Problem statement: To build a CNN based model which can accurately detect melanoma. Melanoma is a type of cancer that can be deadly if not detected early. It accounts for 75% of skin cancer deaths. A solution which can evaluate images and alert the dermatologists about the presence of melanoma has the potential to reduce a lot of manual effort needed in diagnosis.

Importing Skin Cancer Data

To do: Take necessary actions to read the data

Importing all the important libraries

```
In [1]: import pathlib
        import tensorflow as tf
        import matplotlib.pyplot as plt
        import numpy as np
        import pandas as pd
        import os
        import PIL
        from tensorflow import keras
        from tensorflow.keras import layers
        from tensorflow.keras.models import Sequential
        from tensorflow.keras.layers import Dense, Dropout, Flatten, Conv2D, MaxPooling2D
```

Uncomment to read images from Google Drive

```
In [2]: ## If you are using the data by mounting the google drive, use the following:
        #from google.colab import drive
        #drive.mount('/content/gdrive')
        ##Ref:https://towardsdatascience.com/downloading-datasets-into-google-drive-via-google-colab-bcb1b30b016
```

This assignment uses a dataset of about 2357 images of skin cancer types. The dataset contains 9 sub-directories in each train and test subdirectories. The 9 sub-directories contains the images of 9 skin cancer types respectively.

```
In [3]: # Defining the path for train and test images
        ## Todo: Update the paths of the train and test dataset
        # data dir train = pathlib.Path("gdrive/My Drive/UpgradExer5CNNSkin/SkinImg/Train/")
        # data_dir_test = pathlib.Path('gdrive/My Drive/UpgradExer5CNNSkin/SkinImg/Test')
        data_dir_train = pathlib.Path("./UpgradExer5CNNSkin/SkinImg/Train/")
        data_dir_test = pathlib.Path('./UpgradExer5CNNSkin/SkinImg/Test')
```

Train (split for Validation data) and Test data set

```
In [4]: | image_count_train = len(list(data_dir_train.glob('*/*.jpg')))
        print(image_count_train)
        image_count_test = len(list(data_dir_test.glob('*/*.jpg')))
        print(image_count_test)
        2239
        118
```

Load using keras.preprocessing

Let's load these images off disk using the helpful image dataset from directory utility.

Create a dataset

Define some parameters for the loader:

```
In [5]: batch_size = 32
        img height = 180
        img width = 180
```

Use 80% of the images for training, and 20% for validation.

```
In [6]: ## Write your train dataset here
        ## Note use seed=123 while creating your dataset using tf.keras.preprocessing.image dataset from director
        ## Note, make sure your resize your images to the size img height*img width, while writting the dataset
        #train_ds = ##todo
        train_ds = tf.keras.preprocessing.image_dataset_from_directory(
          data dir train,
          seed=123,
          validation_split = 0.2,
          subset = "training", ## Todo choose the correct parameter value, so that only training data is refered
          image_size = (img_height, img_width),
          batch_size = batch_size)
        Found 2239 files belonging to 9 classes.
        Using 1792 files for training.
```

```
In [7]: ## Write your validation dataset here
        ## Note use seed=123 while creating your dataset using tf.keras.preprocessing.image_dataset_from_director
        ## Note, make sure your resize your images to the size img_height*img_width, while writting the dataset
        val_ds = tf.keras.preprocessing.image_dataset_from_directory(
          data_dir_train,
          seed=123,
          validation_split = 0.2,
          subset = "validation", ## Todo choose the correct parameter value, so that only training data is refere
          image_size = (img_height, img_width),
          batch_size = batch_size)
```

Found 2239 files belonging to 9 classes. Using 447 files for validation.

List of classes

```
In [8]: # List out all the classes of skin cancer and store them in a list.
        # You can find the class names in the class names attribute on these datasets.
        # These correspond to the directory names in alphabetical order.
        class_names = train_ds.class_names
        print(class_names)
        #len = len(class_names)
```

['actinic keratosis', 'basal cell carcinoma', 'dermatofibroma', 'melanoma', 'nevus', 'pigmented benign keratosis', 'seborrheic keratosis', 'squamous cell carcinoma', 'vascular lesion']

