## → Import all the Dependencies

```
import numpy as np
import cv2
import PIL.Image as Image
import shutil
import os
from IPython.display import HTML
import matplotlib.pylab as plt
import tensorflow as tf
import tensorflow_hub as hub
from tensorflow import keras
from tensorflow.keras import models, layers
from tensorflow.keras.models import Sequential
import time
```

# Downloading Data from link

```
!mkdir -p ~/.kaggle
!cp kaggle.json ~/.kaggle/
!kaggle datasets download -d fanconic/skin-cancer-malignant-vs-benign

Warning: Your Kaggle API key is readable by other users on this system! To fix this, you can run 'chmod 600 /root/.kaggle/kaggle.json'
    Downloading skin-cancer-malignant-vs-benign.zip to /content
    94% 304M/325M [00:04<00:00, 30.4MB/s]
    100% 325M/325M [00:04<00:00, 73.6MB/s]

import zipfile
zip_ref = zipfile.ZipFile('/content/skin-cancer-malignant-vs-benign.zip', 'r')
zip_ref.extractall('/content')
zip_ref.close()

curr_dir = os.getcwd()
curr_dir
    '/content'</pre>
```

### Setting up directories

```
# creating folders

datasetFolder = curr_dir + "/DATASET"
    os.makedirs(datasetFolder)
    os.makedirs(datasetFolder + "/benign")
    os.makedirs(datasetFolder + "/malignant")

# defining source and destination folders paths

src1 = curr_dir + "/test/benign"

src2 = curr_dir + "/train/benign"

src3 = curr_dir + "/test/malignant"

src4 = curr_dir + "/train/malignant"

benign_src = [src1, src2]

malignant_src = [src3, src4]
```

```
benign_dest = curr_dir + "/DATASET/benign"
malignant_dest = curr_dir + "/DATASET/malignant"
## copying files
for src in benign_src:
 for dirs, subdirs, files in os.walk(src):
   print(" Total benign files : ", len(files))
   for file in files:
     if file.endswith('.jpg'):
       filename = os.path.join(src, dirs, file)
       if os.path.exists(filename):
          # print(filename)
          shutil.copy(filename, benign_dest)
for src in malignant_src:
 for dirs, subdirs, files in os.walk(src):
   print(" Total malignant files : ", len(files))
   for file in files:
     if file.endswith('.jpg'):
       filename = os.path.join(src, dirs, file)
       if os.path.exists(filename):
          # print(filename)
          shutil.copy(filename, malignant_dest)
     Total benign files : 360
      Total benign files : 1440
     Total malignant files: 300
     Total malignant files : 1197
## deleting old folders
shutil.rmtree(curr_dir + "/data")
shutil.rmtree(curr_dir + "/test")
shutil.rmtree(curr_dir + "/train")
Total_images = 0
for dirs, subdirs, files in (os.walk(benign_dest)):
 print(f'Benign : {len(files)}')
 Total_images = Total_images + len(files)
for dirs, subdirs, files in (os.walk(malignant_dest)):
 print(f'Malignant : {len(files)}')
 Total_images = Total_images + len(files)
print(f'\nTotal images : {Total_images}')
     Benign: 1800
    Malignant: 1497
    Total images: 3297
```

### Setting Constants

```
BATCH_SIZE = 32

IMAGE_SIZE = 224

CHANNELS = 3

EPOCHS = 50

dataset = tf.keras.preprocessing.image_dataset_from_directory(
    "./DATASET",
    seed = 123,
    shuffle = True,
    image_size = (IMAGE_SIZE,IMAGE_SIZE),
    batch_size = BATCH_SIZE
)

Found 3297 files belonging to 2 classes.
```

