```
In [67]:
import seaborn as sns
import pandas as pd
import matplotlib.pyplot as plt
from imblearn.over sampling import RandomOverSampler
import numpy as np
from sklearn.model selection import train test split
import os, cv2
import tensorflow as tf
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Conv2D, Flatten, Dense, MaxPool2D
In [68]:
data = pd.read csv('/kaggle/input/skin-cancer-mnist-ham10000/hmnist 28 28 RGB.csv')
data.head()
Out[68]:
  pixel0000 pixel0001 pixel0002 pixel0003 pixel0004 pixel0005 pixel0006 pixel0007 pixel0008 pixel0009 ... pixel2343 pix
0
       192
                153
                        193
                                 195
                                          155
                                                  192
                                                           197
                                                                    154
                                                                            185
                                                                                     202 ...
                                                                                                 173
1
        25
                 14
                         30
                                  68
                                           48
                                                   75
                                                           123
                                                                     93
                                                                            126
                                                                                     158 ...
                                                                                                 60
2
       192
                138
                         153
                                 200
                                          145
                                                  163
                                                           201
                                                                    142
                                                                            160
                                                                                     206 ...
                                                                                                 167
3
        38
                         30
                                  95
                                           59
                                                                    103
                 19
                                                   72
                                                           143
                                                                            119
                                                                                     171 ...
                                                                                                 44
       158
                113
                         139
                                 194
                                          144
                                                                    162
                                                                            191
                                                                                                209
                                                  174
                                                           215
                                                                                     225 ...
5 rows × 2353 columns
In [69]:
data['label'].unique()
Out[69]:
array([2, 4, 3, 6, 5, 1, 0])
In [70]:
y = data['label']
x = data.drop(columns = ['label'])
In [71]:
data.isnull().sum().sum() #no null values present
Out[71]:
0
In [72]:
meta data = pd.read csv('/kaggle/input/skin-cancer-mnist-ham10000/HAM10000 metadata.csv')
meta_data.head()
Out[72]:
       lesion_id
                  image_id dx dx_type age
                                           sex localization
0 HAM_0000118 ISIC_0027419 bkl
                                 histo 80.0 male
                                                    scalp
```

1 HAM\_0000118 ISIC\_0025030 bkl

2 HAM\_0002730 ISIC\_0026769 bkl

2 HAM DODOTOD ISIC DOSESS NO

histo 80.0 male

histo 80.0 male

hista On A mala

scalp

scalp

```
lesion_id image_id dx dx_type age sex localization

-4 HAM_0001466 ISIC_0031633 bkl histo 75.0 male ear
```

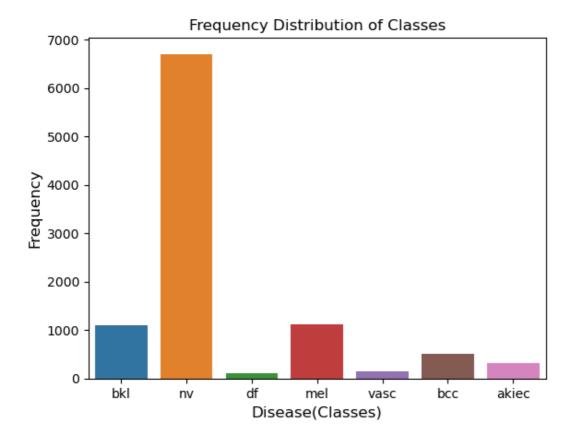
```
In [73]:
```

```
meta_data['dx'].unique()
Out[73]:
array(['bkl', 'nv', 'df', 'mel', 'vasc', 'bcc', 'akiec'], dtype=object)
In [74]:
```

```
sns.countplot(x = 'dx', data = meta_data)
plt.xlabel('Disease(Classes)', size=12)
plt.ylabel('Frequency', size=12)
plt.title('Frequency Distribution of Classes')
```

### Out[74]:

Text(0.5, 1.0, 'Frequency Distribution of Classes')



### In [75]:

```
print(x.shape, y.shape)
# To overcome class imbalace
oversample = RandomOverSampler()
x,y = oversample.fit_resample(x,y)
print(x.shape, y.shape)

(10015, 2352) (10015,)
(46935, 2352) (46935,)

In []:
sns.countplot(x = 'dx', data = meta_data)
plt.xlabel('Disease(Classes)', size=12)
```

