```
Start coding or generate with AI.
import pandas as pd
import -----
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                 tion="ignore")
warning.......
import matplotlib.pyplot as plt
%matplotlib inline
from keras import backend as K
from sklearn.model_selection import train_test_split
from sklearn.metrics import classification_report
import tensorflow as tf
from tensorflow import keras
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Conv2D, MaxPooling2D, Flatten, Dense, Flatten
from tensorflow.keras.optimizers import Adam
from tensorflow.keras.losses import SparseCategoricalCrossentropy
from tensorflow.keras.preprocessing.image import ImageDataGenerator
import sklearn.metrics as metrics
from keras.callbacks import LearningRateScheduler
annealer = LearningRateScheduler(lambda x: 1e-3 * 0.95 ** x, verbose=0)
import os
import pandas as pd
healthy_dirs = [
    r'/kaggle/input/chest-xray-pneumonia/chest_xray/test/NORMAL',
    r'/kaggle/input/chest-xray-pneumonia/chest_xray/train/NORMAL',
    r'/kaggle/input/chest-xray-pneumonia/chest_xray/val/NORMAL',
]
pneumonia_dir = [
    r'/kaggle/input/chest-xray-pneumonia/chest_xray/test/PNEUMONIA',
    r'/kaggle/input/chest-xray-pneumonia/chest_xray/train/PNEUMONIA',
    r'/kaggle/input/chest-xray-pneumonia/chest_xray/val/PNEUMONIA'
filepaths = []
labels = []
dict_lists = [healthy_dirs, pneumonia_dir]
class_labels = ['Normal', 'Pneumonia']
for i, dir_list in enumerate(dict_lists):
    for j in dir_list:
        flist = os.listdir(j)
        for f in flist:
            fpath = os.path.join(j, f)
            filepaths.append(fpath)
            labels.append(class_labels[i])
Fseries = pd.Series(filepaths, name="filepaths")
Lseries = pd.Series(labels, name="labels")
pneumonia_data = pd.concat([Fseries, Lseries], axis=1)
pneumonia_df = pd.DataFrame(pneumonia_data)
print(pneumonia_df.head())
print(pneumonia_df["labels"].value_counts())
pneumonia_df.shape
train_images, test_images = train_test_split(pneumonia_df, test_size=0.3, random_state=42)
train_set, val_set = train_test_split(pneumonia_df, test_size=0.2, random_state=42)
print(train_set.shape)
print(test_images.shape)
print(val_set.shape)
print(train_images.shape)
```

```
image_gen = ImageDataGenerator(preprocessing_function= tf.keras.applications.mobilenet_v2.preprocess_input)
train = image_gen.flow_from_dataframe(dataframe= train_set,x_col="filepaths",y_col="labels",
                                    target_size=(244,244),
                                    color_mode='rgb',
                                    class_mode="categorical", #used for Sequential Model
                                    batch_size=4,
       Insert code cell below
                                    shuffle=False
                                                            #do not shuffle data
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test
                         _dataframe(dataframe= test_images,x_col="filepaths", y_col="labels",
                                   target_size=(244,244),
                                   color_mode='rgb',
                                   class_mode="categorical",
                                   batch_size=4,
                                   shuffle= False
target_size=(244,244),
                                  color_mode= 'rgb',
                                  class_mode="categorical",
                                  batch_size=4,
                                  shuffle=False
classes=list(train.class_indices.keys())
print (classes)
  Generate
                                                                                                                        Q
                                                                                                                              Close
               10 random numbers using numpy
def show_knee_images(image_gen):
   test_dict = test.class_indices
    classes = list(test_dict.keys())