### → Data Visualization

```
class_names = dataset.class_names
class_names
    ['benign', 'malignant']
for image_batch, labels_batch in dataset.take(1):
   print(image_batch.shape)
   print(labels_batch.numpy())
     (32, 224, 224, 3)
    [1\ 0\ 1\ 0\ 1\ 1\ 0\ 1\ 0\ 0\ 0\ 0\ 1\ 0\ 1\ 0\ 1\ 0\ 1\ 1\ 1\ 0\ 1\ 0\ 1\ 0\ 1\ 1\ 0]
plt.figure(figsize=(10, 10))
for image_batch, labels_batch in dataset.take(1):
   for i in range(12):
       ax = plt.subplot(3, 4, i + 1)
       plt.imshow(image_batch[i].numpy().astype("uint8"))
       plt.title(class_names[labels_batch[i]])
       plt.axis("off")
           malignant
                                                                   benign
                                                                                              benign
                                      malignant
             benign
                                      malignant
                                                                   benign
                                                                                              benign
            malignant
                                                                                            malignant
                                      malignant
                                                                   benign
len(dataset)
    104
train_size = 0.8
len(dataset)*train_size
```

```
train_ds = dataset.take(54)
len(train_ds)
     54
test_ds = dataset.skip(54)
len(test_ds)
     50
val_size=0.1
len(dataset)*val_size
     10.4
val_size=0.1
len(dataset)*val_size
     10.4
test_ds = test_ds.skip(6)
len(test_ds)
     44
def get_dataset_partitions_tf(ds, train_split=0.8, val_split=0.1, test_split=0.1, shuffle=True, shuffle_size=10000):
    assert (train_split + test_split + val_split) == 1
   ds_size = len(ds)
    if shuffle:
        ds = ds.shuffle(shuffle_size, seed=12)
   train_size = int(train_split * ds_size)
   val_size = int(val_split * ds_size)
   train_ds = ds.take(train_size)
   val_ds = ds.skip(train_size).take(val_size)
   test_ds = ds.skip(train_size).skip(val_size)
   return train_ds, val_ds, test_ds
train_ds, val_ds, test_ds = get_dataset_partitions_tf(dataset)
len(train_ds)
     83
len(val_ds)
     10
len(test_ds)
     11
actual_label_test = []
for image_batch, labels_batch in test_ds:
   temp = labels_batch.numpy()
   for j in temp:
        actual_label_test.append(j)
# print(len(actual_label_test))
# print(actual_label_test)
```

# - Catching, Prefetching and setting resize rescale layers

```
train_ds = train_ds.cache().shuffle(1000).prefetch(buffer_size=tf.data.AUTOTUNE)
val_ds = val_ds.cache().shuffle(1000).prefetch(buffer_size=tf.data.AUTOTUNE)
test_ds = test_ds.cache().shuffle(1000).prefetch(buffer_size=tf.data.AUTOTUNE)

resize_and_rescale = tf.keras.Sequential([
    layers.experimental.preprocessing.Resizing(IMAGE_SIZE, IMAGE_SIZE),
    layers.experimental.preprocessing.Rescaling(1./255),
])
```

### Data Augmentation

### Model Building

```
input_shape = (BATCH_SIZE, IMAGE_SIZE, IMAGE_SIZE, CHANNELS)
n_{classes} = 2
model = models.Sequential([
   resize_and_rescale,
   layers.Conv2D(32, kernel_size = (3,3), activation='relu', input_shape=input_shape),
   layers.MaxPooling2D((2, 2)),
   layers.Conv2D(64, kernel_size = (3,3), activation='relu'),
   layers.MaxPooling2D((2, 2)),
   layers.Conv2D(64, kernel_size = (3,3), activation='relu'),
   layers.MaxPooling2D((2, 2)),
   layers.Conv2D(64, (3, 3), activation='relu'),
   layers.MaxPooling2D((2, 2)),
   layers.Conv2D(64, (3, 3), activation='relu'),
   layers.MaxPooling2D((2, 2)),
   layers.Conv2D(64, (3, 3), activation='relu'),
   layers.MaxPooling2D((2, 2)),
   layers.Flatten(),
   layers.Dense(64, activation='relu'),
   layers.Dense(n_classes, activation='softmax'),
])
model.build(input_shape=input_shape)
model.summary()
```

Model: "sequential 3"

Layer (type)	Output Shape	Param #
sequential_2 (Sequential)	(32, 224, 224, 3)	0
conv2d_6 (Conv2D)	(32, 222, 222, 32)	896
<pre>max_pooling2d_6 (MaxPooling 2D)</pre>	(32, 111, 111, 32)	0
conv2d_7 (Conv2D)	(32, 109, 109, 64)	18496
max_pooling2d_7 (MaxPooling 2D)	(32, 54, 54, 64)	0
conv2d_8 (Conv2D)	(32, 52, 52, 64)	36928
max_pooling2d_8 (MaxPooling	(32, 26, 26, 64)	0

```
2D)
conv2d_9 (Conv2D)
                         (32, 24, 24, 64)
                                                36928
max_pooling2d_9 (MaxPooling (32, 12, 12, 64)
conv2d_10 (Conv2D)
                         (32, 10, 10, 64)
                                                36928
max_pooling2d_10 (MaxPoolin (32, 5, 5, 64)
conv2d_11 (Conv2D)
                         (32, 3, 3, 64)
                                                36928
max_pooling2d_11 (MaxPoolin (32, 1, 1, 64)
                                                0
flatten_1 (Flatten)
                         (32, 64)
                                                0
dense_2 (Dense)
                         (32, 64)
                                                4160
dense_3 (Dense)
                         (32, 2)
                                                130
______
Total params: 171,394
Trainable params: 171,394
Non-trainable params: 0
```

```
import time
t0 = time.time()

model.compile(
    optimizer='adam',
    loss=tf.keras.losses.SparseCategoricalCrossentropy(from_logits=False),
    metrics=['accuracy']
)

history = model.fit(
    train_ds,
    batch_size = BATCH_SIZE,
    validation_data = val_ds,
    verbose = 1,
    epochs = EPOCHS,
)
```