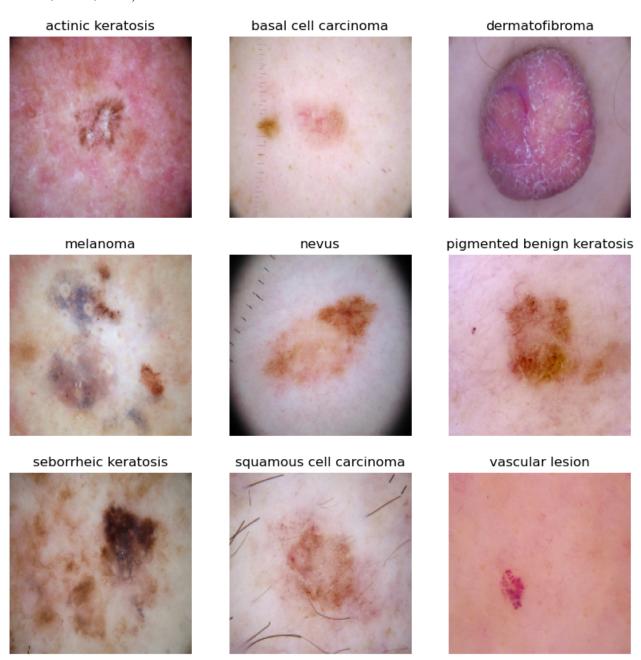
Visualize the data

Todo, create a code to visualize one instance of all the nine classes present in the dataset

```
In [9]: import matplotlib.pyplot as plt
        ### your code goes here, you can use training or validation data to visualize
        plt.figure(figsize=(10, 10))
        for i in range(len(class_names)):
            filtered_ds = train_ds.filter(lambda x, 1: tf.math.equal(l[0], i)).take(1)
            for images, labels in filtered_ds:
                ax = plt.subplot(3, 3, i + 1)
                plt.imshow(images[0].numpy().astype("uint8"))
                plt.title(class_names[labels[0]])
                plt.axis("off")
```

 $WARNING: tensorflow: From d: \Users \admin\anaconda $$\lib\site-packages \tensorflow \python \autograph\pyct\starrow \site-packages \tensorflow \python \autograph\pyct\starrow \python \pyt$ tic_analysis\liveness.py:83: Analyzer.lamba_check (from tensorflow.python.autograph.pyct.static_analysi s.liveness) is deprecated and will be removed after 2023-09-23. Instructions for updating:

Lambda fuctions will be no more assumed to be used in the statement where they are used, or at least in the same block. https://github.com/tensorflow/tensorflow/issues/56089 (https://github.com/tensorflow/te nsorflow/issues/56089)



The image_batch is a tensor of the shape (32, 180, 180, 3). This is a batch of 32 images of shape 180x180x3 (the last dimension refers to color channels RGB). The label_batch is a tensor of the shape (32,), these are corresponding labels to the 32 images.

```
In [10]: for image batch, labels batch in train ds:
            print(image_batch.shape)
            print(labels_batch.shape)
            break
          (32, 180, 180, 3)
          (32,)
           Dataset.cache() keeps the images in memory after they're loaded off disk during the first epoch.
```

```
In [11]: AUTOTUNE = tf.data.experimental.AUTOTUNE
         train_ds = train_ds.cache().shuffle(1000).prefetch(buffer_size=AUTOTUNE)
         val_ds = val_ds.cache().prefetch(buffer_size=AUTOTUNE)
```

Dataset.prefetch() overlaps data preprocessing and model execution while training.

Create the model

Todo: Create a CNN model, which can accurately detect 9 classes present in the dataset. Use layers.experimental.preprocessing.Rescaling to normalize pixel values between (0,1). The RGB channel values are in the [0, 255] range. This is not ideal for a neural network. Here, it is good to standardize values to be in the [0, 1]

```
In [12]: ### Your code goes here
         # model = tf.keras.Sequential([
         # tf.keras.layers.Rescaling(1./255),
         # tf.keras.layers.Conv2D(32, 3, activation='relu'),
         # tf.keras.Layers.MaxPooling2D(),
         # tf.keras.layers.Dropout(0.1),
         # tf.keras.layers.Conv2D(32, 3, activation='relu'),
         # tf.keras.layers.MaxPooling2D(),
         # tf.keras.layers.Dropout(0.1),
         # tf.keras.layers.Conv2D(32, 3, activation='relu'),
         # tf.keras.layers.MaxPooling2D(),
         # tf.keras.Layers.Dropout(0.1),
         # tf.keras.layers.Flatten(),
         # tf.keras.layers.Dense(128, activation='relu'),
         # tf.keras.layers.Dense(num_classes)
         # tf.keras.layers.Dropout(0.1)
         num_classes = 9
         model = Sequential()
         model.add(Conv2D(16,3,padding="same", activation="relu"))
         model.add(MaxPooling2D())
         model.add(Dropout(0.1))
         model.add(Conv2D(32, 3, padding="same", activation="relu"))
         model.add(MaxPooling2D())
         model.add(Dropout(0.1))
         model.add(Conv2D(64, 3, padding="same", activation="relu"))
         model.add(MaxPooling2D())
         model.add(Dropout(0.1))
         model.add(Flatten())
         model.add(Dense(128,activation="relu"))
         model.add(Dense(num_classes))
         model.add(Dropout(0.1))
```

Compile the model

Choose an appropirate optimiser and loss function for model training

```
In [13]: ### Todo, choose an appropirate optimiser and loss function
         model.compile(optimizer='adam',
                       loss=tf.keras.losses.SparseCategoricalCrossentropy(from_logits=True),
                       metrics=['accuracy'])
```

