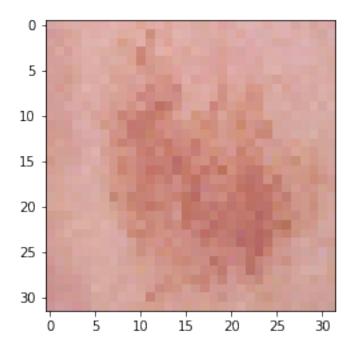
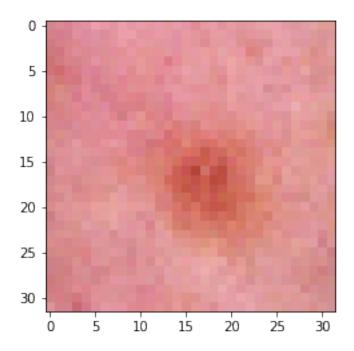
pro1

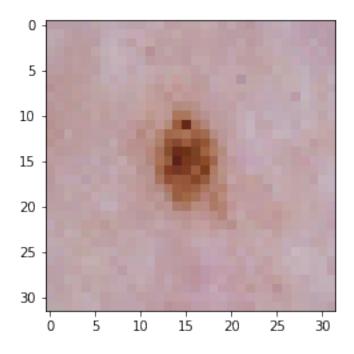
June 23, 2021

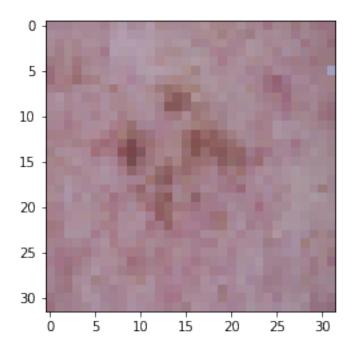
```
[37]: import pandas as pd
      import os
      import shutil
      import matplotlib.pyplot as plt
      import numpy as np
      from glob import glob
      import seaborn as sns
      from PIL import Image
      from multiprocessing import Queue
      np.random.seed(42)
      from sklearn.metrics import confusion_matrix
      import keras
      from keras.utils.np_utils import to_categorical # used for converting labels to_
      \hookrightarrow one-hot-encoding
      from keras.models import Sequential
      from keras.layers import Dense, Dropout, Flatten, Conv2D, MaxPooling2D,
      →BatchNormalization
      from keras import regularizers
      from keras.applications.resnet50 import ResNet50
      from tensorflow.keras.metrics import Recall
      from tensorflow.keras.callbacks import EarlyStopping,ReduceLROnPlateau
      from sklearn.model_selection import train_test_split
      from scipy import stats
      from sklearn.preprocessing import LabelEncoder
[38]: data_dir = os.getcwd() + "/all images"
[39]: dest_dir = os.getcwd() + "/organized"
[40]: skin_df2 = pd.read_csv('/home/coder/Desktop/project/skin disease/
       →HAM10000_metadata.csv')
      skin_df2['dx'].value_counts()
[40]: nv
               6705
     mel
               1113
```

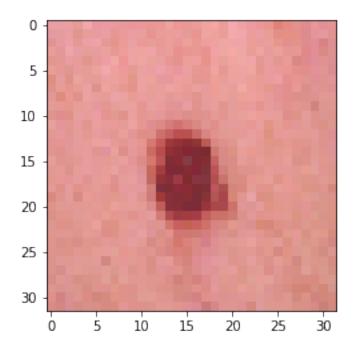
```
1099
     bkl
                514
      bcc
      akiec
                327
                142
      vasc
      df
                115
     Name: dx, dtype: int64
[41]: label=skin_df2['dx'].unique().tolist()
      label_images=[]
[42]: # Copy images to new folders
      for i in label:
          os.mkdir(dest_dir + str(i) + "/")
          sample = skin_df2[skin_df2['dx'] == i]['image_id']
          label_images.extend(sample)
          for id in label_images:
              shutil.copyfile((data_dir + "/"+ id +".jpg"), (dest_dir + i + "/"+id+".
       →jpg"))
          label_images=[]
[44]: from keras.preprocessing.image import ImageDataGenerator
      import os
      from matplotlib import pyplot as plt
[45]: datagen = ImageDataGenerator()
[46]: train_dir = os.getcwd() + "/organized"
      #USe flow_from_directory
      train_data_keras = datagen.flow_from_directory(directory=train_dir,
                                               class_mode='categorical',
                                               batch_size=16, #16 images at a time
                                               target_size=(32,32)) #Resize images
     Found 10015 images belonging to 7 classes.
[47]: x, y = next(train_data_keras)
[48]: for i in range (0,15):
          image = x[i].astype(int)
          plt.imshow(image)
          plt.show()
```

