image_path'])
df_resampled['category_encoded'] = y_resampled
print("\nClass distribution after oversampling:")
print(df_resampled['category_encoded'].value_counts())
Class distribution after oversampling:
category_encoded
```

5000

5000

df_resampled

Name: count, dtype: int64

0

```
image_path category_encoded
      /kaggle/input/lung-and-colon-cancer-histopatho...
0
1
      /kaggle/input/lung-and-colon-cancer-histopatho...
                                                                         0
2
      /kaggle/input/lung-and-colon-cancer-histopatho...
                                                                         0
3
      /kaggle/input/lung-and-colon-cancer-histopatho...
                                                                         0
4
      /kaggle/input/lung-and-colon-cancer-histopatho...
                                                                         0
                                                                       . . .
9995
     /kaggle/input/lung-and-colon-cancer-histopatho...
                                                                         1
9996 /kaggle/input/lung-and-colon-cancer-histopatho...
                                                                         1
9997 /kaggle/input/lung-and-colon-cancer-histopatho...
                                                                         1
9998 /kaggle/input/lung-and-colon-cancer-histopatho...
                                                                         1
9999 /kaggle/input/lung-and-colon-cancer-histopatho...
                                                                         1
[10000 rows x 2 columns]
df resampled['category encoded'] =
df_resampled['category_encoded'].astype(str)
from sklearn.model selection import train test split
from sklearn.metrics import confusion_matrix, classification_report
import tensorflow as tf
from tensorflow import keras
from tensorflow.keras.models import Sequential
from tensorflow.keras.optimizers import Adam
from tensorflow.keras.preprocessing.image import ImageDataGenerator
from tensorflow.keras.layers import Conv2D, MaxPooling2D, Flatten, Dense,
Activation, Dropout, BatchNormalization
from tensorflow.keras import regularizers
import warnings
warnings.filterwarnings("ignore")
print ('check')
check
train_df_new, temp_df_new = train_test_split(
    df_resampled,
    train size=0.8,
    shuffle=True,
    random state=42,
    stratify=df_resampled['category_encoded']
)
valid_df_new, test_df_new = train_test_split(
    temp_df_new,
    test size=0.5,
    shuffle=True,
    random_state=42,
```

```
stratify=temp df new['category encoded']
)
batch_size = 16
img size = (256, 256)
channels = 3
img_shape = (img_size[0], img_size[1], channels)
tr_gen = ImageDataGenerator(rescale=1./255)
ts_gen = ImageDataGenerator(rescale=1./255)
train_gen_new = tr_gen.flow_from_dataframe(
    train_df_new,
    x_col='image_path',
    y_col='category_encoded',
    target_size=img_size,
    class_mode='binary',
    color_mode='rgb',
    shuffle=True,
    batch_size=batch_size
)
valid_gen_new = ts_gen.flow_from_dataframe(
    valid df new,
    x_col='image_path',
    y_col='category_encoded',
    target_size=img_size,
    class_mode='binary',
    color mode='rgb',
    shuffle=True,
    batch_size=batch_size
)
test_gen_new = ts_gen.flow_from_dataframe(
    test_df_new,
    x_col='image_path',
    y_col='category_encoded',
    target_size=img_size,
    class_mode='binary',
    color_mode='rgb',
    shuffle=False,
    batch_size=batch_size
)
Found 8000 validated image filenames belonging to 2 classes.
Found 1000 validated image filenames belonging to 2 classes.
Found 1000 validated image filenames belonging to 2 classes.
from tensorflow.keras.layers import Input, Conv2D, GlobalAveragePooling2D,
Dense, Multiply, Reshape, BatchNormalization, Activation
```