# In [ ]:

plt.ylabel('Frequency', size=12)

plt.title('Frequency Distribution of Classes')

```
In [76]:
```

```
# reshaping the data so that it can be taken by convolution neural network(without distur
bing the no. of samples)
x = np.array(x).reshape(-1,28,28,3)
print('Shape of X :',x.shape)
print('Shape of y :',y.shape)
```

Shape of X: (46935, 28, 28, 3)Shape of y: (46935,)

### In [77]:

```
# Splitting Data
X_train, X_test, Y_train, Y_test = train_test_split(x,y, test_size=0.2, random_state=1)
print(X_train.shape,Y_train.shape)
print(X_test.shape , Y_test.shape)
```

(37548, 28, 28, 3) (37548,) (9387, 28, 28, 3) (9387,)

### In [78]:

```
model = Sequential()
model.add(Conv2D(16, kernel_size = (3,3), input_shape = (28, 28, 3), activation = 'relu'))
model.add(Conv2D(32, kernel_size = (3,3), activation = 'relu'))
model.add(MaxPool2D(pool_size = (2,2)))

model.add(Conv2D(32, kernel_size = (3,3), activation = 'relu'))
model.add(Conv2D(64, kernel_size = (3,3), activation = 'relu'))
model.add(MaxPool2D(pool_size = (2,2)))

model.add(Flatten())
model.add(Dense(64, activation='relu'))
model.add(Dense(7, activation='softmax'))
model.summary()
```

# Model: "sequential 1"

Layer (type)	Output Shape	Param #
conv2d_4 (Conv2D)	(None, 26, 26, 16)	448
conv2d_5 (Conv2D)	(None, 24, 24, 32)	4640
<pre>max_pooling2d_2 (MaxPooling 2D)</pre>	(None, 12, 12, 32)	0
conv2d_6 (Conv2D)	(None, 10, 10, 32)	9248
conv2d_7 (Conv2D)	(None, 8, 8, 64)	18496
<pre>max_pooling2d_3 (MaxPooling 2D)</pre>	(None, 4, 4, 64)	0
flatten_1 (Flatten)	(None, 1024)	0
dense_5 (Dense)	(None, 64)	65600
dense_6 (Dense)	(None, 7)	455