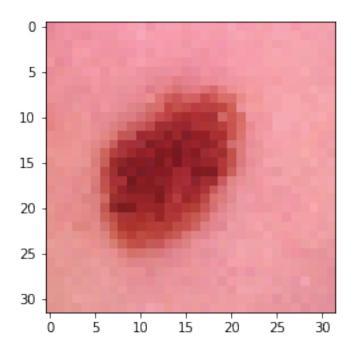


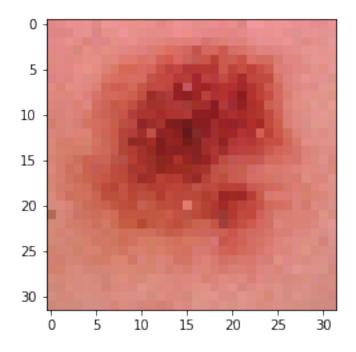


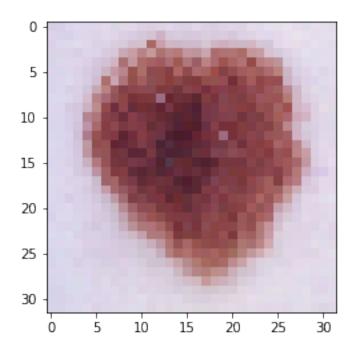


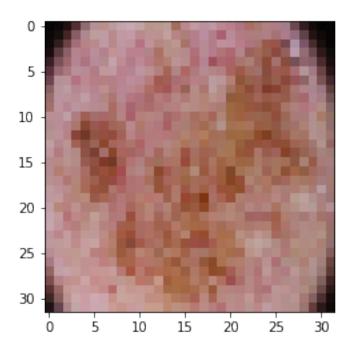


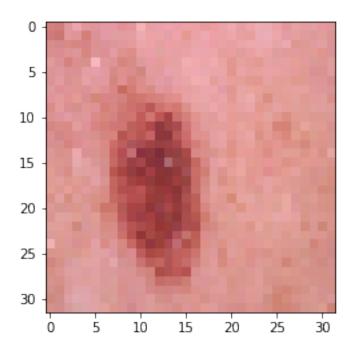


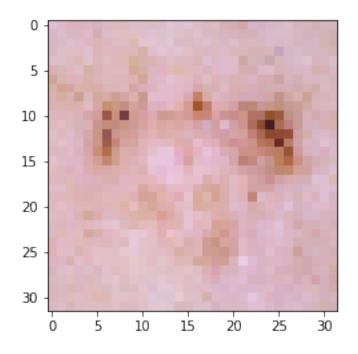


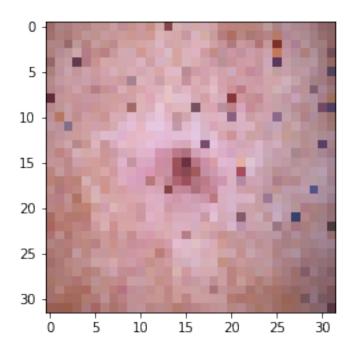


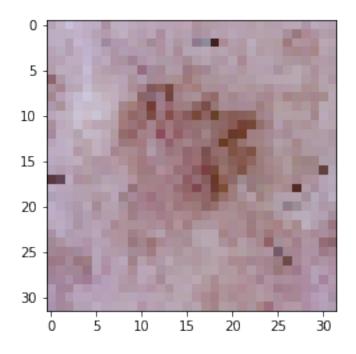


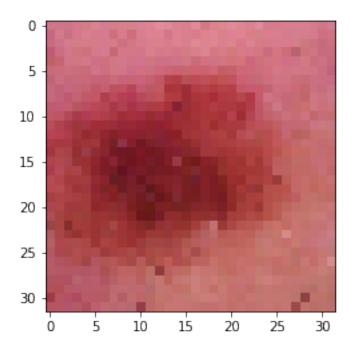


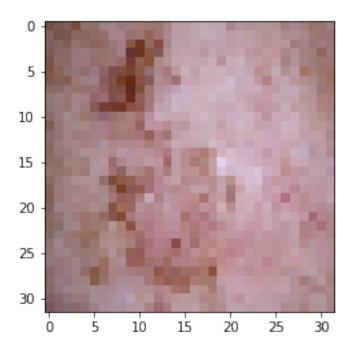






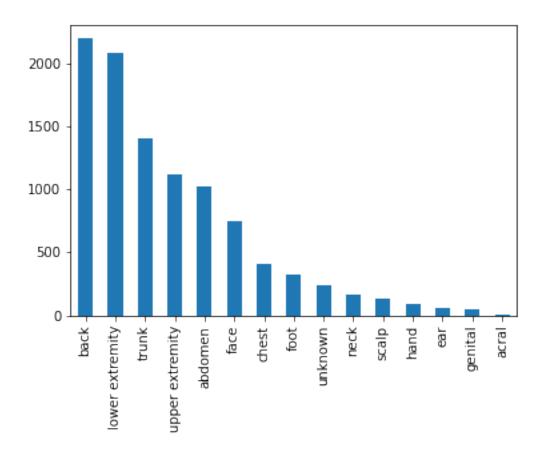






```
[49]: SIZE=32
      # label encoding to numeric values from text
      le = LabelEncoder()
      le.fit(skin_df2['dx'])
      LabelEncoder()
      print(list(le.classes_))
     ['akiec', 'bcc', 'bkl', 'df', 'mel', 'nv', 'vasc']
[50]: skin df2['label'] = le.transform(skin df2["dx"])
      print(skin_df2.sample(10))
              lesion_id
                             image_id
                                         dx
                                               dx_type
                                                         age
                                                                   sex
                         ISIC_0033631
           HAM 0000948
     971
                                       bkl
                                            consensus
                                                         NaN
                                                              unknown
           HAM_0005865
     606
                         ISIC_0031522
                                       bkl
                                                 histo
                                                        70.0
                                                                  male
           HAM_0005116
                         ISIC_0032953
                                                         5.0
                                                               female
     7198
                                        nv
                                                 histo
           HAM_0003015
     695
                         ISIC_0025083
                                       bkl
                                                 histo
                                                        55.0
                                                                  male
     787
           HAM_0002042
                         ISIC_0028294
                                       bkl
                                              confocal
                                                        75.0
                                                               female
           HAM_0004869
                         ISIC_0025370
                                                               female
     6936
                                                 histo
                                                        15.0
