```
from google.colab import drive
drive.mount('/content/drive', force_remount=True)
    Mounted at /content/drive
import pandas as pd
import numpy as np
import tensorflow as tf
from tensorflow import keras
from keras import Sequential
import matplotlib.pyplot as plt
from PIL import Image
import os
import pathlib
import random
import glob
# Create data directory in your google MyDrive folder.
# Copy all three folders in your data directory to MyDrive->data folder in your google drive
# Then Execute below code
def get data dir path():
  path = '/content/drive/MyDrive/data/'
  data_dir_path= pathlib.Path(path)
  data_dir_file = str(pathlib.Path(path)) + "/*.*"
  return data dir path
# Create filter path to retive all the png files
def filter path():
  data_dir_path_benign = str(pathlib.Path(os.path.join(get_data_dir_path(),'benign'))) + "/*.png"
  data_dir_path_normal = str(pathlib.Path(os.path.join(get_data_dir_path(),'normal'))) + "/*.png"
  data dir path malignant = str(pathlib.Path(os.path.join(get data dir path(), 'malignant'))) + "/*.png"
  return data_dir_path_benign,data_dir_path_normal,data_dir_path_malignant
# Use the filter to retrieve the corresponsing image counts
def image count():
  data_dir_path_benign,data_dir_path_normal,data_dir_path_malignant = filter_path()
  img count benign = glob.glob(data dir path benign,recursive=True)
  img_count_normal = glob.glob(data_dir_path_normal,recursive=True)
  img_count_malignant = glob.glob(data_dir_path_malignant,recursive=True)
  img total count = len(img count benign) + len(img count normal) + len(img count malignant)
  print("Total Images: ", img_total_count)
  print("Benign (non-dangerous) Images: {}({})".format(len(img_count_benign), round(len(img_count_benign)*1
  print("Malignant (dangerous) Images: {}({})".format(len(img count normal), round(len(img count normal)*10
  print("Normal (No Traces) Images: {}({})".format(len(img_count_malignant), round(len(img_count_malignant)
  #return img total count,len(img count benign),len(img count normal),len(img count malignant)
#Display the image totals and respective class image counts
image count()
\rightarrow
    Total Images: 1587
     Benign (non-dangerous) Images: 900(56.71)
     Malignant (dangerous) Images: 266(16.76)
```

model = create model()

model.summary()

cnn_bc_detection.ipynb - Colab Normal (No Traces) Images: 421(26.53) # Configure batch size,img size and create train, validation split use function call configure_for_performa def create data sets(): batch_size = 40 $img_height = 224$ img_width = 224 train_ds = tf.keras.utils.image_dataset_from_directory(get_data_dir_path(),validation_split=0.3,subset="t val_ds = tf.keras.utils.image_dataset_from_directory(get_data_dir_path(),validation_split=0.3,subset="val val ds.class names return train_ds,val_ds train ds,val ds = create data sets() → Found 1587 files belonging to 3 classes. Using 1111 files for training. Found 1587 files belonging to 3 classes. Using 476 files for validation. def create_model(img_height=224,img_width=224): model = tf.keras.Sequential([tf.keras.layers.Rescaling(1./255, input_shape=(img_height, img_width, 3)), tf.keras.layers.Conv2D(16, 3, padding='same', activation='relu'), tf.keras.layers.MaxPooling2D(), tf.keras.layers.Conv2D(32, 3, padding='same', activation='relu'), tf.keras.layers.MaxPooling2D(), tf.keras.layers.Conv2D(64, 3, padding='same', activation='relu'), tf.keras.layers.MaxPooling2D(), tf.keras.layers.Conv2D(128, 3, padding='same', activation='relu'), tf.keras.layers.MaxPooling2D(), tf.keras.layers.Dropout(0.3), tf.keras.layers.Flatten(), tf.keras.layers.Dense(64, activation='relu'), tf.keras.layers.Dense(3,activation="softmax") 1) return model

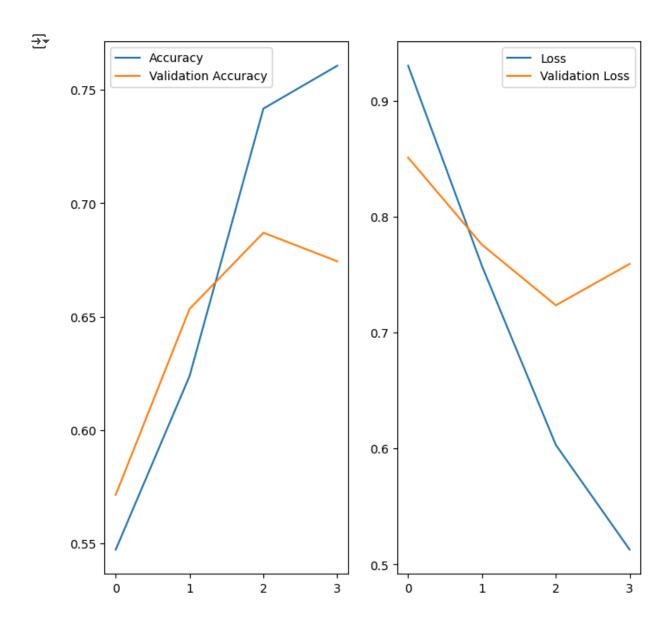
→ Model: "sequential_7"