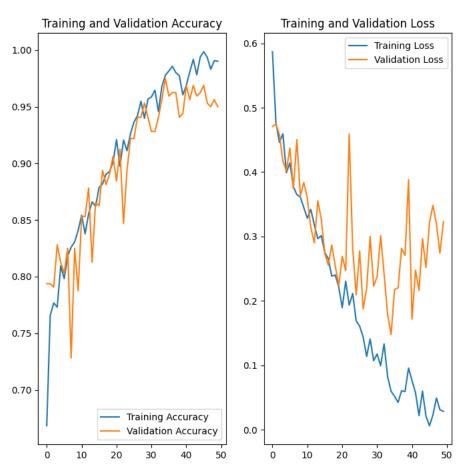
```
Epoch 39/50
Epoch 40/50
Epoch 41/50
Epoch 42/50
Epoch 43/50
Fnoch 44/50
83/83 [============] - 4s 44ms/step - loss: 0.0602 - accuracy: 0.9782 - val_loss: 0.2967 - val_accuracy: 0.9594
Epoch 45/50
Epoch 46/50
Epoch 47/50
Epoch 48/50
Fnoch 49/50
83/83 [=====
  Epoch 50/50
83/83 [=============] - 4s 44ms/step - loss: 0.0286 - accuracy: 0.9902 - val_loss: 0.3232 - val_accuracy: 0.9500
```

## Model Analysis

```
t1 = time.time()
                                                                                                        print("CNN Model Training time: ", (t1-t0)/60 , "minutes")
   CNN Model Training time: 3.990845835208893 minutes
                                                                                                        scores = model.evaluate(test_ds)
   scores
   [0.3968188166618347, 0.9431818127632141]
                                                                                                        predicted = model.predict(test_ds)
   11/11 [======== ] - 1s 20ms/step
                                                                                                        import numpy as np
confidence = np.max(predicted, axis=1)
predictions = np.argmax(predicted, axis=1)
```

```
plt.figure(figsize=(8, 8))
plt.subplot(1, 2, 1)
plt.plot(range(EPOCHS), acc, label='Training Accuracy')
plt.plot(range(EPOCHS), val_acc, label='Validation Accuracy')
plt.legend(loc='lower right')
plt.title('Training and Validation Accuracy')

plt.subplot(1, 2, 2)
plt.plot(range(EPOCHS), loss, label='Training Loss')
plt.plot(range(EPOCHS), val_loss, label='Validation Loss')
plt.legend(loc='upper right')
plt.title('Training and Validation Loss')
plt.show()
```

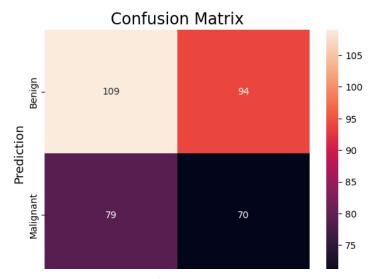


```
from sklearn.metrics import confusion_matrix
import seaborn as sns
import matplotlib.pyplot as plt

cm = confusion_matrix(actual_label_test, predictions)

sns.heatmap(
    cm,
    annot=True,
    fmt='g',
    xticklabels=['Benign','Malignant'],
    yticklabels=['Benign','Malignant'])
)

plt.ylabel('Prediction',fontsize=13)
plt.xlabel('Actual',fontsize=13)
plt.title('Confusion Matrix',fontsize=17)
plt.show()
```



from sklearn.metrics import classification\_report
print(classification\_report(actual\_label\_test, predictions))

	precision	recall	f1-score	support
0	0.58 0.43	0.54 0.47	0.56 0.45	203 149
accuracy			0.51	352
macro avg	0.50	0.50	0.50	352
weighted avg	0.52	0.51	0.51	352

# → Saving Model

```
# import os
```

✓ 0s completed at 10:40 AM

 $<sup>\</sup>label{eq:continuous} \begin{tabular}{ll} \# model\_version=max([int(i) \ for \ i \ in \ os.listdir("../savedmodels") \ + \ [0]])+1 \end{tabular}$ 

<sup>#</sup> model.save(f"/content/savedmodels/{model\_version}")