                                        nv
     6292
           HAM_0004162
                         ISIC_0031944
                                             follow_up
                                                        65.0
                                                                  male
                                        nv
           HAM_0006553
                         ISIC_0026942
                                             consensus
                                                               female
     9323
                                                        20.0
                                        nv
           HAM_0007306
     7480
                         ISIC_0026810
                                                        30.0
                                                                  male
                                                 histo
                                        nv
           HAM_0004518
     8886
                         ISIC_0024578
                                                        35.0
                                                               female
                                                 histo
                                        nv
               localization label
     971
                    unknown
```

```
606
                     chest
                                 2
     7198 lower extremity
                                 5
     695
                      back
                                 2
     787
                      face
                                 2
                                 5
     6936 lower extremity
     6292
                      back
                                 5
                      face
                                 5
     9323
                                 5
     7480
                      back
     8886 upper extremity
                                 5
[51]: fig = plt.figure(figsize=(12,8))
     <Figure size 864x576 with 0 Axes>
[52]: ax1 = fig.add_subplot(221)
      skin_df2['dx'].value_counts().plot(kind='bar', ax=ax1)
      ax1.set_ylabel('Count')
      ax1.set_title('Cell Type')
[52]: Text(0.5, 1.0, 'Cell Type')
[53]: ax2 = fig.add_subplot(222)
      skin_df2['sex'].value_counts().plot(kind='bar', ax=ax2)
      ax2.set_ylabel('Count', size=15)
      ax2.set_title('Sex')
[53]: Text(0.5, 1.0, 'Sex')
[54]: ax3 = fig.add_subplot(223)
      skin_df2['localization'].value_counts().plot(kind='bar')
      ax3.set_ylabel('Count',size=12)
      ax3.set_title('Localization')
[54]: Text(0.5, 1.0, 'Localization')
```

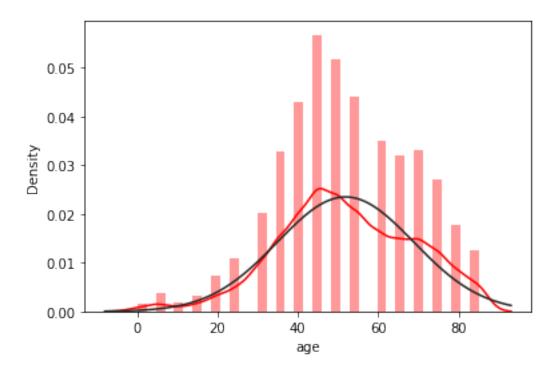


```
[55]: ax4 = fig.add_subplot(224)
sample_age = skin_df2[pd.notnull(skin_df2['age'])]
sns.distplot(sample_age['age'], fit=stats.norm, color='red');
ax4.set_title('Age')
```

