In [20]:	Importing the liabraries # Importing the liabraries
	<pre>import tensorflow as tf import pandas as pd import cv2 import warnings import matplotlib.pyplot as plt import numpy as np</pre>
	<pre>import matplotlib.pyplot as plt warnings.filterwarnings("ignore") from tqdm import tqdm from tensorflow.keras.models import Sequential, Model from tensorflow.keras.optimizers import Adam</pre>
	<pre>from tensorflow.keras.layers import Dense, Activation, Conv2D, MaxPooling2D, MaxPool2D, Flatten, Dropout, Batcl from tensorflow.keras.preprocessing.image import ImageDataGenerator from tensorflow.keras.callbacks import EarlyStopping, ModelCheckpoint from tensorflow.keras.utils import to_categorical from imblearn.over_sampling import SMOTE from tensorflow import keras from IPython.display import clear output</pre>
	<pre>from sklearn.model_selection import train_test_split from imblearn.under_sampling import RandomUnderSampler from sklearn.preprocessing import LabelEncoder from sklearn.metrics import confusion_matrix,ConfusionMatrixDisplay, accuracy_score,fl_score,classification_rep from tensorflow.keras.preprocessing.image import ImageDataGenerator from tensorflow.keras.callbacks import ReduceLROnPlateau</pre>
	Loading and Analysis of Data
In [21]: Out[21]:	df= pd.read_csv('skin_lesions.csv') df image_id
	1ISIC_0025661Benign keratosis-like lesions1.0scalp2ISIC_0027850Benign keratosis-like lesions1.0ear3ISIC_0025837Benign keratosis-like lesions1.0back4ISIC_0025209Benign keratosis-like lesions1.0back
	5367 ISIC_0026722 Melanoma 1.0 NaN 5368 ISIC_0029038 Melanocytic nevi 1.0 NaN 5369 ISIC_0032603 Benign keratosis-like lesions 1.0 NaN 5370 rows × 4 columns
In [3]:	<pre># check for the missing values in the csv df.isna().sum()</pre>
Out[3]: In [4]:	<pre>image_id 0 cell_type 100 is_benign 100 localization 13 dtype: int64 # drop the rows with missing values</pre>
In [4]:	<pre>df = df.dropna() # again check for missing values</pre>
Out[5]:	<pre>df.isna().sum() image_id</pre>
In [6]:	<pre>#check the class distribution print(df['cell_type'].value_counts()) df['cell_type'].value_counts().plot.bar() Melanocytic nevi 3528</pre>
	Benign keratosis-like lesions 580 Melanoma 576 Basal cell carcinoma 268 Actinic keratoses 169 Vascular lesions 74 Dermatofibroma 62
Out[6]:	Name: cell_type, dtype: int64 <axessubplot:> 3500 3000</axessubplot:>
	2500 - 2000 - 1500 -
	1000 - 500 - 100 ow a ow a sign was sign was sign with a sign was sign was sign with a sign was sign was sign with a sign was
	Melanocytic nevi Benign keratosis-like lesions Actinic keratoses Vascular lesions Dermatofibroma
	Observations • The dataset is imbalanced.
In [7]: In [8]:	
	<pre># now lets load the images all_data=[] labels=[] for i in tqdm (range(len(df))): path="/Users/priyank_7/Documents/Jupiter/Skinlesions/skin_lesions_images/"+df["image_id"][i]+".jpg" #print(path)</pre>
	<pre>img=cv2.imread(path) img=cv2.resize(img,(150,150)) img=cv2.cvtColor(img, cv2.COLOR_BGR2RGB) img=img*(1/255) all_data.append(img) labels.append(df['cell type'][i])</pre>
In [9]:	100% 5257/5257 [00:26<00:00, 201.77it/s]
In [10]:	<pre># (Synthetic Minority Over Sampling Technique) sm = SMOTE() all_data = all_data.reshape(all_data.shape[0],-1) all_data, labels = sm.fit_resample(all_data, labels)</pre>
In [11]:	<pre>all_data =all_data.reshape(all_data.shape[0],150,150,3) print("Max pixel value = ",all_data.max()) # one hot encoding for class label</pre>
	<pre>labels = to_categorical(labels) print("Shape of images = ",all_data.shape) print("Shape of labels = ",labels.shape) Max pixel value = 1.0 Shape of images = (24696, 150, 150, 3) Shape of labels = (24696, 7)</pre>
In [12]:	
	<pre>i=0 for img , ax in zip(images_arr, axes): ax.imshow(img) ax.axis('off') title="Class Label ="+str(labels[i]) ax.set_title(title)</pre>
	<pre>i+=1 plt.tight_layout() plt.show() plot_images(all_data[:10], labels[:10])</pre>
	Class Label =[0. 0. 1. 0. 0. 0. 0.] Class Label =[0. 0. 1. 0. 0. 0. 0.] Class Label =[0. 0. 1. 0. 0. 0. 0.]
