```
In [ ]:
```

## **Dataset Overview**

HAM10000 ("Human Against Machine with 10000 training images") dataset - a large collection of multi-source dermatoscopic images of pigmented lesions

The dermatoscopic images are collected from different populations, acquired and stored by different modalities. The final dataset consists of 10015 dermatoscopic images.

It has 7 different classes of skin cancer which are listed below:

- Melanocytic nevi
- Melanoma
- Benign keratosis-like lesions
- · Basal cell carcinoma
- Actinic keratoses
- Vascular lesions
- Dermatofibroma

## **Importing libraries**

```
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, Activation
from sklearn.metrics import classification_report, accuracy_score
```

## **Reading the Data**

```
In []:
data = pd.read_csv('/kaggle/input/skin-cancer-mnist-ham10000/hmnist_28_28_RGB.csv')
data.head()
```

# **Data Preprocessing**

# **Data Cleaning**

In [ ]:

```
data['label'].unique()

In []:

y = data['label']
x = data.drop(columns = ['label'])
```

```
In [ ]:
data.isnull().sum().sum() #no null values present
In [ ]:
meta data = pd.read csv('/kaggle/input/skin-cancer-mnist-ham10000/HAM10000 metadata.csv')
meta data.head()
In [ ]:
meta data['dx'].unique()
In [ ]:
y = data['label']
x = data.drop(columns = ['label'])
In [ ]:
data.isnull().sum().sum() #no null values present
In [ ]:
meta data = pd.read csv('/kaggle/input/skin-cancer-mnist-ham10000/HAM10000 metadata.csv')
meta data.head()
In [ ]:
meta_data['dx'].unique()
Exploratory Data Analysis
In [ ]:
sns.countplot(x = 'dx', data = meta data)
plt.xlabel('Disease(Classes)', size=12)
plt.ylabel('Frequency', size=12)
plt.title('Frequency Distribution of Classes')
In [ ]:
sns.set style('whitegrid')
colors = ['#87ace8','#e3784d', 'green']
fig,axes = plt.subplots(figsize=(8,8))
ax = sns.countplot(x='sex',data=meta data, palette = 'Paired')
for container in ax.containers:
   ax.bar label(container)
plt.title('Gender-wise Distribution')
plt.xticks(rotation=45)
plt.show()
In [ ]:
sns.set style('whitegrid')
fig,axes = plt.subplots(figsize=(12,8))
ax = sns.countplot(x='dx',data=meta data, order = meta data['dx'].value counts().index,
palette = 'Paired')
for container in ax.containers:
   ax.bar label(container)
plt.title('Cell Types Skin Cancer Affected patients')
plt.xticks(rotation=45)
plt.show()
In [ ]:
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```
In [ ]:
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)
In [ ]:
sns.set style('whitegrid')
fig,axes = plt.subplots(figsize=(12,8))
ax = sns.countplot(x='dx', hue='sex', data=meta data, order = meta data['dx'].value count
s().index, palette = 'Paired')
for container in ax.containers:
    ax.bar_label(container)
plt.title('Cell Types Frequencies')
plt.xticks(rotation=45)
plt.show()
In [ ]:
sns.set style('whitegrid')
fig,axes = plt.subplots(figsize=(12,8))
ax = sns.countplot(x='localization', data=meta data, order = meta data['localization'].val
ue_counts().index, palette = 'crest')
for container in ax.containers:
    ax.bar label(container)
plt.title('Localization Area Frequencies')
plt.xticks(rotation=45)
plt.show()
In [ ]:
sns.set style('whitegrid')
fig,axes = plt.subplots(figsize=(12,8))
ax = sns.histplot(data=meta data, x='age')
plt.title('Age Histogram')
plt.show()
In [ ]:
sns.set style('whitegrid')
fig,axes = plt.subplots(figsize=(12,8))
ax = sns.histplot(data=meta data, x='age', hue='sex', multiple='stack')
plt.title('Age Histogram Gender Wise')
plt.show()
In [ ]:
print(x.shape, y.shape)
# To overcome class imbalace
oversample = RandomOverSampler()
x,y = oversample.fit resample(x,y)
print(x.shape, y.shape)
In [ ]:
# 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)
In [ ]:
# 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)
```

#### In [ ]:

```
model CNN = Sequential()
model CNN.add(Conv2D(16, kernel size = (3,3), input shape = (28, 28, 3), activation = 'r
elu', padding = 'same'))
model CNN.add(MaxPool2D(pool size = (2,2)))
model_CNN.add(Conv2D(32, kernel_size = (3,3), activation = 'relu', padding = 'same'))
model CNN.add(MaxPool2D(pool_size = (2,2), padding = 'same'))
model CNN.add(Conv2D(64, kernel size = (3,3), activation = 'relu', padding = 'same'))
model CNN.add(MaxPool2D(pool size = (2,2), padding = 'same'))
model CNN.add(Conv2D(128, kernel size = (3,3), activation = 'relu', padding = 'same'))
model CNN.add(MaxPool2D(pool size = (2,2), padding = 'same'))
model CNN.add(Flatten())
model CNN.add(Dense(64, activation = 'relu'))
model CNN.add(Dense(32))
model_CNN.add(Activation(activation='relu'))
model_CNN.add(Dense(7))
model CNN.add(Activation(activation='softmax'))
optimizer = tf.keras.optimizers.Adam(learning rate = 0.001)
model CNN.compile(loss = 'sparse categorical crossentropy',
                 optimizer = optimizer,
                 metrics = ['accuracy'])
print(model CNN.summary())
```

