```
from google.colab import drive
drive.mount('/content/gdrive')
 Mounted at /content/gdrive
!pwd
     /content
import matplotlib.pyplot as plt
import numpy as np
import pandas as pd
import os
from glob import glob
import seaborn as sns
from PIL import Image
from sklearn.metrics import confusion_matrix
import keras
from keras.utils.np_utils import to_categorical # used for converting labels to one-hot-encoding
from keras.models import Sequential
from keras.layers import Dense, Dropout, Flatten, Conv2D, MaxPool2D, BatchNormalization
from sklearn.model_selection import train_test_split
from scipy import stats
from sklearn.preprocessing import LabelEncoder
np.random.seed(42)
skin_df = pd.read_csv('/content/gdrive/MyDrive/IIITA Documents/M.Tech._2nd_SEM_Project_(mit2020068)/Datasets/HAM16
SIZE=32
# label encoding to numeric values from text
le = LabelEncoder()
le.fit(skin_df['dx'])
LabelEncoder()
print(list(le.classes_))
skin_df['label'] = le.transform(skin_df["dx"])
print(skin_df.sample(10))
     ['akiec', 'bcc', 'bkl', 'df', 'mel', 'nv', 'vasc']
    lesion_id image_id dx ...
1617 HAM_0007180 ISIC_0033272 mel ...
                                                          localization label
                                                 male
                                                                  face
     8128 HAM_0007195 ISIC_0031923 nv \dots female lower extremity
     2168 HAM_0001835 ISIC_0026652 mel ...
                                               male
                                                                  back
                                                                           4
     1090 HAM_0000465 ISIC_0030583 bkl ... female
                                                                           2
                                                                 trunk
     7754 HAM 0001720 ISIC 0034010
                                     nv ... male
                                                               abdomen
                                                                           5
    8071 HAM_0006333 ISIC_0024424 nv ...
                                                male
                                                                 trunk
                                                                           5
     7423 HAM_0004548 ISIC_0032832 nv ... female upper extremity
     8984 HAM_0006526 ISIC_0026671 nv ... male lower extremity
                                                                           5
     2310 HAM_0003102 ISIC_0032389 mel ...
                                                                           4
                                                male
                                                                  face
     7256 HAM_0004260 ISIC_0025525 nv ...
                                                 male
                                                                  back
                                                                           5
     [10 rows x 8 columns]
# Data distribution visualization
fig = plt.figure(figsize=(12,8))
ax1 = fig.add_subplot(221)
skin_df['dx'].value_counts().plot(kind='bar', ax=ax1)
ax1.set_ylabel('Count')
ax1.set_title('Cell Type');
```

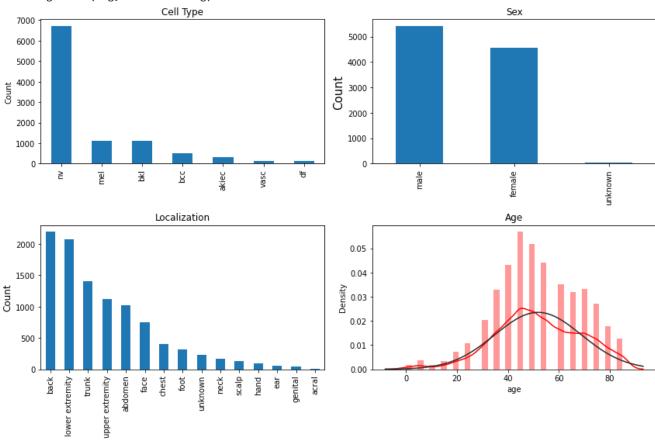
```
ax2 = fig.add_subplot(222)
skin_df['sex'].value_counts().plot(kind='bar', ax=ax2)
ax2.set_ylabel('Count', size=15)
ax2.set_title('Sex');

ax3 = fig.add_subplot(223)
skin_df['localization'].value_counts().plot(kind='bar')
ax3.set_ylabel('Count',size=12)
ax3.set_title('Localization')

ax4 = fig.add_subplot(224)
sample_age = skin_df[pd.notnull(skin_df['age'])]
sns.distplot(sample_age['age'], fit=stats.norm, color='red');
ax4.set_title('Age')

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

/usr/local/lib/python3.7/dist-packages/seaborn/distributions.py:2557: FutureWarning: `distplot` is a depreca warnings.warn(msg, FutureWarning)