Train the model

```
In [14]: | # Training and Validation happens Parallel in Batches (epouch).
         # As long as the training accuracy gets better we can let the iterations continue.
         # In case the training is not accurate we can retune the model
         epochs = 20
         history = model.fit(
          train_ds,
           validation_data=val_ds,
           epochs=epochs
```

```
Epoch 1/20
2.1039 - val_accuracy: 0.3781
Epoch 2/20
56/56 [============= ] - 42s 747ms/step - loss: 1.8411 - accuracy: 0.3549 - val loss:
1.9932 - val accuracy: 0.2483
Epoch 3/20
2.1288 - val_accuracy: 0.2215
Epoch 4/20
56/56 [============= ] - 42s 747ms/step - loss: 1.6806 - accuracy: 0.4129 - val loss:
1.9054 - val accuracy: 0.3445
Epoch 5/20
56/56 [====================] - 42s 757ms/step - loss: 1.5984 - accuracy: 0.4291 - val_loss:
2.0002 - val_accuracy: 0.3333
Epoch 6/20
1.9125 - val_accuracy: 0.3199
Epoch 7/20
1.9445 - val_accuracy: 0.3356
Epoch 8/20
1.8135 - val_accuracy: 0.3982
Epoch 9/20
1.6633 - val accuracy: 0.3870
Epoch 10/20
1.9272 - val_accuracy: 0.3244
Epoch 11/20
56/56 [===============] - 42s 758ms/step - loss: 1.3582 - accuracy: 0.5123 - val_loss:
1.8587 - val_accuracy: 0.3400
Epoch 12/20
56/56 [===============] - 45s 800ms/step - loss: 1.3533 - accuracy: 0.5089 - val_loss:
1.6263 - val_accuracy: 0.4631
Epoch 13/20
56/56 [====================] - 42s 746ms/step - loss: 1.3156 - accuracy: 0.5335 - val_loss:
1.7482 - val_accuracy: 0.4072
Epoch 14/20
56/56 [============= ] - 44s 780ms/step - loss: 1.2398 - accuracy: 0.5575 - val loss:
1.6596 - val_accuracy: 0.4720
Epoch 15/20
1.7573 - val_accuracy: 0.4094
Epoch 16/20
56/56 [============= ] - 42s 746ms/step - loss: 1.1881 - accuracy: 0.5770 - val loss:
1.7437 - val_accuracy: 0.4273
Epoch 17/20
56/56 [==============] - 46s 818ms/step - loss: 1.1671 - accuracy: 0.5859 - val_loss:
1.9293 - val_accuracy: 0.4318
Epoch 18/20
56/56 [================= ] - 44s 789ms/step - loss: 1.0772 - accuracy: 0.6239 - val loss:
1.8437 - val_accuracy: 0.4251
Epoch 19/20
56/56 [================ ] - 43s 761ms/step - loss: 0.9794 - accuracy: 0.6602 - val loss:
1.7959 - val_accuracy: 0.4720
Epoch 20/20
56/56 [============ ] - 42s 759ms/step - loss: 0.9580 - accuracy: 0.6607 - val loss:
1.8658 - val accuracy: 0.4430
```

In [15]: # View the summary of all layers model.summary()

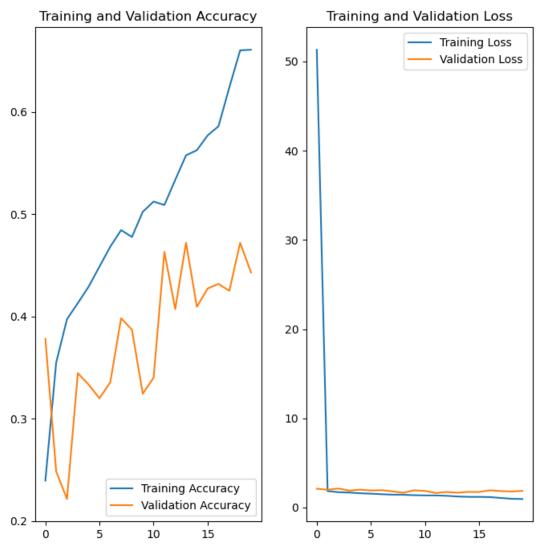
Model: "sequential"

| Layer (type) | Output Shape | Param # |
|--|--------------------------|---------|
| conv2d (Conv2D) | (None, 180, 180, 16) | |
| <pre>max_pooling2d (MaxPooling2D)</pre> | (None, 90, 90, 16) | 0 |
| dropout (Dropout) | (None, 90, 90, 16) | 0 |
| conv2d_1 (Conv2D) | (None, 90, 90, 32) | 4640 |
| <pre>max_pooling2d_1 (MaxPooling 2D)</pre> | (None, 45, 45, 32) | 0 |
| dropout_1 (Dropout) | (None, 45, 45, 32) | 0 |
| conv2d_2 (Conv2D) | (None, 45, 45, 64) | 18496 |
| <pre>max_pooling2d_2 (MaxPooling 2D)</pre> | (None, 22, 22, 64) | 0 |
| dropout_2 (Dropout) | (None, 22, 22, 64) | 0 |
| flatten (Flatten) | (None, 30976) | 0 |
| dense (Dense) | (None, 128) | 3965056 |
| dense_1 (Dense) | (None, 9) | 1161 |
| dropout_3 (Dropout) | (None, 9) | 0 |
| | | |