Layer (type)	Output Shape	Param #
rescaling_7 (Rescaling)	(None, 224, 224, 3)	0
conv2d_23 (Conv2D)	(None, 224, 224, 16)	448
max_pooling2d_23 (MaxPooling2D)	(None, 112, 112, 16)	0
conv2d_24 (Conv2D)	(None, 112, 112, 32)	4,640
max_pooling2d_24 (MaxPooling2D)	(None, 56, 56, 32)	0
conv2d_25 (Conv2D)	(None, 56, 56, 64)	18,496
max_pooling2d_25 (MaxPooling2D)	(None, 28, 28, 64)	0
conv2d_26 (Conv2D)	(None, 28, 28, 128)	73,856
max_pooling2d_26 (MaxPooling2D)	(None, 14, 14, 128)	0
dropout_7 (Dropout)	(None, 14, 14, 128)	0
flatten_7 (Flatten)	(None, 25088)	0
dense_14 (Dense)	(None, 64)	1,605,696
dense_15 (Dense)	(None, 3)	195

```
Total params: 1,703,331 (6.50 MB)
      Trainable narams: 1.703.331 (6.50 MR)
def train_model(batch_size = 32,epochs=4):
  train ds,val ds = create data sets()
  model.compile(optimizer="Adam",
            loss=tf.keras.losses.SparseCategoricalCrossentropy(from_logits=True),
            metrics=["accuracy"])
  history = model.fit(train_ds,
                    epochs=epochs,
                    validation_data=val_ds,
                    batch size=batch size)
  return history
#Train the model
history=train_model(batch_size = 32,epochs=4)
history.history.keys()
\rightarrow Found 1587 files belonging to 3 classes.
    Using 1111 files for training.
    Found 1587 files belonging to 3 classes.
    Using 476 files for validation.
    Epoch 1/4
    /usr/local/lib/python3.11/dist-packages/keras/src/backend/tensorflow/nn.py:708: UserWarning: "`sparse_ca
      output, from_logits = _get_logits(
                              - 90s 3s/step - accuracy: 0.4895 - loss: 0.9795 - val_accuracy: 0.5714 - val_lc
    28/28 -
    Epoch 2/4
                              - 147s 3s/step - accuracy: 0.6105 - loss: 0.7821 - val_accuracy: 0.6534 - val_l
    28/28 -
    Epoch 3/4
    28/28 -
                              - 142s    3s/step - accuracy: 0.7500 - loss: 0.5849 - val_accuracy: 0.6870 - val_l
    Epoch 4/4
                              - 143s 3s/step - accuracy: 0.7586 - loss: 0.5129 - val_accuracy: 0.6744 - val_l
    28/28 -
```

```
dict_keys(['accuracy', 'loss', 'val_accuracy', 'val_loss'])
```

```
#Plot the graph acc/val accuracy versus loss/validation loss
epochs=4
acc = history.history['accuracy']
val acc = history.history['val accuracy']
loss = history.history['loss']
val_loss = history.history['val_loss']
ep range = range(epochs)
plt.figure(figsize=(8,8))
plt.subplot(1,2,1)
plt.plot(ep_range,acc,label='Accuracy')
plt.plot(ep_range,val_acc,label="Validation Accuracy")
plt.legend()
plt.subplot(1,2,2)
plt.plot(ep_range,loss,label='Loss')
plt.plot(ep range,val loss,label="Validation Loss")
plt.legend()
plt.show()
```



```
#Evaluate model with validation data set
train_ds,val_ds = create_data_sets()
loss, acc = model.evaluate(val ds)
print('model, accuracy: {:5.2f}%'.format(100 * acc))
Found 1587 files belonging to 3 classes.
     Using 1111 files for training.
     Found 1587 files belonging to 3 classes.
     Using 476 files for validation.
     12/12 ---
                              - 18s 847ms/step - accuracy: 0.6652 - loss: 0.7885
     model, accuracy: 67.44%
#Pedict the model using Validation data
plt.figure(figsize=(15, 15))
class_names = val_ds.class_names
result = ' | False'
for images, labels in val_ds.take(1):
    for i in range(25):
        ax = plt.subplot(5, 5, i + 1)
        img = images[i].numpy().astype("uint8")
        img = tf.expand_dims(img, axis=0)
        predication=""
        predictions = model.predict(img)
        predicted class = np.argmax(predictions)
        if class_names[predicted_class] == class_names[labels[i]]:
            result = ' | TRUE'
        plt.imshow(images[i].numpy().astype("uint8"))
        plt.title('Predicated:'+ class_names[predicted_class]+result)
        plt.axis("off")
```


 0s
 63ms/step

 0s
 62ms/step

 0s
 60ms/step

 0s
 57ms/step
 1/1 -1/1 -1/1 -
 Os
 56ms/step

 Os
 71ms/step

 Os
 55ms/step
 1/1 -1/1 -1/1 ----- **0s** 60ms/step 1/1 -1/1 ______ 0s 56ms/step 1/1 _____ 0s 58ms/step 1/1 -1/1 -**1/1** ——— **0s** 56ms/step 1/1 ---- 0s 64ms/step

 0s
 64ms/step

 0s
 59ms/step

 0s
 59ms/step

 0s
 61ms/step

 0s
 58ms/step

 1/1 -1/1 -**Os** 72ms/step **———— 0s** 91ms/step 1/1 ------ **0s** 81ms/step 1/1 -**Os** 86ms/step **Os** 83ms/step 1/1 -Predicated:normal | TRUE | Predicated:benign | TRUE Predicated:benign | TRUE | Predicated:benign | TRUE Predicated:normal | TRUE Predicated:benign | TRUE | Predicated:normal | TRUE Predicated:benign | TRUE | Predicated:benign | TRUE Predicated:benign | TRUE



new_model.summary()



new_model = tf.keras.models.load_model(data_dir_file_path)







WARNING:absl:Compiled the loaded model, but the compiled metrics have yet to be built. `model.compile_m Model: "sequential_7"

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