/home/coder/anaconda3/lib/python3.8/site-packages/seaborn/distributions.py:2551: FutureWarning: `distplot` is a deprecated function and will be removed in a future version. Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).

warnings.warn(msg, FutureWarning)

[55]: Text(0.5, 1.0, 'Age')



```
[56]: plt.tight_layout()
      plt.show()
     <Figure size 432x288 with 0 Axes>
[57]: from sklearn.utils import resample
      print(skin_df2['label'].value_counts())
     5
          6705
     4
          1113
     2
          1099
           514
     1
     0
           327
     6
           142
     3
           115
     Name: label, dtype: int64
[58]: df_0 = skin_df2[skin_df2['label'] == 0]
      df_1 = skin_df2[skin_df2['label'] == 1]
      df_2 = skin_df2[skin_df2['label'] == 2]
      df_3 = skin_df2[skin_df2['label'] == 3]
      df_4 = skin_df2[skin_df2['label'] == 4]
      df_5 = skin_df2[skin_df2['label'] == 5]
      df_6 = skin_df2[skin_df2['label'] == 6]
```

```
[59]: 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,_u
       \rightarrowrandom_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,_u
      →random_state=42)
      df_4_balanced = resample(df_4, replace=True, n_samples=n_samples,_u
      →random_state=42)
      df_5_balanced = resample(df_5, replace=True, n_samples=n_samples,_u
      →random_state=42)
      df_6_balanced = resample(df_6, replace=True, n_samples=n_samples,_u
       →random_state=42)
[60]: 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])
[61]: print(skin_df_balanced['label'].value_counts())
     5
          500
          500
     3
          500
     1
          500
     6
     4
          500
     2
          500
          500
     0
     Name: label, dtype: int64
[62]: | image_path = {os.path.splitext(os.path.basename(x))[0]: x
                           for x in glob(os.path.join('/home/coder/Desktop/project/
       →skin disease', '*', '*.jpg'))}
[63]: | skin_df_balanced['path'] = skin_df2['image_id'].map(image_path.get)
[64]: | skin_df_balanced['image'] = skin_df_balanced['path'].map(lambda x: np.
      →asarray(Image.open(x).resize((SIZE,SIZE))))
      \# skin_df_balanced['image'] = lambda xs: K.reshape(xs[0], [xs[1][0], xs[1][1]],
       →output_shape=(SIZE,SIZE))([x,skin_df_balanced['image']])
[65]: n_{samples} = 5
      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')
```



```
[66]: 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_\( \infty multiclass classification problem\)
#Split to training and testing
x_train, x_test, y_train, y_test = train_test_split(X, Y_cat, test_size=0.028,\( \infty \) random_state=42)
```

```
[71]: input_shape=(32,32,3)
      model=Sequential()
      model.add(Conv2D(64,(2,2),input_shape=(32,32,3),activation='relu'))
      model.add(MaxPooling2D(pool_size=(2, 2)))
      model.add(BatchNormalization())
     model.add(Conv2D(512,(2,2),input_shape=(32,32,3),activation='relu'))
      model.add(MaxPooling2D(pool_size=(2, 2)))
      model.add(Dropout(0.3))
      model.add(Conv2D(1024,(2,2),input_shape=(32,32,3),activation='relu'))
      model.add(MaxPooling2D(pool_size=(2, 2)))
      model.add(BatchNormalization())
      model.add(Dropout(0.4))
      model.add(Conv2D(1024,(1,1),input_shape=(32,32,3),activation='relu'))
      model.add(MaxPooling2D(pool size=(1, 1)))
      model.add(Dropout(0.4))
      model.add(Flatten())
      model.add(Dense(256,activation='relu',kernel_regularizer=regularizers.12(0.01)))
      model.add(Dropout(0.5))
      model.add(Dense(7,activation='softmax'))
       →compile(optimizer='adam',loss='categorical_crossentropy',metrics=['accuracy',Recall()])
```

[72]: model.summary()

Model:	"sequential	5"
--------	-------------	----

Layer (type)	Output Shape	Param #
conv2d_20 (Conv2D)	(None, 31, 31, 64)	832
max_pooling2d_20 (MaxPooling	(None, 15, 15, 64)	0
batch_normalization_10 (Batc	(None, 15, 15, 64)	256
conv2d_21 (Conv2D)	(None, 14, 14, 512)	131584
max_pooling2d_21 (MaxPooling	(None, 7, 7, 512)	0
dropout_20 (Dropout)	(None, 7, 7, 512)	0
conv2d_22 (Conv2D)	(None, 6, 6, 1024)	2098176
max_pooling2d_22 (MaxPooling	(None, 3, 3, 1024)	0
batch_normalization_11 (Batc	(None, 3, 3, 1024)	4096
dropout_21 (Dropout)	(None, 3, 3, 1024)	0
conv2d_23 (Conv2D)	(None, 3, 3, 1024)	1049600
max_pooling2d_23 (MaxPooling	(None, 3, 3, 1024)	0
dropout_22 (Dropout)	(None, 3, 3, 1024)	0
flatten_5 (Flatten)	(None, 9216)	0
dense_10 (Dense)	(None, 256)	2359552
dropout_23 (Dropout)	(None, 256)	0
dense_11 (Dense)	(None, 7)	1799
Total mamma, E 64E 90E		