Total params: 3,989,801 Trainable params: 3,989,801 Non-trainable params: 0

Visualizing training results

```
In [16]: acc = history.history['accuracy']
         val_acc = history.history['val_accuracy']
         loss = history.history['loss']
         val_loss = history.history['val_loss']
         epochs_range = range(epochs)
         plt.figure(figsize=(8, 8))
         plt.subplot(1, 2, 1)
         plt.plot(epochs_range, acc, label='Training Accuracy')
         plt.plot(epochs_range, val_acc, label='Validation Accuracy')
         plt.legend(loc='lower right')
         plt.title('Training and Validation Accuracy')
         plt.subplot(1, 2, 2)
         plt.plot(epochs_range, loss, label='Training Loss')
         plt.plot(epochs_range, val_loss, label='Validation Loss')
         plt.legend(loc='upper right')
         plt.title('Training and Validation Loss')
         plt.show()
```



Todo: Write your findings after the model fit, see if there is an evidence of model overfit or underfit

Training accuracy is around 60% while Validation accuracy stops around 45%. Use of DropOut has reduced Overfitting. Need to boost validation accuracy using data augmentation. Image Rotation & flipping incorporated.

A Sample image with image rotation

```
In [18]: # Todo, visualize how your augmentation strategy works for one instance of training image.
         # Your code goes here
         #image = tf.cast(tf.expand_dims(image_batch, 0), tf.float32)
         plt.figure(figsize=(10, 10))
         for i in range(9):
           augmented_image = data_augmentation(image_batch)
           ax = plt.subplot(3, 3, i + 1)
           #plt.imshow(augmented_image[0])
           plt.imshow((augmented_image[0]).numpy().astype(np.uint8))
           plt.axis("off")
         WARNING:tensorflow:Using a while_loop for converting Bitcast cause there is no registered converter
         WARNING:tensorflow:Using a while loop for converting Bitcast cause there is no registered converter
         for this op.
         WARNING:tensorflow:Using a while_loop for converting StatelessRandomUniformV2 cause there is no regi
         stered converter for this op.
         WARNING:tensorflow:Using a while_loop for converting ImageProjectiveTransformV3 cause there is no re
         gistered converter for this op.
         WARNING:tensorflow:Using a while_loop for converting RngReadAndSkip cause there is no registered con
         verter for this op.
         WARNING:tensorflow:Using a while_loop for converting Bitcast cause there is no registered converter
         for this op.
         WARNING:tensorflow:Using a while_loop for converting Bitcast cause there is no registered converter
         for this op.
         WARNING:tensorflow:Using a while loop for converting StatelessRandomUniformV2 cause there is no regi
         stered converter for this op.
         WARNING:tensorflow:Using a while_loop for converting ImageProjectiveTransformV3 cause there is no re
         gistered converter for this op.
         WARNING:tensorflow:Using a while_loop for converting RngReadAndSkip cause there is no registered con
```

Todo:

Create the model, compile and train the model

```
In [19]: ## You can use Dropout Layer if there is an evidence of overfitting in your findings
         ## Your code goes here
         model = Sequential()
         model.add(Conv2D(16,3,padding="same", activation="relu"))
         model.add(MaxPooling2D())
         model.add(Conv2D(32, 3, padding="same", activation="relu"))
         model.add(MaxPooling2D())
         model.add(Conv2D(64, 3, padding="same", activation="relu"))
         model.add(MaxPooling2D())
         model.add(Dropout(0.2))
         model.add(Flatten())
         model.add(Dense(128,activation="relu"))
         model.add(Dense(num_classes))
```

Compiling the model

```
In [20]: ## Your code goes here
         model.compile(optimizer= 'adam',
                       loss=tf.keras.losses.SparseCategoricalCrossentropy(from_logits=True),
                       metrics=['accuracy'])
```

