Machine Learning Case Study and Project Predicting Fractures and Brain Tumors Using CNN

SY Computer Engineering - Batch C2

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Problem Statement:

<u>Accurate diagnosis of medical conditions</u> such as fractures and brain tumors is critical for effective treatment and patient care. However, the manual interpretation of medical images by healthcare professionals can be time-consuming and prone to errors. Therefore, there is a need to develop <u>automated systems that can analyze medical images</u> and assist in the <u>early</u> detection of fractures and brain tumors.

Introduction:

Medical imaging techniques, such as X-rays and MRI scans, provide valuable insights into the internal structures of the human body. Interpretation of these images traditionally relies on the expertise of radiologists and physicians. However, advancements in machine learning have opened up new possibilities for automating the analysis of medical images.

In this case study, we aim to leverage convolutional neural networks (CNNs), a powerful class of deep learning models, to develop predictive models for detecting fractures and brain tumors from medical images. By training CNN models on labeled datasets containing X-ray and MRI images, we seek to build robust classifiers capable of accurately identifying pathological conditions.

Tech Stack Used:

Platform : Google Colab, Streamlit (possible UI)

Language: Python

Libraries:

- 1. Kaggle API datasets
- 2. Zipfile extraction of dataset files
- 3. Os file operations and directory management

- 4. Numpy Mathematical and Array operations
- 5. Matplotlib.pyplot plotting and visual representation
- 6. Matplotlib.image reading and displaying images
- 7. Cv2 (OpenCV) Computer Vision tasks
- 8. PIL Python Image Library imaging
- 9. Scikit Machine Learning model
- 10. Tensorflow and Keras Training ML models and Building Neural Networks
- 11. Streamlit Possible Web Page Creation for User Interface

Dataset Information:

Source : Kaggle Type : Images

Using a key instead of downloading entire datasets of images.

For X-ray Analysis and Fracture Detection:

```
!pip install kaggle

# configuring the path of Kaggle.json file
!mkdir -p ~/.kaggle
!cp kaggle.json ~/.kaggle/
!chmod 600 ~/.kaggle/kaggle.json

#importing dataset
!kaggle datasets download -d
vuppalaadithyasairam/bone-fracture-detection-using-xrays
!kaggle datasets download -d bebofekry/bone-fracture-atlas-ds-compressed
```

For MRI Analysis and Brain Tumor Detection:

```
!pip install kaggle
# configuring the path of Kaggle.json file
!mkdir -p ~/.kaggle
!cp kaggle.json ~/.kaggle/
!chmod 600 ~/.kaggle/kaggle.json
```

```
!kaggle datasets download -d shreyag1103/brain-mri-scans-for-brain-tumor-classification
```

Code and Output:

For X-ray Analysis and Fracture Detection: (Google Colab PDF is attached externally.) Image Processing:

```
data=[]
fractured path1='/content/archive (6)/train/fractured/'
fractured path2='/content/FracAtlas/FracAtlas/images/Train/Fractured/'
for img file in fractured files1:
  image = Image.open(fractured path1 + img file)
  image = image.resize((128, 128))
 image = image.convert('RGB')
  image = np.array(image)
 data.append(image)
for img file in fractured files2:
  image = Image.open(fractured path2 + img file)
 image = image.resize((128,128))
  image = image.convert('RGB')
 image = np.array(image)
 data.append(image)
not fractured path1='/content/archive (6)/train/not fractured/'
not fractured path2='/content/FracAtlas/FracAtlas/images/Train/Non fractur
ed/'
for img file in not fractured files1:
 image = Image.open(not fractured path1 + img file)
  image = image.resize((128, 128))
 image = image.convert('RGB')
  image = np.array(image)
```

```
data.append(image)

for img_file in not_fractured_files2:
   image = Image.open(not_fractured_path2 + img_file)
   image = image.resize((128,128))
   image = image.convert('RGB')
   image = np.array(image)
   data.append(image)
```