Total params: 5,645,895 Trainable params: 5,643,719 Non-trainable params: 2,176

[33]: early=EarlyStopping(monitor='accuracy',patience=4,mode='auto')

```
reduce_lr = ReduceLROnPlateau(monitor='accuracy', factor=0.5, patience=2, u overbose=1,cooldown=0, mode='auto',min_delta=0.0001, min_lr=0)
```

[34]: history=model. →fit(x_train,y_train,epochs=50,batch_size=90,validation_data=(x_test,_ →y_test),callbacks=[early,reduce_lr])

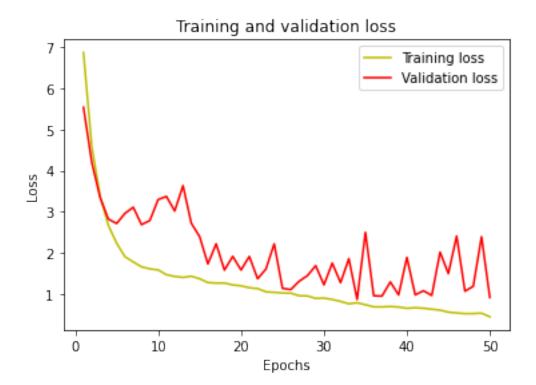
```
Epoch 1/50
0.2419 - recall: 0.0629 - val_loss: 5.5418 - val_accuracy: 0.1429 - val_recall:
0.0000e+00
Epoch 2/50
0.3075 - recall: 0.0841 - val_loss: 4.2043 - val_accuracy: 0.1429 - val_recall:
0.0000e+00
Epoch 3/50
38/38 [============== ] - 93s 2s/step - loss: 3.3887 - accuracy:
0.3471 - recall: 0.1264 - val_loss: 3.3529 - val_accuracy: 0.1633 - val_recall:
0.0000e+00
Epoch 4/50
38/38 [=============== ] - 91s 2s/step - loss: 2.6701 - accuracy:
0.3810 - recall: 0.1487 - val_loss: 2.8259 - val_accuracy: 0.1429 - val_recall:
0.0000e+00
Epoch 5/50
38/38 [============== ] - 83s 2s/step - loss: 2.2391 - accuracy:
0.4068 - recall: 0.1796 - val_loss: 2.7171 - val_accuracy: 0.1837 - val_recall:
0.0000e+00
Epoch 6/50
0.4606 - recall: 0.2287 - val_loss: 2.9629 - val_accuracy: 0.1429 - val_recall:
0.0000e+00
Epoch 7/50
0.4636 - recall: 0.2378 - val loss: 3.1126 - val accuracy: 0.1939 - val recall:
0.1837
Epoch 8/50
0.4774 - recall: 0.2716 - val_loss: 2.6893 - val_accuracy: 0.1429 - val_recall:
0.0000e+00
Epoch 9/50
38/38 [=============== ] - 82s 2s/step - loss: 1.6139 - accuracy:
0.4979 - recall: 0.2851 - val_loss: 2.7890 - val_accuracy: 0.1224 - val_recall:
0.0204
Epoch 10/50
0.5100 - recall: 0.2969 - val_loss: 3.2974 - val_accuracy: 0.2755 - val_recall:
0.0306
```

```
Epoch 11/50
0.5300 - recall: 0.3272 - val loss: 3.3762 - val accuracy: 0.2551 - val recall:
0.1735
Epoch 12/50
38/38 [=============== ] - 55s 1s/step - loss: 1.4314 - accuracy:
0.5456 - recall: 0.3360 - val_loss: 3.0216 - val_accuracy: 0.2143 - val_recall:
0.0816
Epoch 13/50
0.5626 - recall: 0.3595 - val loss: 3.6369 - val accuracy: 0.1939 - val recall:
0.1224
Epoch 14/50
0.5755 - recall: 0.3798 - val_loss: 2.7244 - val_accuracy: 0.2551 - val_recall:
0.1735
Epoch 15/50
0.5879 - recall: 0.4077 - val_loss: 2.4006 - val_accuracy: 0.3061 - val_recall:
0.2245
Epoch 16/50
0.6123 - recall: 0.4342 - val_loss: 1.7348 - val_accuracy: 0.4082 - val_recall:
0.3163
Epoch 17/50
0.6314 - recall: 0.4647 - val_loss: 2.2243 - val_accuracy: 0.3673 - val_recall:
0.2959
Epoch 18/50
38/38 [=============== ] - 70s 2s/step - loss: 1.2693 - accuracy:
0.6308 - recall: 0.4644 - val_loss: 1.5848 - val_accuracy: 0.4796 - val_recall:
0.3673
Epoch 19/50
0.6455 - recall: 0.4871 - val loss: 1.9190 - val accuracy: 0.3980 - val recall:
0.3571
Epoch 20/50
0.6599 - recall: 0.5085 - val_loss: 1.5868 - val_accuracy: 0.5714 - val_recall:
0.4592
Epoch 21/50
0.6822 - recall: 0.5317 - val_loss: 1.9164 - val_accuracy: 0.4490 - val_recall:
0.3571
Epoch 22/50
38/38 [=============== ] - 55s 1s/step - loss: 1.1354 - accuracy:
0.6796 - recall: 0.5603 - val_loss: 1.3746 - val_accuracy: 0.6633 - val_recall:
0.4898
```