Training the model

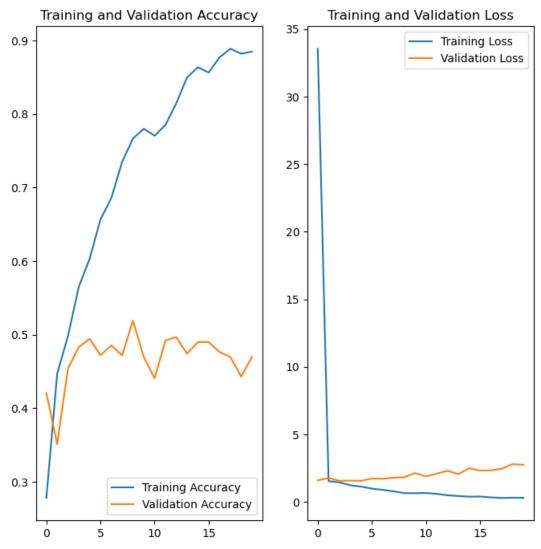
```
Epoch 1/20
1.6129 - val_accuracy: 0.4206
Epoch 2/20
56/56 [============= ] - 38s 683ms/step - loss: 1.5482 - accuracy: 0.4470 - val loss:
1.7805 - val accuracy: 0.3512
Epoch 3/20
56/56 [===================] - 37s 666ms/step - loss: 1.4616 - accuracy: 0.4983 - val_loss:
1.5813 - val_accuracy: 0.4541
Epoch 4/20
56/56 [============= ] - 37s 667ms/step - loss: 1.2441 - accuracy: 0.5653 - val loss:
1.5885 - val accuracy: 0.4832
Epoch 5/20
56/56 [====================] - 40s 710ms/step - loss: 1.1469 - accuracy: 0.6032 - val_loss:
1.5691 - val_accuracy: 0.4944
Epoch 6/20
1.7483 - val_accuracy: 0.4720
Epoch 7/20
1.7251 - val_accuracy: 0.4855
Epoch 8/20
56/56 [================ ] - 43s 773ms/step - loss: 0.7994 - accuracy: 0.7349 - val loss:
1.8041 - val_accuracy: 0.4720
Epoch 9/20
1.8401 - val accuracy: 0.5190
Epoch 10/20
2.1573 - val_accuracy: 0.4698
Epoch 11/20
56/56 [==============] - 38s 676ms/step - loss: 0.6751 - accuracy: 0.7706 - val_loss:
1.8979 - val_accuracy: 0.4407
Epoch 12/20
56/56 [==============] - 38s 671ms/step - loss: 0.6081 - accuracy: 0.7852 - val_loss:
2.1137 - val_accuracy: 0.4922
Epoch 13/20
2.3080 - val_accuracy: 0.4966
Epoch 14/20
56/56 [===============] - 41s 729ms/step - loss: 0.4501 - accuracy: 0.8499 - val_loss:
2.0700 - val_accuracy: 0.4743
Epoch 15/20
56/56 [=============] - 39s 700ms/step - loss: 0.4000 - accuracy: 0.8638 - val loss:
2.5104 - val_accuracy: 0.4899
Epoch 16/20
56/56 [============= ] - 39s 703ms/step - loss: 0.4119 - accuracy: 0.8566 - val loss:
2.3285 - val_accuracy: 0.4899
Epoch 17/20
56/56 [==============] - 41s 727ms/step - loss: 0.3507 - accuracy: 0.8772 - val_loss:
2.3516 - val_accuracy: 0.4765
Epoch 18/20
56/56 [================= ] - 40s 722ms/step - loss: 0.3045 - accuracy: 0.8890 - val loss:
2.4708 - val_accuracy: 0.4698
Epoch 19/20
56/56 [================= ] - 44s 787ms/step - loss: 0.3240 - accuracy: 0.8823 - val loss:
2.8071 - val_accuracy: 0.4430
Epoch 20/20
56/56 [============ ] - 44s 783ms/step - loss: 0.3168 - accuracy: 0.8850 - val loss:
2.7608 - val accuracy: 0.4698
```

```
In [22]: test_ds = tf.keras.preprocessing.image_dataset_from_directory(
           data_dir_test,
           seed=123,
           image_size = (img_height, img_width),
           batch_size = batch_size)
         loss, acc = model.evaluate(test_ds)
         print("Accuracy", acc)
```

```
Found 118 files belonging to 9 classes.
4/4 [=========== ] - 5s 164ms/step - loss: 5.7768 - accuracy: 0.2966
Accuracy 0.29661017656326294
```

Visualizing the results

```
In [23]: acc = history.history['accuracy']
         val_acc = history.history['val_accuracy']
         loss = history.history['loss']
         val_loss = history.history['val_loss']
         epochs_range = range(epochs)
         plt.figure(figsize=(8, 8))
         plt.subplot(1, 2, 1)
         plt.plot(epochs_range, acc, label='Training Accuracy')
         plt.plot(epochs_range, val_acc, label='Validation Accuracy')
         plt.legend(loc='lower right')
         plt.title('Training and Validation Accuracy')
         plt.subplot(1, 2, 2)
         plt.plot(epochs_range, loss, label='Training Loss')
         plt.plot(epochs_range, val_loss, label='Validation Loss')
         plt.legend(loc='upper right')
         plt.title('Training and Validation Loss')
         plt.show()
```



Todo: Write your findings after the model fit, see if there is an evidence of model overfit or underfit. Do you think there is some improvement now as compared to the previous model run?

Training accuracy has improved. But Validation accuracy is still low. So its still overfitting.

Todo: Find the distribution of classes in the training dataset.