CNN:

```
import tensorflow as tf
from tensorflow import keras
num of classes = 2
model = keras.Sequential()
model.add(keras.layers.Conv2D(32, kernel size=(3,3), activation='relu',
input shape=(128,128,3)))
model.add(keras.layers.MaxPooling2D(pool size=(2,2)))
model.add(keras.layers.Conv2D(64, kernel size=(3,3), activation='relu'))
model.add(keras.layers.MaxPooling2D(pool size=(2,2)))
model.add(keras.layers.Flatten())
model.add(keras.layers.Dense(128, activation='relu'))
model.add(keras.layers.Dropout(0.5))
model.add(keras.layers.Dense(64, activation='relu'))
model.add(keras.layers.Dropout(0.5))
model.add(keras.layers.Dense(num of classes, activation='sigmoid'))
```

```
# compile the neural network
model.compile(optimizer='adam',
              metrics=['acc'])
history = model.fit(X train scaled, Y train, validation split=0.1,
epochs=5)
# Input from User and Prediction
input image path = input('Path of the image to be predicted: ')
input image = cv2.imread(input image path)
cv2 imshow(input image)
input image resized = cv2.resize(input image, (128,128))
input image scaled = input image resized/255
input_image_reshaped = np.reshape(input image scaled, [1,128,128,3])
input prediction = model.predict(input image reshaped)
print(input prediction)
input pred label = np.argmax(input prediction)
print(input pred label)
if input pred label == 1:
 print('The bone is fractured')
```

```
else:
print('The bone is not fractured')
```

For MRI Analysis and Brain Tumor Detection: (Google Colab PDF is attached externally.)

CNN:

```
X train, X test, Y train, Y test = train test split(X, Y, test size=0.2,
random_state=2)
X train scaled = X train/255
X \text{ test scaled} = X \text{ test/255}
import tensorflow as tf
from tensorflow import keras
num of classes = 4
model = keras.Sequential()
model.add(keras.layers.Conv2D(32, kernel_size=(3,3), activation='relu',
input shape=(128,128,3)))
model.add(keras.layers.MaxPooling2D(pool size=(2,2)))
model.add(keras.layers.Conv2D(64, kernel size=(3,3), activation='relu'))
model.add(keras.layers.MaxPooling2D(pool size=(2,2)))
model.add(keras.layers.Flatten())
model.add(keras.layers.Dense(128, activation='relu'))
model.add(keras.layers.Dropout(0.5))
model.add(keras.layers.Dense(64, activation='relu'))
```

```
model.add(keras.layers.Dropout(0.5))
model.add(keras.layers.Dense(num of classes, activation='sigmoid'))
model.compile(optimizer='adam',
              metrics=['acc'])
# training the neural network
history = model.fit(X train scaled, Y train, validation split=0.1,
epochs=50)
input image path = input('Path of the image to be predicted: ')
input image = cv2.imread(input image path)
cv2 imshow(input image)
input image resized = cv2.resize(input image, (128,128))
input image scaled = input image resized/255
input_image_reshaped = np.reshape(input image scaled, [1,128,128,3])
input prediction = model.predict(input image reshaped)
print(input prediction)
input pred label = np.argmax(input prediction)
print(input pred label)
if input pred label == 1:
 print('The person has tumor')
```

```
elif input_pred_label == 2:
    print('The person has tumor')

elif input_pred_label == 3:
    print('The person has tumor')

elif input_pred_label == 4:
    print('The person does not have tumor')

else:
    print('The person does not have tumor.')
```

Conclusion:

The development of CNN-based models for predicting fractures and brain tumors from medical images represents a significant advancement in healthcare technology. These models have the potential to assist healthcare professionals in making accurate diagnoses and providing timely treatment to patients.

By leveraging machine learning techniques, we can improve the efficiency and accuracy of medical image analysis, leading to better patient outcomes. However, further research and validation are necessary to ensure the reliability and generalization of these models across different populations and imaging modalities.

In conclusion, CNN-based models offer promising opportunities for enhancing diagnostic capabilities in medical imaging and advancing the field of radiology. With continued research and innovation, we can harness the power of machine learning to improve healthcare delivery and patient care.