	Class Label =[0. 0. 1. 0. 0. 0. 0.] Class Label =[0. 0. 1. 0. 0. 0. 0.] Class Label =[0. 0. 1. 0. 0. 0. 0.]
In [13]:	<pre>X_train, X_test, y_train, y_test = train_test_split(all_data, labels, test_size=0.3, random_state=42) del(all_data) del(labels)</pre>
	<pre>print("Shape of training images = ",X_train.shape) print("Shape of training labels = ",y_train.shape) print("Shape of validation images = ",X_test.shape) print("Shape of validation labels = ",y_test.shape) Shape of training images = (17287, 150, 150, 3) Shape of training labels = (17287, 7)</pre>
In [14]:	<pre>Shape of validation images = (7409, 150, 150, 3) Shape of validation labels = (7409, 7) # plot accuracy, loss curve (graph) class PlotLearning(keras.callbacks.Callback): def on_train_begin(self, logs={}): self.i = 0</pre>
	<pre>self.r = 0 self.x = [] self.losses = [] self.val_losses = [] self.acc = [] self.val_acc = [] self.fig = plt.figure()</pre>
	<pre>self.logs = [] def on_epoch_end(self, epoch, logs={}): self.logs.append(logs)</pre>
	<pre>self.x.append(self.i) self.losses.append(logs.get('loss')) self.val_losses.append(logs.get('val_loss')) self.acc.append(logs.get('accuracy')) self.val_acc.append(logs.get('val_accuracy')) self.i += 1 for (ov1 = ov2) = plt supplets(1 = 2 = abarov=###we)</pre>
	<pre>f, (ax1, ax2) = plt.subplots(1, 2, sharex=True) clear_output(wait=True) ax1.set_yscale('log') ax1.plot(self.x, self.losses, label="loss") ax1.plot(self.x, self.val losses, label="val loss")</pre>
	<pre>ax1.legend() ax2.plot(self.x, self.acc, label="accuracy") ax2.plot(self.x, self.val_acc, label="validation accuracy") ax2.legend()</pre>
	<pre>plt.show(); accuracy_loss_plot = PlotLearning() Model</pre>
In [15]:	<pre># initializing the model input_shape = (150, 150, 3) num_classes = 7</pre>
	<pre># creating sequential model model = Sequential() # adding extra conv2D, pooling layer(s) model.add(Conv2D(32, kernel_size=(3, 3),activation='relu',padding = 'Same',input_shape=input_shape))</pre>
	<pre>model.add(Conv2D(32,kernel_size=(3, 3),activation='relu',padding = 'Same')) model.add(MaxPool2D(pool_size = (2, 2))) model.add(Dropout(0.25)) model.add(Conv2D(64, (3, 3), activation='relu',padding ='Same')) model.add(Conv2D(64, (3, 3), activation='relu',padding ='Same')) model.add(MaxPool2D(pool_size = (2, 2)))</pre>
	<pre>model.add(MaxFoo12D(poo1_s12e = (2, 2))) model.add(Dropout(0.40)) model.add(Flatten()) model.add(Dense(128, activation='relu')) model.add(Dropout(0.5)) model.add(Dense(num classes, activation='softmax'))</pre>
	<pre>model.add(Dense(num_classes, activation='softmax')) model.summary() Metal device set to: Apple M1 Pro Model: "sequential" Layer (type)</pre>
	conv2d (Conv2D) (None, 150, 150, 32) 896 conv2d_1 (Conv2D) (None, 150, 150, 32) 9248 max_pooling2d (MaxPooling2D (None, 75, 75, 32) 0)
	dropout (Dropout) (None, 75, 75, 32) 0 conv2d_2 (Conv2D) (None, 75, 75, 64) 18496 conv2d_3 (Conv2D) (None, 75, 75, 64) 36928
	max_pooling2d_1 (MaxPooling (None, 37, 37, 64) 0 2D) dropout_1 (Dropout) (None, 37, 37, 64) 0 flatten (Flatten) (None, 87616) 0
	dense (Dense) (None, 128) 11214976 dropout_2 (Dropout) (None, 128) 0 dense 1 (Dense) (None, 7) 903
	Total params: 11,281,447 Trainable params: 11,281,447 Non-trainable params: 0
	2022-04-24 19:31:07.529836: I tensorflow/core/common_runtime/pluggable_device/pluggable_device_factory.cc:305] Could not identify NUMA node of platform GPU ID 0, defaulting to 0. Your kernel may not have been built with NU MA support. 2022-04-24 19:31:07.530228: I tensorflow/core/common_runtime/pluggable_device/pluggable_device_factory.cc:271] Created TensorFlow device (/job:localhost/replica:0/task:0/device:GPU:0 with 0 MB memory) -> physical Pluggable Device (device: 0, name: METAL, pci bus id: <undefined>)</undefined>
	<pre>model.compile(loss='categorical_crossentropy',optimizer='adam',metrics=["accuracy"]) Define callbacks # Set a learning mate appealer</pre>
In [17]:	<pre>learning_rate_reduction = ReduceLROnPlateau(monitor='val_loss',</pre>
In [18]:	<pre>callbacks = [ModelCheckpoint('.mdl_wts.hdf5', monitor='val_loss', save_best_only=True),accuracy_loss_plot,learn Training the model datagen = ImageDataGenerator(</pre>
111 [10].	rotation_range=10, # randomly rotate images in the range (degrees, 0 to 180) zoom_range = 0.1 # Randomly zoom image # Fit the model epochs = 15
	<pre>batch_size = 128 history = model.fit(datagen.flow(X_train,y_train, batch_size=batch_size),</pre>
	10°] 0.9 - 0.8 - 0.7 -
	0.6 - 0.5 - 0.5 - 0.4
	3×10 ⁻¹ 0.3 validation accuracy 135/135 [====================================
In [19]:	Performance Evaluation from sklearn.metrics import classification_report # Get predictions on test data y_predict=model.predict(X_test)
	<pre>y_predict = np.argmax(y_predict, axis=-1) y_true= np.argmax(y_test, axis=-1) # plot confusion matrix cm=confusion_matrix(y_true, y_predict) plt.figure(figsize=(15,10))</pre>
	<pre>ConfusionMatrixDisplay(cm).plot() # check performance print("Accuracy on test data",accuracy_score(y_true,y_predict)) print(classification_report(y_true,y_predict))</pre>
	2022-04-24 19:54:32.347804: I tensorflow/core/grappler/optimizers/custom_graph_optimizer_registry.cc:113] Plugi n optimizer for device_type GPU is enabled. Accuracy on test data 0.8782561749223917 precision recall f1-score support 0 0.93 0.96 0.94 1044 1 0.92 0.95 0.93 1060
	1 0.92 0.95 0.93 1060 2 0.79 0.82 0.80 1067 3 0.98 0.99 0.99 1040 4 0.92 0.57 0.70 1071 5 0.69 0.87 0.77 1077 6 0.98 1.00 0.99 1050
	accuracy 0.88 7409 macro avg 0.89 0.88 0.88 7409 weighted avg 0.89 0.88 0.88 7409
	0 - 1001 29 7 2 0 5 0
	4 - 16
	0 1 2 3 4 5 6 Predicted label