```
# Distribution of data into various classes
from sklearn.utils import resample
print("label Frequency")
print(skin_df['label'].value_counts())
print()
print("Class Frequency")
print(skin_df['dx'].value_counts())
```

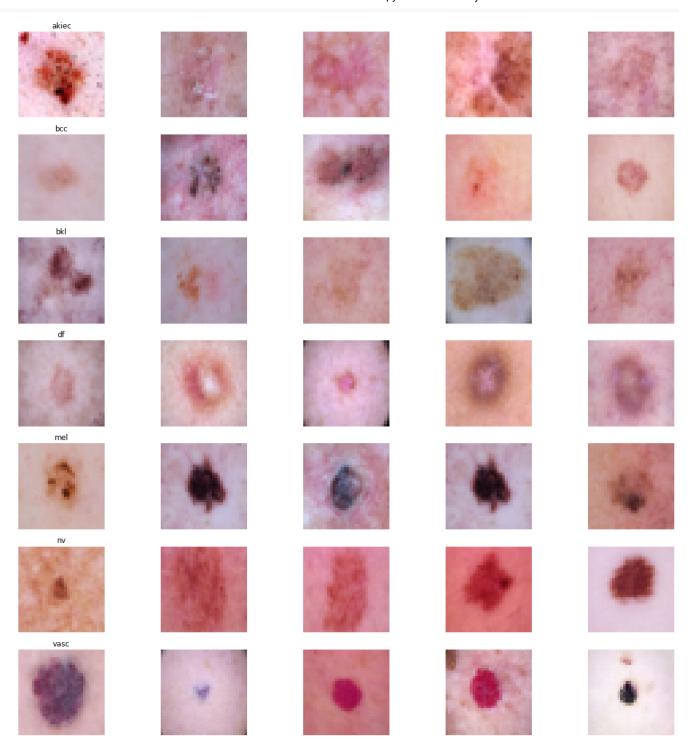
```
label Frequency
5
     6705
4
     1113
2
     1099
1
      514
0
       327
6
      142
3
      115
Name: label, dtype: int64
```

```
Class Frequency
nv
         6705
me1
         1113
bk1
         1099
          514
bcc
          327
akiec
vasc
          142
df
          115
Name: dx, dtype: int64
```