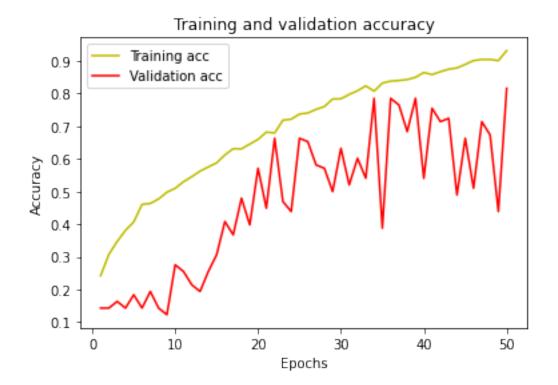
```
Epoch 23/50
0.7187 - recall: 0.6026 - val loss: 1.6091 - val accuracy: 0.4694 - val recall:
0.3980
Epoch 24/50
38/38 [=============== ] - 55s 1s/step - loss: 1.0444 - accuracy:
0.7219 - recall: 0.6035 - val_loss: 2.2228 - val_accuracy: 0.4388 - val_recall:
0.3265
Epoch 25/50
38/38 [=============== ] - 55s 1s/step - loss: 1.0276 - accuracy:
0.7375 - recall: 0.6364 - val loss: 1.1418 - val accuracy: 0.6633 - val recall:
0.5918
Epoch 26/50
38/38 [================= ] - 56s 1s/step - loss: 1.0239 - accuracy:
0.7407 - recall: 0.6476 - val_loss: 1.1114 - val_accuracy: 0.6531 - val_recall:
0.5816
Epoch 27/50
38/38 [=============== ] - 55s 1s/step - loss: 0.9616 - accuracy:
0.7519 - recall: 0.6678 - val_loss: 1.3127 - val_accuracy: 0.5816 - val_recall:
0.5102
Epoch 28/50
38/38 [================== ] - 55s 1s/step - loss: 0.9576 - accuracy:
0.7604 - recall: 0.6764 - val_loss: 1.4491 - val_accuracy: 0.5714 - val_recall:
0.5000
Epoch 29/50
38/38 [=============== ] - 55s 1s/step - loss: 0.8970 - accuracy:
0.7837 - recall: 0.7078 - val_loss: 1.6921 - val_accuracy: 0.5000 - val_recall:
0.4184
Epoch 30/50
38/38 [=============== ] - 55s 1s/step - loss: 0.9041 - accuracy:
0.7842 - recall: 0.7099 - val_loss: 1.2262 - val_accuracy: 0.6327 - val_recall:
0.5816
Epoch 31/50
38/38 [=============== ] - 55s 1s/step - loss: 0.8704 - accuracy:
0.7978 - recall: 0.7316 - val loss: 1.7539 - val accuracy: 0.5204 - val recall:
0.4796
Epoch 32/50
38/38 [================= ] - 55s 1s/step - loss: 0.8238 - accuracy:
0.8089 - recall: 0.7399 - val_loss: 1.2800 - val_accuracy: 0.6020 - val_recall:
0.5714
Epoch 33/50
38/38 [=============== ] - 55s 1s/step - loss: 0.7651 - accuracy:
0.8239 - recall: 0.7704 - val_loss: 1.8625 - val_accuracy: 0.5408 - val_recall:
0.5204
Epoch 34/50
38/38 [================ ] - 55s 1s/step - loss: 0.7912 - accuracy:
0.8078 - recall: 0.7504 - val_loss: 0.8719 - val_accuracy: 0.7857 - val_recall:
0.7245
```

```
Epoch 35/50
accuracy: 0.8325 - recall: 0.7725 - val_loss: 2.5024 - val_accuracy: 0.3878 -
val_recall: 0.3776
Epoch 36/50
0.8386 - recall: 0.7907 - val_loss: 0.9617 - val_accuracy: 0.7857 - val_recall:
0.6837
Epoch 37/50
0.8404 - recall: 0.7922 - val loss: 0.9535 - val accuracy: 0.7653 - val recall:
0.6837
Epoch 38/50
0.8433 - recall: 0.8013 - val_loss: 1.2980 - val_accuracy: 0.6837 - val_recall:
0.6837
Epoch 39/50
38/38 [=============== ] - 43s 1s/step - loss: 0.6883 - accuracy:
0.8501 - recall: 0.8113 - val_loss: 0.9811 - val_accuracy: 0.7857 - val_recall:
0.7653
Epoch 40/50
0.8648 - recall: 0.8280 - val_loss: 1.8942 - val_accuracy: 0.5408 - val_recall:
0.5000
Epoch 41/50
38/38 [=============== ] - 45s 1s/step - loss: 0.6700 - accuracy:
0.8586 - recall: 0.8154 - val_loss: 0.9828 - val_accuracy: 0.7551 - val_recall:
0.6939
Epoch 42/50
0.8674 - recall: 0.8266 - val_loss: 1.0834 - val_accuracy: 0.7143 - val_recall:
0.6429
Epoch 43/50
0.8751 - recall: 0.8389 - val loss: 0.9683 - val accuracy: 0.7245 - val recall:
0.7143
Epoch 44/50
38/38 [=================== ] - 43s 1s/step - loss: 0.6117 - accuracy:
0.8789 - recall: 0.8445 - val_loss: 2.0208 - val_accuracy: 0.4898 - val_recall:
0.4694
Epoch 45/50
0.8898 - recall: 0.8624 - val_loss: 1.4999 - val_accuracy: 0.6633 - val_recall:
0.6122
Epoch 46/50
0.9015 - recall: 0.8762 - val_loss: 2.4124 - val_accuracy: 0.5102 - val_recall:
0.4796
```