### In [ ]:

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

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

### In [ ]:

```
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()
```

#### In [ ]:

```
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()
In [ ]:
from sklearn.metrics import confusion matrix , classification report
y true CNN = list(Y test)
y pred CNN = model CNN.predict(X test)
y pred CNN = list(map(lambda x: np.argmax(x), y pred CNN))
print("Predicting First Ten Rows:")
print('Y Actual Values :' , y_true_CNN[0:10])
print('Y Predicted Values :' , y_pred_CNN[0:10])
In [ ]:
cm_CNN = confusion_matrix(y_true_CNN,y_pred_CNN,labels=classes_labels)
print(confusion matrix(y true CNN, y pred CNN, labels=classes labels))
sns.heatmap(cm CNN, annot = True, fmt='')
In [ ]:
#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()
In [ ]:
#predicting
y pred CNN = model CNN.predict(X test)
target names = [f"{classes[i]}" for i in range(7)]
y pred CNN = list(map(lambda x: np.argmax(x), y_pred_CNN))
print("CNN Model Prediction Results")
print(classification report(Y test , y pred CNN, target names=target names))
In [ ]:
# Layers definitions
from keras import backend as K
for l in range(len(model CNN.layers)):
   print(l, model CNN.layers[l])
In [ ]:
model CNN.layers[-2]
In [ ]:
os.environ["KERAS BACKEND"] = "tensorflow"
kerasBKED = os.environ["KERAS BACKEND"]
print(kerasBKED)
```

## **Separating Features Layers from the CNN Model**

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## **Extracting Features from CNN Model**

```
In [ ]:
# Extract features from input data using the CNN model
X train cnn = cnn model features.predict(X train)
X test cnn = cnn model features.predict(X test)
```

# Integrating CNN with SVM Classifier using Grid Search for Best

```
Perameters
In [ ]:
import numpy as np
from sklearn.svm import SVC
from sklearn.model_selection import GridSearchCV
parameters = {'kernel':['rbf'],
              'C':[1, 10, 100, 1000],
              'gamma':[1e-3, 1e-4]}
clf = GridSearchCV(SVC(), parameters)
clf.fit(X train cnn, Y train)
# Evaluate the combined CNN-SVM model on a test dataset
svm accuracy = clf.score(X test cnn, Y test)
print('SVM Accuracy:', svm accuracy*100)
y testSVM = clf.predict(X test cnn)
In [ ]:
svm accuracy = clf.score(X test cnn, Y test)
print('SVM Accuracy:', svm accuracy*100)
svmclf = clf.best estimator
print(svmclf)
svmclf.fit(X train cnn, Y train)
print("Accuracy: {0}".format(accuracy_score(Y_test, y_testSVM)*100))
In [ ]:
y testSVM = svmclf.predict(X test cnn)
from sklearn.metrics import confusion matrix, classification report, accuracy score
print(classification report(Y test, y testSVM, target names=target names))
print("Accuracy: {0}".format(accuracy_score(Y_test, y_testSVM)*100))
In [ ]:
```

## Integrating CNN with Random Forest Classifier using Grid Search for **Best Perameters**

```
In [ ]:
from sklearn.ensemble import RandomForestClassifier
from sklearn.model selection import GridSearchCV
parameters = {"max depth": [3, None],
              "max features": [1, 3, 10],
              "min_samples_split": [1.0, 3, 10],
              "min samples leaf": [1, 3, 10],
              "bootstrap": [True, False],
              "criterion": ["gini", "entropy"],
              "n estimators": [10, 20, 50]}
rclf = RandomForestClassifier()
rgclf = GridSearchCV(rclf, param grid=parameters)
rgclf.fit(X train cnn, Y train)
```

```
print('Random Forest Classifier Accuracy:', RFC_accuracy*100)
y_test_RF = rgclf.predict(X_test_cnn)
print("Accuracy: {0}".format(accuracy_score(Y_test, y_test_RF)*100))

In []:

y_test_RF = rgclf.predict(X_test_cnn)
print("Accuracy: {0}".format(accuracy_score(Y_test, y_test_RF)*100))
RFclf = rgclf.best_estimator_
RFclf.fit(X_test_cnn, Y_test)
print(RFclf)
y_testRFC = RFclf.predict(X_test_cnn)
from sklearn.metrics import confusion_matrix, classification_report, accuracy_score

print(classification_report(Y_test, y_testRFC, target_names=target_names))
print("Accuracy: {0}".format(accuracy_score(Y_test, y_testRFC)*100))
```

# Integrating CNN with KNN Classifier using Grid Search for Best Perameters

RFC\_accuracy = rgclf.score(X\_test\_cnn, Y\_test)

```
In []:

y_testKNN = kgclf.predict(X_test_cnn)
KNNclf = kgclf.best_estimator_
print(KNNclf)
from sklearn.metrics import confusion_matrix, classification_report, accuracy_score

print(classification_report(Y_test, y_testKNN, target_names=target_names))
print("Accuracy Score: {0}".format(accuracy_score(Y_test, y_testKNN)*100))
```

# Integrating CNN with Logistic Regression Classifier using Grid Search for Best Perameters

| <pre>grid_search_LR.fit(X_train_cnn, Y_train)</pre>  |
|--|
| <pre># Print the best hyperparameters and the corresponding accuracy score print("Best hyperparameters: ", grid_search_LR.best_params_) y_test_LR = grid_search_LR.predict(X_test_cnn)</pre> |
| <pre>print(classification_report(Y_test, y_test_LR, target_names=target_names)) print("Accuracy: {0}".format(accuracy_score(Y_test, y_test_LR)*100))</pre>                                   |
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