```
#Balance data.
df_0 = skin_df[skin_df['label'] == 0]
df_1 = skin_df[skin_df['label'] == 1]
df_2 = skin_df[skin_df['label'] == 2]
df_3 = skin_df[skin_df['label'] == 3]
df_4 = skin_df[skin_df['label'] == 4]
df_5 = skin_df[skin_df['label'] == 5]
df_6 = skin_df[skin_df['label'] == 6]
n_samples=500
df_0_balanced = resample(df_0, replace=True, n_samples=n_samples, random_state=42)
df_1_balanced = resample(df_1, replace=True, n_samples=n_samples, random_state=42)
df_2_balanced = resample(df_2, replace=True, n_samples=n_samples, random_state=42)
df_3_balanced = resample(df_3, replace=True, n_samples=n_samples, random_state=42)
\label{eq:df_dpalanced} $$ df_4$ palanced = resample(df_4$, replace=True, n_samples=n_samples, random_state=42) $$
df_5_balanced = resample(df_5, replace=True, n_samples=n_samples, random_state=42)
df_6_balanced = resample(df_6, replace=True, n_samples=n_samples, random_state=42)
#Combined back to a single dataframe
skin_df_balanced = pd.concat([df_0_balanced, df_1_balanced,
                               df_2_balanced, df_3_balanced,
                               df_4_balanced, df_5_balanced, df_6_balanced])
#Check the distribution. All classes should be balanced now.
print(skin_df_balanced['label'].value_counts())
          500
     5
     3
          500
     1
          500
     6
          500
     4
          500
     2
          500
     0
          500
     Name: label, dtype: int64
time to read images based on image ID from the CSV file
is the safest way to read images as it ensures the right image is read for the right ID
_path = {os.path.splitext(os.path.basename(x))[0]: x
                for x in glob(os.path.join('/content/gdrive/MyDrive/IIITA Documents/M.Tech._2nd_SEM_Project_(mit20)
#Define the path and add as a new column
skin_df_balanced['path'] = skin_df['image_id'].map(image_path.get)
#Use the path to read images.
skin_df_balanced['image'] = skin_df_balanced['path'].map(lambda x: np.asarray(Image.open(x).resize((SIZE,SIZE))))
n_samples = 5 # number of samples for plotting
# Plotting
fig, m_axs = plt.subplots(7, n_samples, figsize = (4*n_samples, 3*7))
for n_axs, (type_name, type_rows) in zip(m_axs,
                                          skin_df_balanced.sort_values(['dx']).groupby('dx')):
    n_axs[0].set_title(type_name)
    for c_ax, (_, c_row) in zip(n_axs, type_rows.sample(n_samples, random_state=1234).iterrows()):
```

c_ax.imshow(c_row['image'])

c_ax.axis('off')



[#]Convert dataframe column of images into numpy array
X = np.asarray(skin_df_balanced['image'].tolist())

X = X/255. # Scale values to 0-1. You can also used standardscaler or other scaling methods.

Y=skin_df_balanced['label'] #Assign label values to Y

Y_cat = to_categorical(Y, num_classes=7) #Convert to categorical as this is a multiclass classification problem

```
#Split to training and testing
x_train, x_test, y_train, y_test = train_test_split(X, Y_cat, test_size=0.25, random_state=42)
```

```
#Define the model.
num classes = 7
model = Sequential()
model.add(Conv2D(256, (3, 3), activation="relu", input_shape=(SIZE, SIZE, 3)))
#model.add(BatchNormalization())
model.add(MaxPool2D(pool_size=(2, 2)))
model.add(Dropout(0.3))
model.add(Conv2D(128, (3, 3),activation='relu'))
#model.add(BatchNormalization())
model.add(MaxPool2D(pool_size=(2, 2)))
model.add(Dropout(0.3))
model.add(Conv2D(64, (3, 3),activation='relu'))
#model.add(BatchNormalization())
model.add(MaxPool2D(pool_size=(2, 2)))
model.add(Dropout(0.3))
model.add(Flatten())
model.add(Dense(32))
model.add(Dense(7, activation='softmax'))
model.summary()
model.compile(loss='categorical_crossentropy', optimizer='Adam', metrics=['acc'])
```

Model: "sequential_1"

Layer (type)	Output Shape	Param #
conv2d_3 (Conv2D)	(None, 30, 30, 256)	7168
max_pooling2d_3 (MaxPooling2	(None, 15, 15, 256)	0
dropout_3 (Dropout)	(None, 15, 15, 256)	0
conv2d_4 (Conv2D)	(None, 13, 13, 128)	295040
max_pooling2d_4 (MaxPooling2	(None, 6, 6, 128)	0
dropout_4 (Dropout)	(None, 6, 6, 128)	0
conv2d_5 (Conv2D)	(None, 4, 4, 64)	73792
max_pooling2d_5 (MaxPooling2	(None, 2, 2, 64)	0
dropout_5 (Dropout)	(None, 2, 2, 64)	0
flatten_1 (Flatten)	(None, 256)	0
dense_2 (Dense)	(None, 32)	8224
dense 3 (Dense)	(None, 7)	231

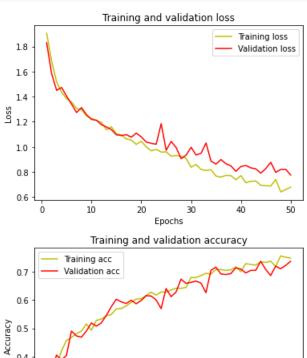
Total params: 384,455 Trainable params: 384,455 Non-trainable params: 0

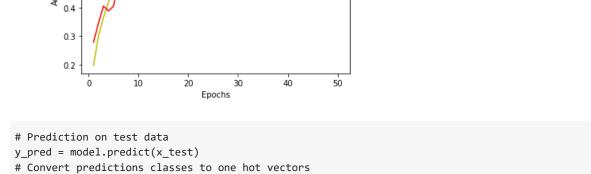
```
# Train
#You can also use generator to use augmentation during training.
batch_size = 16
epochs = 50
history = model.fit(
```