Context: Many times real life datasets can have class imbalance, one class can have proportionately higher number of samples compared to the others. Class imbalance can have a detrimental effect on the final model quality. Hence as a sanity check it becomes important to check what is the distribution of classes in the data.

```
In [24]: ## Your code goes here.
         from glob import glob
         train_image_names = glob(r'./UpgradExer5CNNSkin/SkinImg/Train/*/*.jpg')
         print("Total number of training images: ", len(train_image_names))
         #print(train image names)
         # make train image names as serie object
         train_image_names = pd.Series(train_image_names)
         # train_df: a dataframe with 2 field: Filename, ClassId
         train_df = pd.DataFrame()
         # aenerate Filename field
         train_df['Filename'] = train_image_names.map(lambda img_name: img_name.split("\\")[-1])
         # generate ClassId field
         train_df['ClassId'] = train_image_names.map(lambda img_name: (img_name.split("\\")[-2]))
         train_df.head()
         class id distribution = train df['ClassId'].value counts()
         class_id_distribution.head(10)
         Total number of training images: 2239
```

```
Out[24]: pigmented benign keratosis 462
melanoma 438
basal cell carcinoma 376
nevus 357
squamous cell carcinoma 181
vascular lesion 139
actinic keratosis 114
dermatofibroma 95
```

Todo: Write your findings here:

Name: ClassId, dtype: int64

seborrheic keratosis

- Which class has the least number of samples?
- Which classes dominate the data in terms proportionate number of samples?

77

Least Samples in seborrheic keratosis 77

Following classes dominate

pigmented benign keratosis 462

melanoma 438

basal cell carcinoma 376

nevus 357

Todo: Rectify the class imbalance

Context: You can use a python package known as Augmentor (https://augmentor.readthedocs.io/en/master/ (https://augmentor.readthedocs.io/en/master/) to add more samples across all classes so that none of the classes have very few samples.

In [25]: #!pip install Augmentor

To use Augmentor , the following general procedure is followed:

- 1. Instantiate a Pipeline object pointing to a directory containing your initial image data set.
- 2. Define a number of operations to perform on this data set using your Pipeline object.
- 3. Execute these operations by calling the Pipeline's sample() method.

```
In [26]: path to training dataset="./UpgradExer5CNNSkin/SkinImg/Train/"
        import Augmentor
        for i in class names:
           p = Augmentor.Pipeline(path_to_training_dataset + i)
           p.rotate(probability=0.7, max_left_rotation=10, max_right_rotation=10)
           p.sample(500) ## We are adding 500 samples per class to make sure that none of the classes are sparse
        Initialised with 114 image(s) found.
        Output directory set to ./UpgradExer5CNNSkin/SkinImg/Train/actinic keratosis\output.
        Processing <PIL.Image.Image image mode=RGB size=600x450 at 0x1DC826DA3A0>: 100% | ■ | 500/500 [00:08<00:0
        0, 56.70 Samples/
        Initialised with 376 image(s) found.
        Output directory set to ./UpgradExer5CNNSkin/SkinImg/Train/basal cell carcinoma\output.
        Processing <PIL.JpegImagePlugin.JpegImageFile image mode=RGB size=600x450 at 0x1DC8265A430>: 100%  ■ 50
        0/500 [00:09<00:
        Initialised with 95 image(s) found.
        Output directory set to ./UpgradExer5CNNSkin/SkinImg/Train/dermatofibroma\output.
        Processing <PIL.Image.Image image mode=RGB size=600x450 at 0x1DC84399130>: 100% | ■ | 500/500 [00:08<00:0
        0, 55.97 Samples/
        Initialised with 438 image(s) found.
        Output directory set to ./UpgradExer5CNNSkin/SkinImg/Train/melanoma\output.
        0/500 [00:47<00:
        Initialised with 357 image(s) found.
        Output directory set to ./UpgradExer5CNNSkin/SkinImg/Train/nevus\output.
        00, 11.01 Sample
        Initialised with 462 image(s) found.
        Output directory set to ./UpgradExer5CNNSkin/SkinImg/Train/pigmented benign keratosis\output.
        0, 49.06 Samples/
        Initialised with 77 image(s) found.
        Output directory set to ./UpgradExer5CNNSkin/SkinImg/Train/seborrheic keratosis\output.
        Processing <PIL.Image.Image image mode=RGB size=1024x768 at 0x1DC842CB760>: 100%| | 500/500 [00:22<00:0
        0, 22.65 Samples
        Initialised with 181 image(s) found.
        Output directory set to ./UpgradExer5CNNSkin/SkinImg/Train/squamous cell carcinoma\output.
        Processing <PIL.Image.Image image mode=RGB size=600x450 at 0x1DC84359D30>: 100% | ■ 500/500 [00:09<00:0
        0, 53.13 Samples/
        Initialised with 139 image(s) found.
        Output directory set to ./UpgradExer5CNNSkin/SkinImg/Train/vascular lesion\output.
        Processing <PIL.Image.Image image mode=RGB size=600x450 at 0x1DC8432EEE0>: 100% | ■ | 500/500 [00:10<00:0
        0, 49.91 Samples/
```

Augmentor has stored the augmented images in the output sub-directory of each of the sub-directories of skin cancer types... Lets take a look at total count of augmented images.

```
In [27]: | image_count_train = len(list(data_dir_train.glob('*/output/*.jpg')))
         print(image_count_train)
```

4500

Lets see the distribution of augmented data after adding new images to the original training data.

```
In [28]: from glob import glob
    path_list = [x for x in glob(os.path.join(data_dir_train, '*', '*.jpg'))]
    path_list_new = [x for x in glob(os.path.join(data_dir_train, '*', 'output', '*.jpg'))]

In [29]: lesion_list = [os.path.basename((os.path.dirname(y))) for y in glob(os.path.join(data_dir_train, '*' lesion_list_new = [os.path.basename(os.path.dirname(os.path.dirname(y))) for y in glob(os.path.join(data_dir_train, '*')
    lesion_list_new = [os.path.basename(os.path.dirname(y))) for y in glob(os.path.join(data_dir_train, '*'
    lesion_list_new = [os.path.dirname(os.path.dirname(y))) for y in glob(os.path.join(data_dir_train, '*'
    lesion_list_new = [os.path.basename(os.path.dirname(y))) for y in glob(os.path.dirname(y)) for y in glob(os.path.join(data_dir_train, '*'
    lesion_list_new = [os.path.basename(os.path.dirname(y))) for y in glob(os.path.join(data_dir_train, '*'
    lesion_list_new = [os.path.basename(os.path.dirname(y
```

Distribution of data across classes including augmented data

```
In [32]: new_df['Label'].value_counts()
Out[32]: pigmented benign keratosis
                                         962
         melanoma
                                         938
         basal cell carcinoma
                                         876
         nevus
                                         857
         squamous cell carcinoma
                                        681
         vascular lesion
                                        639
         actinic keratosis
                                        614
         dermatofibroma
                                         595
         seborrheic keratosis
                                        577
         Name: Label, dtype: int64
```

So, now we have added 500 images to all the classes to maintain some class balance. We can add more images as we want to improve training process.

Todo: Train the model on the data created using Augmentor

```
In [33]: batch_size = 32
    img_height = 180
    img_width = 180
```

Todo: Create a training dataset

Found 6739 files belonging to 9 classes. Using 5392 files for training.

Todo: Create a validation dataset

```
In [35]: val_ds = tf.keras.preprocessing.image_dataset_from_directory(
    data_dir_train,
    seed=123,
    validation_split = 0.2,
    subset = 'validation', ## Todo choose the correct parameter value, so that only validation data is refer image_size=(img_height, img_width),
    batch_size=batch_size)
Found 6739 files belonging to 9 classes.
Using 1347 files for validation.
```

Todo: Create your model (make sure to include normalization)

```
In [36]: ## your code goes here

model = tf.keras.Sequential([
    tf.keras.layers.Rescaling(1./255),
    tf.keras.layers.Conv2D(32, 3, activation='relu'),
    tf.keras.layers.MaxPooling2D(),
    tf.keras.layers.Conv2D(32, 3, activation='relu'),
    tf.keras.layers.MaxPooling2D(),
    tf.keras.layers.Conv2D(32, 3, activation='relu'),
    tf.keras.layers.MaxPooling2D(),
    tf.keras.layers.MaxPooling2D(),
    tf.keras.layers.Flatten(),
    tf.keras.layers.Dense(128, activation='relu'),
    tf.keras.layers.Dense(num_classes)
])
```

Todo: Compile your model (Choose optimizer and loss function appropriately)

Todo: Train your model

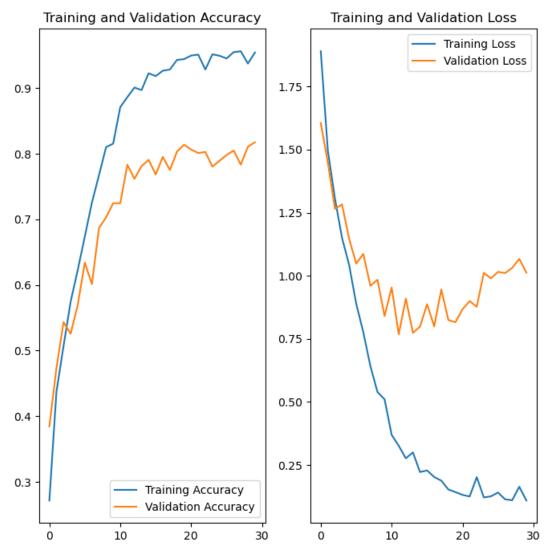
```
In [38]: epochs = 30
          history = model.fit(
train_ds,
             validation_data=val_ds,
             epochs=epochs
```

```
Epoch 1/30
s: 1.6057 - val_accuracy: 0.3846
Epoch 2/30
s: 1.4491 - val accuracy: 0.4736
Epoch 3/30
s: 1.2650 - val_accuracy: 0.5434
Epoch 4/30
s: 1.2830 - val_accuracy: 0.5256
Epoch 5/30
s: 1.1470 - val_accuracy: 0.5694
Epoch 6/30
s: 1.0483 - val_accuracy: 0.6340
Epoch 7/30
s: 1.0870 - val_accuracy: 0.6013
Epoch 8/30
s: 0.9604 - val_accuracy: 0.6867
Epoch 9/30
s: 0.9842 - val_accuracy: 0.7030
Epoch 10/30
s: 0.8404 - val_accuracy: 0.7246
Epoch 11/30
s: 0.9539 - val_accuracy: 0.7246
Epoch 12/30
s: 0.7684 - val_accuracy: 0.7832
Epoch 13/30
s: 0.9096 - val_accuracy: 0.7617
Epoch 14/30
s: 0.7740 - val_accuracy: 0.7810
Epoch 15/30
s: 0.7988 - val_accuracy: 0.7906
Epoch 16/30
s: 0.8876 - val_accuracy: 0.7684
Epoch 17/30
s: 0.7992 - val_accuracy: 0.7951
Epoch 18/30
s: 0.9459 - val_accuracy: 0.7751
Epoch 19/30
s: 0.8246 - val_accuracy: 0.8033
Epoch 20/30
s: 0.8164 - val_accuracy: 0.8137
Epoch 21/30
s: 0.8676 - val_accuracy: 0.8062
Epoch 22/30
s: 0.8999 - val_accuracy: 0.8010
Epoch 23/30
s: 0.8774 - val_accuracy: 0.8025
Epoch 24/30
```

```
s: 1.0118 - val_accuracy: 0.7803
Epoch 25/30
s: 0.9898 - val_accuracy: 0.7892
Epoch 26/30
s: 1.0156 - val accuracy: 0.7981
Epoch 27/30
s: 1.0109 - val_accuracy: 0.8048
Epoch 28/30
s: 1.0315 - val_accuracy: 0.7832
Epoch 29/30
s: 1.0671 - val_accuracy: 0.8107
Epoch 30/30
s: 1.0126 - val_accuracy: 0.8174
```

Todo: Visualize the model results

```
In [39]:
         acc = history.history['accuracy']
         val_acc = history.history['val_accuracy']
         loss = history.history['loss']
         val_loss = history.history['val_loss']
         epochs_range = range(epochs)
         plt.figure(figsize=(8, 8))
         plt.subplot(1, 2, 1)
         plt.plot(epochs_range, acc, label='Training Accuracy')
         plt.plot(epochs_range, val_acc, label='Validation Accuracy')
         plt.legend(loc='lower right')
         plt.title('Training and Validation Accuracy')
         plt.subplot(1, 2, 2)
         plt.plot(epochs_range, loss, label='Training Loss')
         plt.plot(epochs_range, val_loss, label='Validation Loss')
         plt.legend(loc='upper right')
         plt.title('Training and Validation Loss')
         plt.show()
```



Todo: Analyze your results here. Did you get rid of underfitting/overfitting? Did class rebalance help?

Rebalancing the class must have helped address underfitting of few classes with very less data (images).

Overall Overfitting is addressed as we see the Validation Accuracy has increased.

```
In [41]: from keras.preprocessing.image import ImageDataGenerator
    test_generator = ImageDataGenerator()
    test_data_generator = test_generator.flow_from_directory(
        data_dir_test , # Put your path here
            target_size=(img_width, img_height),
            batch_size=32,
            shuffle=False)
    test_steps_per_epoch = np.math.ceil(test_data_generator.samples / test_data_generator.batch_size)
    predictions = model.predict_generator(test_data_generator, steps=test_steps_per_epoch)
    # Get most Likely class
    predicted_classes = np.argmax(predictions, axis=1)
```

Found 118 images belonging to 9 classes.

C:\Users\admin\AppData\Local\Temp\ipykernel_16544\2016176624.py:10: UserWarning: `Model.predict_generat or` is deprecated and will be removed in a future version. Please use `Model.predict`, which supports g enerators.

predictions = model.predict_generator(test_data_generator, steps=test_steps_per_epoch)

```
In [42]: true_classes = test_data_generator.classes
    class_labels = list(test_data_generator.class_indices.keys())
```

```
In [43]: import sklearn.metrics as metrics
    report = metrics.classification_report(true_classes, predicted_classes, target_names=class_labels)
    print(report)
```

| | precision | recall | f1-score | support |
|----------------------------|-----------|--------|----------|---------|
| actinic keratosis | 0.00 | 0.00 | 0.00 | 16 |
| basal cell carcinoma | 0.39 | 0.44 | 0.41 | 16 |
| dermatofibroma | 0.67 | 0.12 | 0.21 | 16 |
| melanoma | 0.17 | 0.12 | 0.14 | 16 |
| nevus | 0.31 | 0.94 | 0.46 | 16 |
| pigmented benign keratosis | 0.39 | 0.56 | 0.46 | 16 |
| seborrheic keratosis | 0.00 | 0.00 | 0.00 | 3 |
| squamous cell carcinoma | 0.29 | 0.12 | 0.17 | 16 |
| vascular lesion | 0.67 | 0.67 | 0.67 | 3 |
| | | | | |
| accuracy | | | 0.33 | 118 |
| macro avg | 0.32 | 0.33 | 0.28 | 118 |
| weighted avg | 0.32 | 0.33 | 0.27 | 118 |

| In []: | |
|---------|--|
| | |
| In []: | |