    images, labels=next(image_gen) # get a sample batch from the generator
   plt.figure(figsize=(20,20))
    length = len(labels)
    if length<25:
       r=length
    else:
       r=25
    for i in range(r):
       plt.subplot(5,5,i+1)
       image=(images[i]+1)/2 #scale images between 0 and 1
       plt.imshow(image)
       index=np.argmax(labels[i])
       class_name=classes[index]
       plt.title(class_name, color="green",fontsize=16)
       plt.axis('off')
    plt.show()
    show_knee_images(train)
```

```
model = keras.models.Sequential([
    keras.layers.Conv2D(filters=128, kernel_size=(8, 8), strides=(3, 3), activation='relu', input_shape=(224, 224, 3)),
    keras.layers.BatchNormalization(),
    keras.layers.Conv2D(filters=256, kernel_size=(5, 5), strides=(1, 1), activation='relu', padding="same"),
    keras lavers RatchNormalization(),
    kei Insert code cell below (pool_size=(3, 3)),
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    kei
                         _lters=256, kernel_size=(3, 3), strides=(1, 1), activation='relu', padding="same"),
    keras.layers.BatchNormalization(),
    keras.layers.Conv2D(filters=256, kernel_size=(1, 1), strides=(1, 1), activation='relu', padding="same"),
    keras.layers.BatchNormalization(),
    keras.layers.Conv2D(filters=256, kernel_size=(1, 1), strides=(1, 1), activation='relu', padding="same"),
    keras.layers.BatchNormalization(),
    keras.layers.Conv2D(filters=512, kernel_size=(3, 3), activation='relu', padding="same"),
    keras.layers.BatchNormalization(),
    keras.layers.MaxPool2D(pool_size=(2, 2)),
    keras.layers.Conv2D(filters=512, kernel_size=(3, 3), activation='relu', padding="same"),
    keras.layers.BatchNormalization(),
    keras.layers.Conv2D(filters=512, kernel_size=(3, 3), activation='relu', padding="same"),
    keras.layers.BatchNormalization(),
    keras.layers.MaxPool2D(pool_size=(2, 2)),
    keras.layers.Conv2D(filters=512, kernel_size=(3, 3), activation='relu', padding="same"),
    keras.layers.BatchNormalization(),
    keras.layers.MaxPool2D(pool_size=(2, 2)),
    keras.layers.Flatten(),
    keras.layers.Dense(1024, activation='relu'),
    keras.layers.Dropout(0.5),
    keras.layers.Dense(1024, activation='relu'),
    keras.layers.Dropout(0.5),
    keras.layers.Dense(2, activation='softmax')
])
model.compile(
    loss='categorical_crossentropy',
    optimizer=tf.optimizers.SGD(learning_rate=0.001),
    metrics=['accuracy']
)
model.summary()
from keras.utils import plot_model
plot_model(model, to_file='model_plot.png', show_shapes=True, show_layer_names=True)
history = model.fit(train, epochs=10, validation_data=val, verbose=1)
model.evaluate(test, verbose=1)
model.save("pneumonia_Model.h5")
pred = model.predict(test)
pred = np.argmax(pred, axis=1) #pick class with highest probability
labels = (train.class_indices)
labels = dict((v,k) \text{ for } k,v \text{ in labels.items()})
pred2 = [labels[k] for k in pred]
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()
```

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plt.plot(history.history['loss'])
plt.plot(history.history['val_loss'])
plt.title('model loss')
plt.ylabel('loss')
plt.xl; denote |
plt.le Insert code cell below ], loc='upper left')
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from sklearn.metrics import confusion_matrix, accuracy_score
y_test = test_images.labels
print(classification_report(y_test, pred2))
print("Accuracy of the Model:","{:.1f}%".format(accuracy_score(y_test, pred2)*100))
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.metrics import confusion_matrix
class_labels = ['Normal', 'Pneumonia']
cm = confusion_matrix(y_test, pred2)
plt.figure(figsize=(10, 5))
sns.heatmap(cm, annot=True, fmt='g', vmin=0, cmap='Blues')
plt.xticks(ticks=[0.5, 1.5], labels=class_labels)
plt.yticks(ticks=[0.5, 1.5], labels=class_labels)
plt.xlabel("Predicted")
plt.ylabel("Actual")
plt.title("Confusion Matrix")
plt.show()
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