```
x_train, y_train,
    epochs=epochs,
    batch_size = batch_size,
    validation_data=(x_test, y_test),
    verbose=2)
score = model.evaluate(x test, y test)
print('Test accuracy:', score[1])
     165/165 - 22s - loss: 0.9683 - acc: 0.6274 - val_loss: 1.0277 - val_acc: 0.6137
     165/165 - 23s - loss: 0.9795 - acc: 0.6168 - val_loss: 1.0212 - val_acc: 0.6000
     Epoch 24/50
     165/165 - 22s - loss: 0.9582 - acc: 0.6278 - val_loss: 1.1848 - val_acc: 0.5691
     Epoch 25/50
     165/165 - 22s - loss: 0.9599 - acc: 0.6274 - val_loss: 0.9722 - val_acc: 0.6400
     Epoch 26/50
     165/165 - 22s - loss: 0.9256 - acc: 0.6347 - val loss: 1.0433 - val_acc: 0.6114
     Epoch 27/50
     165/165 - 22s - loss: 0.9315 - acc: 0.6411 - val_loss: 0.9946 - val_acc: 0.6274
     Epoch 28/50
     165/165 - 22s - loss: 0.9243 - acc: 0.6408 - val_loss: 0.9055 - val_acc: 0.6731
     Epoch 29/50
     165/165 - 23s - loss: 0.9086 - acc: 0.6438 - val_loss: 0.9335 - val_acc: 0.6583
     Epoch 30/50
     165/165 - 22s - loss: 0.8373 - acc: 0.6792 - val_loss: 0.9966 - val_acc: 0.6617
     Epoch 31/50
     165/165 - 23s - loss: 0.8593 - acc: 0.6785 - val_loss: 0.9349 - val_acc: 0.6663
     Epoch 32/50
     165/165 - 23s - loss: 0.8195 - acc: 0.6857 - val_loss: 0.9484 - val_acc: 0.6594
     Epoch 33/50
     165/165 - 22s - loss: 0.8113 - acc: 0.6937 - val_loss: 1.0306 - val_acc: 0.6251
     Epoch 34/50
     165/165 - 22s - loss: 0.8177 - acc: 0.6910 - val_loss: 0.8861 - val_acc: 0.7040
     Epoch 35/50
     165/165 - 22s - loss: 0.7668 - acc: 0.7070 - val loss: 0.8626 - val acc: 0.7154
     Epoch 36/50
     165/165 - 22s - loss: 0.7589 - acc: 0.7070 - val_loss: 0.8988 - val_acc: 0.6914
     Epoch 37/50
     165/165 - 22s - loss: 0.7724 - acc: 0.7040 - val_loss: 0.8652 - val_acc: 0.6891
     Epoch 38/50
     165/165 - 22s - loss: 0.7702 - acc: 0.7055 - val_loss: 0.8482 - val_acc: 0.6926
     Epoch 39/50
     165/165 - 22s - loss: 0.7373 - acc: 0.7162 - val_loss: 0.8046 - val_acc: 0.7131
     Epoch 40/50
     165/165 - 22s - loss: 0.7700 - acc: 0.6998 - val_loss: 0.8431 - val_acc: 0.7086
     Epoch 41/50
     165/165 - 22s - loss: 0.7137 - acc: 0.7276 - val_loss: 0.8511 - val_acc: 0.6949
     Epoch 42/50
     165/165 - 22s - loss: 0.7256 - acc: 0.7250 - val loss: 0.8323 - val acc: 0.7040
     Epoch 43/50
     165/165 - 23s - loss: 0.7274 - acc: 0.7223 - val_loss: 0.8241 - val_acc: 0.7040
     Epoch 44/50
     165/165 - 22s - loss: 0.6931 - acc: 0.7341 - val_loss: 0.7903 - val_acc: 0.7360
     Epoch 45/50
     165/165 - 22s - loss: 0.6908 - acc: 0.7318 - val_loss: 0.8270 - val_acc: 0.7074
     Epoch 46/50
     165/165 - 22s - loss: 0.6871 - acc: 0.7364 - val_loss: 0.8765 - val_acc: 0.6857
     Epoch 47/50
     165/165 - 21s - loss: 0.7406 - acc: 0.7181 - val loss: 0.7961 - val_acc: 0.7189
     Epoch 48/50
     165/165 - 22s - loss: 0.6403 - acc: 0.7550 - val_loss: 0.8198 - val_acc: 0.7097
     Epoch 49/50
     165/165 - 21s - loss: 0.6595 - acc: 0.7501 - val_loss: 0.8207 - val_acc: 0.7211
     Epoch 50/50
     165/165 - 21s - loss: 0.6781 - acc: 0.7474 - val loss: 0.7753 - val_acc: 0.7360
     28/28 [============= ] - 2s 61ms/step - loss: 0.7753 - acc: 0.7360
     Test accuracy: 0.7360000014305115
```