Total params: 98,887 Trainable params: 98,887 Non-trainable params: 0

## In [79]:

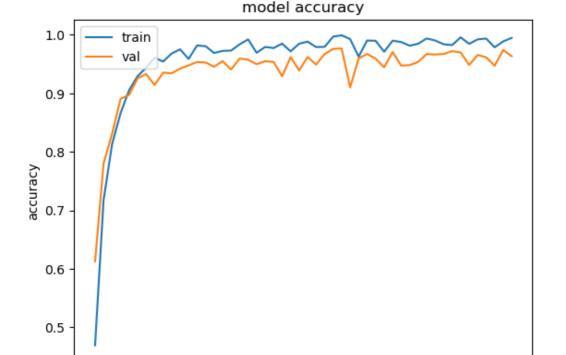
model.compile(loss = 'sparse categorical crossentropy',

```
75 - val loss: 0.6307 - val accuracy: 0.7802
Epoch 3/\overline{50}
29 - val loss: 0.4791 - val accuracy: 0.8308
Epoch 4/\overline{5}0
62 - val loss: 0.3350 - val accuracy: 0.8907
Epoch 5/50
54 - val loss: 0.3125 - val accuracy: 0.8973
Epoch 6/50
94 - val loss: 0.2361 - val accuracy: 0.9257
Epoch 7/50
37 - val loss: 0.2292 - val accuracy: 0.9330
Epoch 8/50
11 - val loss: 0.2561 - val accuracy: 0.9142
Epoch 9/50
43 - val loss: 0.2284 - val accuracy: 0.9354
Epoch 10/50
81 - val loss: 0.2244 - val accuracy: 0.9344
Epoch 11/50
53 - val_loss: 0.2408 - val_accuracy: 0.9422
Epoch 12/50
89 - val loss: 0.2025 - val accuracy: 0.9479
Epoch 13/50
235/235 [=============== ] - 25s 106ms/step - loss: 0.0566 - accuracy: 0.98
20 - val loss: 0.1957 - val accuracy: 0.9537
Epoch 14/50
06 - val loss: 0.2234 - val accuracy: 0.9527
Epoch 15/50
90 - val loss: 0.2037 - val accuracy: 0.9455
Epoch 16/50
25 - val loss: 0.1920 - val accuracy: 0.9551
Epoch 17/50
235/235 [================ ] - 25s 104ms/step - loss: 0.0778 - accuracy: 0.97
32 - val loss: 0.2300 - val accuracy: 0.9407
Epoch 18/50
38 - val loss: 0.1655 - val accuracy: 0.9595
Epoch 19/50
22 - val loss: 0.1860 - val accuracy: 0.9575
Epoch 20/50
94 - val loss: 0.2461 - val accuracy: 0.9499
Epoch 21/50
93 - val loss: 0.2351 - val accuracy: 0.9549
Epoch 22/50
```

```
75 - val loss: 0.2050 - val accuracy: 0.9538
Epoch 23/50
51 - val loss: 0.3018 - val accuracy: 0.9290
Epoch 24/50
18 - val loss: 0.2202 - val accuracy: 0.9621
Epoch 25/50
49 - val loss: 0.2810 - val accuracy: 0.9393
Epoch 26/50
84 - val loss: 0.2186 - val accuracy: 0.9621
Epoch 27/50
91 - val loss: 0.2246 - val accuracy: 0.9493
Epoch 28/50
95 - val loss: 0.2096 - val accuracy: 0.9672
Epoch 29/50
70 - val loss: 0.1647 - val accuracy: 0.9760
Epoch 30/50
93 - val loss: 0.1677 - val accuracy: 0.9767
Epoch 31/50
28 - val loss: 0.4777 - val accuracy: 0.9101
Epoch 32/50
28 - val loss: 0.2033 - val accuracy: 0.9598
Epoch 33/50
03 - val loss: 0.1893 - val accuracy: 0.9674
Epoch 34/50
98 - val loss: 0.2182 - val accuracy: 0.9594
Epoch 35/50
235/235 [=============== ] - 25s 106ms/step - loss: 0.0925 - accuracy: 0.97
12 - val_loss: 0.2705 - val accuracy: 0.9446
Epoch 36/50
01 - val loss: 0.1762 - val accuracy: 0.9706
Epoch 37/50
235/235 [=============== ] - 26s 109ms/step - loss: 0.0393 - accuracy: 0.98
78 - val loss: 0.2368 - val accuracy: 0.9475
Epoch 38/50
15 - val loss: 0.2727 - val accuracy: 0.9482
Epoch 39/50
48 - val loss: 0.2883 - val accuracy: 0.9538
Epoch 40/50
39 - val loss: 0.1856 - val accuracy: 0.9671
Epoch 41/50
04 - val loss: 0.1897 - val accuracy: 0.9664
Epoch 42/50
40 - val loss: 0.2031 - val accuracy: 0.9672
Epoch 43/50
25 - val loss: 0.1851 - val accuracy: 0.9722
Epoch 44/50
58 - val loss: 0.2126 - val accuracy: 0.9699
Epoch 45/50
46 - val loss: 0.2614 - val accuracy: 0.9486
```

Epoch  $46\overline{/}50$ 

```
plt.plot(history.history['accuracy'])
plt.plot(history.history['val_accuracy'])
plt.title('model accuracy')
plt.ylabel('accuracy')
plt.xlabel('epoch')
plt.legend(['train', 'val'], loc='upper left')
plt.show()
```



20

epoch

10

# In [83]:

0

```
plt.plot(history.history['loss'])
plt.plot(history.history['val_loss'])
plt.title('model loss')
plt.ylabel('loss')
plt.xlabel('epoch')
plt.legend(['train', 'val'], loc='upper right')
plt.show()
```

30

40

50

# 2.00 - train val

# In [96]:

```
results = model.evaluate(X_test , Y_test, verbose=0)

print(" Test Loss: {:.5f}".format(results[0]))
print("Test Accuracy: {:.2f}%".format(results[1] * 100))
```

Test Loss: 0.24558
Test Accuracy: 96.27%

### In [97]:

```
from sklearn.metrics import confusion_matrix , classification_report

y_true = list(Y_test)
y_pred = model.predict(X_test)
y_pred = list(map(lambda x: np.argmax(x), y_pred))
print('Y Actual Values:' , y_true[0:10])
print('Y Predicted Values:' , y_pred[0:10])
```