```
from tensorflow.keras.applications import ResNet50
from tensorflow.keras.models import Model
from tensorflow.keras.optimizers import Adam
print("Num GPUs Available: ", len(tf.config.list physical devices('GPU')))
Num GPUs Available: 2
gpus = tf.config.list physical devices('GPU')
if gpus:
    try:
        for gpu in gpus:
            tf.config.experimental.set_memory_growth(gpu, True)
        print("GPU is set for TensorFlow")
    except RuntimeError as e:
        print(e)
GPU is set for TensorFlow
from tensorflow.keras.layers import Add
from tensorflow.keras.layers import Add, GlobalAveragePooling2D, Dense,
BatchNormalization, Reshape, Multiply, Input, UpSampling2D
from tensorflow.keras.layers import Add, GlobalAveragePooling2D, Dense,
BatchNormalization, Reshape, Multiply, Input, Conv2D, UpSampling2D
from tensorflow.keras.applications import ResNet50
from tensorflow.keras.models import Model
from tensorflow.keras.optimizers import Adam
from tensorflow.keras import backend as K
def attention_block(inputs, filters):
    x = GlobalAveragePooling2D()(inputs)
    x = Dense(filters // 16, activation='relu')(x)
    x = Dense(filters, activation='sigmoid')(x)
    x = Reshape((1, 1, filters))(x)
    return Multiply()([inputs, x])
def build mranet(input shape=(256, 256, 3), num classes=2):
    base model = ResNet50(weights='imagenet', include top=False,
input tensor=Input(shape=input shape))
    conv4 block6 out = base model.get layer('conv4 block6 out').output
    conv5_block3_out = base_model.get_layer('conv5_block3_out').output
    attn_block1 = attention_block(conv4_block6_out, filters=1024)
    attn_block2 = attention_block(conv5_block3_out, filters=2048)
    attn_block2 = Conv2D(1024, (1, 1), padding='same',
activation='relu')(attn_block2)
```

```
attn block2 = UpSampling2D((2, 2))(attn block2)
    attn block1 shape = K.int shape(attn block1)
    attn block2 shape = K.int shape(attn block2)
    print("Shape of attn_block1:", attn_block1_shape)
    print("Shape of attn block2:", attn block2 shape)
    merged attention = Add()([attn block1, attn block2])
    x = GlobalAveragePooling2D()(merged_attention)
    x = Dense(512, activation='relu')(x)
    x = Dropout(0.5)(x)
    x = BatchNormalization()(x)
    x = Dense(1, activation='sigmoid')(x)
    model = Model(inputs=base model.input, outputs=x)
    for layer in base model.layers:
        layer.trainable = False
    return model
mranet model = build mranet()
mranet model.compile(
    optimizer=Adam(learning_rate=0.0001),
    loss='binary crossentropy',
    metrics=['accuracy'])
Downloading data from https://storage.googleapis.com/tensorflow/keras-
applications/resnet/resnet50_weights_tf_dim_ordering_tf_kernels_notop.h5
94765736/94765736 ——
                                     — 0s Ous/step
Shape of attn_block1: (None, 16, 16, 1024)
Shape of attn block2: (None, 16, 16, 1024)
from PIL import Image
def my image check(generator):
    while True:
        x, y = next(generator)
        new x = []
        new_y = []
        for i in range(x.shape[0]):
            try:
                img = Image.fromarray((x[i] * 255).astype(np.uint8))
                img.verify()
                new x.append(x[i])
                new y.append(y[i])
            except Exception as e:
                print(f"Error loading image:
{generator.dataframe['image_path'].iloc[generator.index + i] if
```

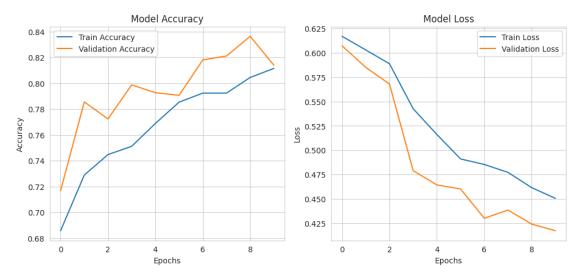
```
(generator.index + i) < len(generator.dataframe) else 'Path information not
available'}")
                print(f"Error: {e}")
        yield np.array(new_x), np.array(new_y)
tr gen = ImageDataGenerator(rescale=1./255)
ts_gen = ImageDataGenerator(rescale=1./255)
train generator = tr gen.flow from dataframe(
    train_df_new,
    x col='image path',
    y_col='category_encoded',
    target_size=img_size,
    class_mode='binary',
    color_mode='rgb',
    shuffle=True,
    batch size=batch_size,
    stratify=df resampled['category encoded']
train_gen_new = my_image_check(train_generator)
valid generator = ts gen.flow from dataframe(
    valid_df_new,
    x_col='image_path',
    y col='category encoded',
    target_size=img_size,
    class_mode='binary',
    color mode='rgb',
    shuffle=True,
    batch size=batch size,
    stratify=df resampled['category encoded']
)
valid gen new = my image check(valid generator)
test generator = ts gen.flow from dataframe(
    test_df_new,
    x_col='image_path',
    y_col='category_encoded',
    target_size=img_size,
    class mode='binary',
    color_mode='rgb',
    shuffle=False,
    batch_size=batch_size,
    stratify=df_resampled['category_encoded']
)
test_gen_new = my_image_check(test_generator)
```