```
#plot the training and validation accuracy and loss at each epoch
loss = history.history['loss']
val_loss = history.history['val_loss']
epochs = range(1, len(loss) + 1)
```

```
plt.plot(epochs, loss, 'y', label='Training loss')
plt.plot(epochs, val_loss, 'r', label='Validation loss')
plt.title('Training and validation loss')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()
plt.show()
acc = history.history['acc']
val_acc = history.history['val_acc']
plt.plot(epochs, acc, 'y', label='Training acc')
plt.plot(epochs, val_acc, 'r', label='Validation acc')
plt.title('Training and validation accuracy')
plt.xlabel('Epochs')
plt.ylabel('Accuracy')
plt.legend()
plt.show()
```



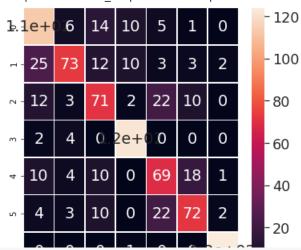


```
y_pred_classes = np.argmax(y_pred, axis = 1)
# Convert test data to one hot vectors
y_true = np.argmax(y_test, axis = 1)

#Print confusion matrix
cm = confusion_matrix(y_true, y_pred_classes)

fig, ax = plt.subplots(figsize=(6,6))
sns.set(font_scale=1.6)
sns.heatmap(cm, annot=True, linewidths=.5, ax=ax)
```

<matplotlib.axes._subplots.AxesSubplot at 0x7fd1b26968d0>



#PLot fractional incorrect misclassifications
incorr_fraction = 1 - np.diag(cm) / np.sum(cm, axis=1)
plt.bar(np.arange(7), incorr_fraction)
plt.xlabel('True Label')
plt.ylabel('Fraction of incorrect predictions')

Text(0, 0.5, 'Fraction of incorrect predictions')