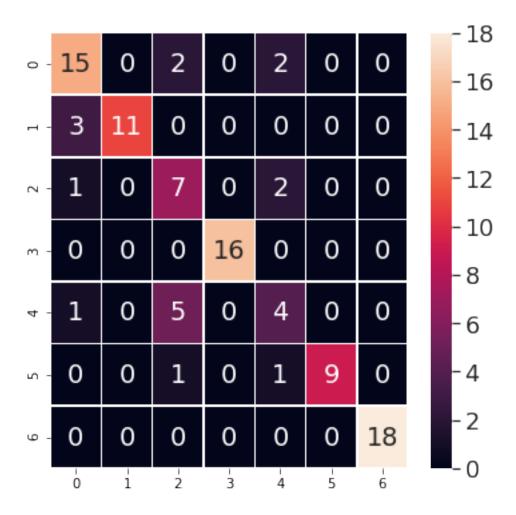
```
Epoch 47/50
   0.9048 - recall: 0.8812 - val loss: 1.0716 - val accuracy: 0.7143 - val recall:
   0.6939
   Epoch 48/50
   38/38 [=============== ] - 71s 2s/step - loss: 0.5286 - accuracy:
   0.9048 - recall: 0.8854 - val_loss: 1.1920 - val_accuracy: 0.6735 - val_recall:
   0.6224
   Epoch 49/50
   0.9015 - recall: 0.8777
   Epoch 00049: ReduceLROnPlateau reducing learning rate to 0.00050000000237487257.
   0.9015 - recall: 0.8777 - val loss: 2.3961 - val accuracy: 0.4388 - val recall:
   0.4082
   Epoch 50/50
   0.9321 - recall: 0.9165 - val loss: 0.9191 - val accuracy: 0.8163 - val recall:
   0.7959
[35]: score = model.evaluate(x_test, y_test)
    print('Test accuracy:', score[1])
   0.8163 - recall: 0.7959
   Test accuracy: 0.8163265585899353
[36]: 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()
```



```
[37]: acc = history.history['accuracy']
    val_acc = history.history['val_accuracy']
    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()
```

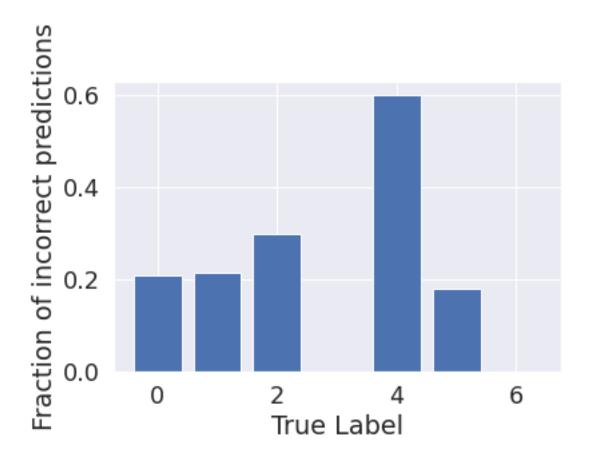


[42]: <AxesSubplot:>



```
[43]: 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')
```

[43]: Text(0, 0.5, 'Fraction of incorrect predictions')



```
[6]: model = ResNet50(weights='imagenet')
    model.save('/home/coder/Desktop/project/models')

INFO:tensorflow:Assets written to: /home/coder/Desktop/project/models/assets

[7]: model.save('model_resnet50.h5')
[]:
```

```
from __future__ import division, print_function
import sys
import os
import os.path
import glob
import re
import numpy as np
from PIL import Image
from keras.applications.imagenet utils import
preprocess_input,decode_predictions
from keras.models import load model
from keras.preprocessing import image
# from keras.applications.resnet50 import decode predictions
from flask import Flask
# from flask.ext.sqlalchemy import SQLAlchemy
from flask import redirect, url for, request, render template
from werkzeug.utils import secure filename
from gevent.pywsgi import WSGIServer
# # Actinic Keratoses
# # Basal cell carcinoma
# Benign keratosis
# Dermatofibroma
# # Melanoma
# # Melanocytic nevi
0.akiec
1.bcc
2.bkl
3.df
# # Dermatofibroma
5.nv
# # Melanocytic nevi
# # Vascular
                                 6.vasc
# app = Flask( name )
MODEL PATH = "/home/coder/Desktop/project/models/model resnet50.h5"
mymodel = load model(MODEL PATH)
mymodel.make predict function()
import os
from flask import request
from flask import Flask
from flask import render template
import PIL
app = Flask(__name__,static_url_path='/static')
MODEL PATH = "/home/coder/Desktop/project/models/model.h5"
@app.route('/',methods=["GET","POST"])
def upload pred():
    if request.method=="POST":
         image file = request.files["image"]
```

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```
54
            predict="No file uploaded"
 55
            if image file:
 56
                base path = os.path.dirname(os.path.abspath( file ))
 57
                file path = os.path.join(base path, 'uploads', image file.filename)
 58
            #
                validator
 59
                # chk , not file = is valid file(file path)
 60
                # if chk:
 61
                image file.save(file path)
 62
                predict = model predict(file path)
 63
                # else:
 64
                # predict="You Uploaded a " + not file + " file !!!"
            return render template("index.html",prediction=predict)
 65
 66
        return render_template("index.html",prediction="Upload Image")
 67
 68
 69 def model predict(file path):
       MODEL PATH = "/home/coder/Desktop/project/models/model.h5"
 70
 71
        mymodel = load model(MODEL PATH)
 72
        mymodel.make predict function()
 73
        class name = ["Actinic keratoses", "Basal cell carcinoma", "Benign
   keratosis-like lesions", "Dernatofibroma", "Vascular
   lesions", "Melanoma", "Vascular"]
 74
        img = image.load img(file path, target size=(32,32))
 75
        data = np.ndarray(shape=(1, 32, 32, 3), dtype=np.float32)
 76
        x=image.img to array(img)
 77
        normalized_image_array = (x.astype(np.float32) / 132.0) - 1
 78
        data[0] = normalized image array
 79
        predict1 = (mymodel.predict(data))*100000000000000
 80
       print(predict1)
        index_max = np.argmax(predict1)
 81
 82
        print(index max)
 83
        return class name[index max]
 84
 85 # def is valid file(file path):
 86
 87 #
          validity=True
 88 #
          not file=""
 89
90 #
          if file path==None :
 91 #
              return False
 92
93 #
          file path = str(file path)
 94
          file_extension_list = list( map(str , file_path.split(".")) )
 95 #
 96 #
          allowed extension = ["jpg" , "png" , "webp"]
 97
 98 #
          if file extension list[len(file extension list)-1] in allowed extension
99 #
              validity = True
100 #
          else:
101 #
              validity = False
102 #
              not file += file extension list[len(file extension list)-1]
103
104 #
          return ( validity , not file )
105
```

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