```
steps_per_epoch = len(train_df_new) // batch_size
validation steps = len(valid df new) // batch size
Found 8000 validated image filenames belonging to 2 classes.
Found 1000 validated image filenames belonging to 2 classes.
Found 1000 validated image filenames belonging to 2 classes.
history = mranet model.fit(
   train_gen_new,
   epochs=10,
   validation data=valid_gen_new,
   steps per epoch=steps per epoch,
   validation_steps=validation_steps,
)
Epoch 1/10
              98s 163ms/step - accuracy: 0.6369 - loss: 0.6445
500/500 -----
- val accuracy: 0.7167 - val loss: 0.6070
Epoch 2/10
              500/500 ----
- val_accuracy: 0.7856 - val_loss: 0.5849
Epoch 3/10
           500/500 ----
- val_accuracy: 0.7724 - val_loss: 0.5678
Epoch 4/10
500/500 ----- 46s 92ms/step - accuracy: 0.7453 - loss: 0.5601
- val accuracy: 0.7988 - val loss: 0.4791
Epoch 5/10
500/500 ----- 46s 92ms/step - accuracy: 0.7732 - loss: 0.5173
- val accuracy: 0.7927 - val loss: 0.4644
Epoch 6/10
              - val accuracy: 0.7907 - val loss: 0.4603
Epoch 7/10
               ------- 44s 88ms/step - accuracy: 0.7960 - loss: 0.4876
500/500 ---
- val accuracy: 0.8181 - val loss: 0.4303
Epoch 8/10
           500/500 ---
- val accuracy: 0.8211 - val loss: 0.4387
- val_accuracy: 0.8364 - val_loss: 0.4243
Epoch 10/10
500/500 ----- 45s 90ms/step - accuracy: 0.8039 - loss: 0.4662
- val_accuracy: 0.8140 - val_loss: 0.4175
import matplotlib.pyplot as plt
def plot history(history):
   plt.figure(figsize=(12, 5))
```

```
plt.subplot(1, 2, 1)
plt.plot(history.history['accuracy'], label='Train Accuracy')
plt.plot(history.history['val_accuracy'], label='Validation Accuracy')
plt.title('Model Accuracy')
plt.xlabel('Epochs')
plt.ylabel('Accuracy')
plt.legend()

plt.subplot(1, 2, 2)
plt.plot(history.history['loss'], label='Train Loss')
plt.plot(history.history['val_loss'], label='Validation Loss')
plt.title('Model Loss')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()
```

plot_history(history)



import seaborn as sns
from sklearn.metrics import confusion_matrix, classification_report

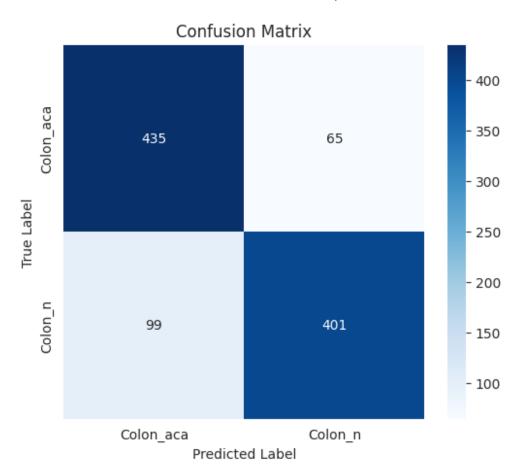
```
y_true = test_generator.classes
y_pred = mranet_model.predict(test_generator)
y_pred_classes = np.round(y_pred).astype(int)

cm = confusion_matrix(y_true, y_pred_classes)

plt.figure(figsize=(6, 5))
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues', xticklabels=['Colon_aca', 'Colon_n'], yticklabels=['Colon_aca', 'Colon_n'])
plt.xlabel('Predicted Label')
```

```
plt.ylabel('True Label')
plt.title('Confusion Matrix')
plt.show()

print("Classification Report:\n", classification_report(y_true, y_pred_classes))
```



Classification Report:

	precision	recall	f1-score	support
0	0.81	0.87	0.84	500
1	0.86	0.80	0.83	500
accuracy			0.84	1000
macro avg	0.84	0.84	0.84	1000
weighted avg	0.84	0.84	0.84	1000

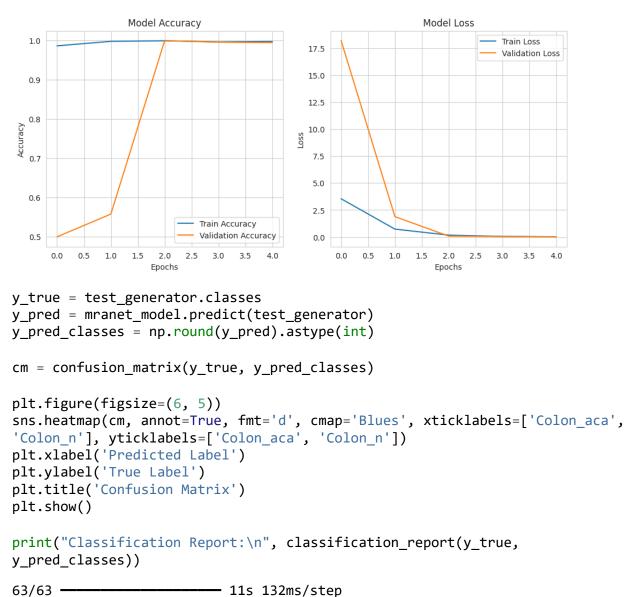
import tensorflow as tf
from tensorflow.keras.layers import Add, GlobalAveragePooling2D, Dense,
BatchNormalization, Reshape, Multiply, Input, Conv2D, UpSampling2D, Dropout

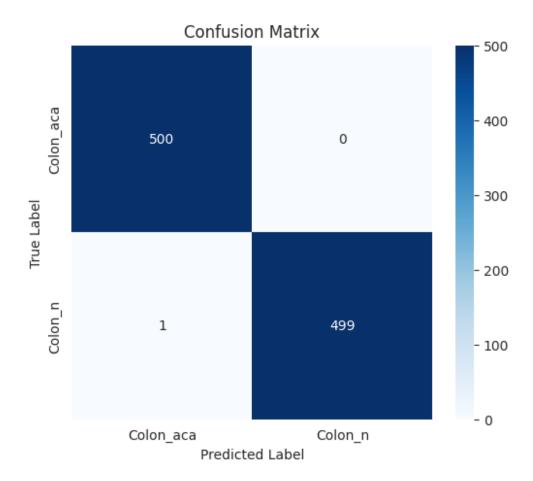
```
from tensorflow.keras.applications import ResNet50
from tensorflow.keras.models import Model
from tensorflow.keras.optimizers import Adam
from tensorflow.keras.regularizers import 12
from tensorflow.keras.callbacks import ReduceLROnPlateau
import numpy as np
def se_block(input_tensor, ratio=8):
    channels = input tensor.shape[-1]
    se = GlobalAveragePooling2D()(input tensor)
    se = Dense(channels // ratio, activation='relu')(se)
    se = Dense(channels, activation='sigmoid')(se)
    se = Reshape((1, 1, channels))(se)
    return Multiply()([input tensor, se])
def build_mranet(input_shape=(256, 256, 3), num_classes=1):
    base model = ResNet50(weights='imagenet', include top=False,
input tensor=Input(shape=input shape))
    conv4 block6 out = base model.get layer('conv4 block6 out').output
    conv5 block3 out = base model.get layer('conv5 block3 out').output
    attn block1 = se block(conv4 block6 out, ratio=8)
    attn_block2 = se_block(conv5_block3_out, ratio=8)
    attn_block2 = Conv2D(1024, (3, 3), dilation_rate=(2, 2), padding='same',
activation='relu')(attn block2)
    attn_block2 = UpSampling2D((2, 2))(attn_block2)
    attn_block1 = Conv2D(1024, (1, 1), padding='same',
activation='relu')(attn_block1)
    merged_attention = Add()([attn_block1, attn_block2])
    x = GlobalAveragePooling2D()(merged_attention)
    x = Dense(512, activation='relu', kernel_regularizer=12(0.01))(x)
    x = Dropout(0.5)(x)
    x = BatchNormalization()(x)
    x = Dense(num_classes, activation='sigmoid')(x)
    model = Model(inputs=base_model.input, outputs=x)
    for layer in base model.layers[-10:]:
        layer.trainable = True
    return model
mranet_model = build_mranet()
```

```
mranet model.compile(
   optimizer=Adam(learning rate=0.0001),
   loss='binary_crossentropy',
   metrics=['accuracy']
)
lr scheduler = ReduceLROnPlateau(monitor='val loss', factor=0.1, patience=5)
history = mranet model.fit(
   train_gen_new,
   epochs=5,
   validation data=valid gen new,
   steps per epoch=steps per epoch,
   validation steps=validation steps,
)
Epoch 1/5
             500/500 <del>---</del>
5.0099 - val accuracy: 0.5000 - val loss: 18.1901
Epoch 2/5
                   ------ 120s 240ms/step - accuracy: 0.9983 - loss:
500/500 -
1.0274 - val_accuracy: 0.5579 - val_loss: 1.8907
Epoch 3/5
                 ------ 116s 233ms/step - accuracy: 0.9991 - loss:
500/500 <del>---</del>
0.2367 - val_accuracy: 1.0000 - val_loss: 0.0838
Epoch 4/5
0.0673 - val_accuracy: 0.9959 - val_loss: 0.0498
Epoch 5/5
500/500 ————— 114s 228ms/step - accuracy: 0.9967 - loss:
0.0344 - val_accuracy: 0.9949 - val_loss: 0.0199
def plot_history(history):
   plt.figure(figsize=(12, 5))
   plt.subplot(1, 2, 1)
   plt.plot(history.history['accuracy'], label='Train Accuracy')
   plt.plot(history.history['val_accuracy'], label='Validation Accuracy')
   plt.title('Model Accuracy')
   plt.xlabel('Epochs')
   plt.ylabel('Accuracy')
   plt.legend()
   plt.subplot(1, 2, 2)
   plt.plot(history.history['loss'], label='Train Loss')
   plt.plot(history.history['val_loss'], label='Validation Loss')
   plt.title('Model Loss')
   plt.xlabel('Epochs')
   plt.ylabel('Loss')
   plt.legend()
```

plt.show()

plot_history(history)





Classification Report:

Classification	precision	recall	f1-score	support
0	1.00	1.00	1.00	500
1	1.00	1.00	1.00	500
accuracy			1.00	1000
macro avg	1.00	1.00	1.00	1000
weighted avg	1.00	1.00	1.00	1000