294/294 [=======] - 3s 9ms/step Y Actual Values : [5, 1, 4, 0, 5, 0, 2, 0, 3, 2] Y Predicted Values : [5, 1, 4, 0, 5, 0, 2, 0, 3, 2]

### In [127]:

```
classes = {2:'bkl', 4:'nv', 3:'df', 6:'mel', 5:'vasc', 1:'bcc', 0:'akiec'}

classes_labels=[]
for key in classes.keys():
    classes_labels.append(key)
print(classes_labels)
```

[2, 4, 3, 6, 5, 1, 0]

### In [136]:

```
classes = {4: ('nv', ' melanocytic nevi'),
    6: ('mel', 'melanoma'),
    2 :('bkl', 'benign keratosis-like lesions'),
    1:('bcc', ' basal cell carcinoma'),
    5: ('vasc', ' pyogenic granulomas and hemorrhage'),
    0: ('akiec', 'Actinic keratoses and intraepithelial carcinomae'),
    3: ('df', 'dermatofibroma')}
```

### In [100]:

Ω

0 1351

Ω

0

0

0]

```
cm = confusion_matrix(y_true,y_pred,labels=classes_labels)
print(confusion_matrix(y_true,y_pred,labels=classes_labels))

[[1235     9     0     15     2     0     1]
     [ 95 1079     2 161     7     25     5]
```

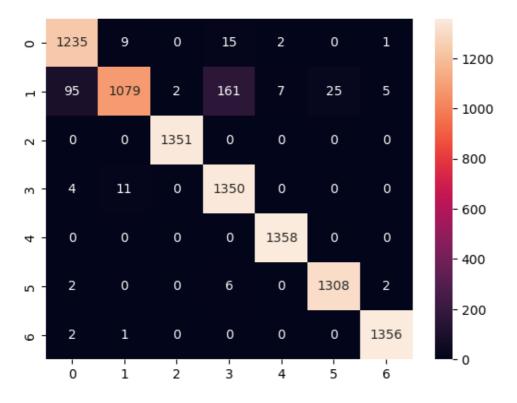
```
0 1350
[
     4
          11
                              0
                                     0
                                           0]
    0
[
           0
                 0
                        0 1358
                                    0
                                           0]
    2
[
           0
                        6
                              0 1308
     2
[
           1
                        0
                              0
                                     0 1356]]
```

# In [101]:

```
sns.heatmap(cm, annot = True, fmt='')
```

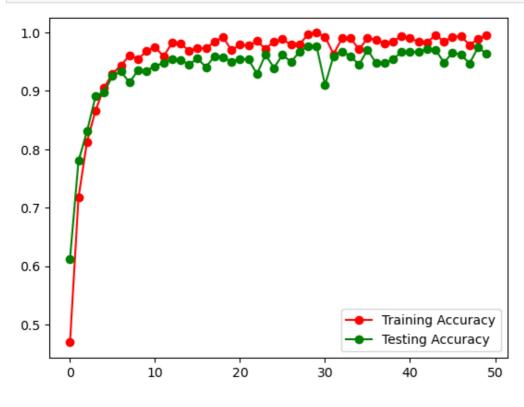
# Out[101]:

<AxesSubplot:>



# In [102]:

```
#training acc vs testing acc graph
plt.plot(history.history["accuracy"] , 'ro-' , label = "Training Accuracy")
plt.plot(history.history["val_accuracy"] , 'go-' , label = "Testing Accuracy")
plt.legend()
plt.show()
```



9387	9387	precision	recall	f1-
score	support	r		
('akie	ec', 'Actinic keratoses and intraepithelial carcinomae') 1359	0.99	1.00	0
0.99	('bcc', ' basal cell carcinoma') 1318	0.98	0.99	
0.95	('bkl', 'benign keratosis-like lesions')	0.92	0.98	
1.00	('df', 'dermatofibroma') 1351	1.00	1.00	
	('nv', ' melanocytic nevi')	0.98	0.79	
0.87	1374 ('vasc', ' pyogenic granulomas and hemorrhage')	0.99	1.00	
1.00	1358 ('mel', 'melanoma')	0.88	0.99	
0.93	1365			
0.96	9387			
0.96	9387 macro avg	0.96	0.96	
0.96	weighted avg	0.96	0.96	

# In [ ]: