Detecting Illneses Using XRays

Classification of Chest X Rays images into one of three classes: Covid, Normal, Pneumonia

Notebook created for Advanced Data Science Capstone Project on Coursera

Dataset from <u>COVID19+PNEUMONIA+NORMAL Chest X-Ray Images</u> (https://www.kaggle.com/sachinkumar413/covid-pneumonia-normal-chest-xray-images) on Kaggle

Importing Libraries

Using PyTorch version 1.10.1

Importing Data

```
In [2]: #downloading data from Kaggle
import opendatasets as od
od.download("https://www.kaggle.com/sachinkumar413/covid-pneumonia-normal-che

Please provide your Kaggle credentials to download this dataset. Learn mor
e: http://bit.ly/kaggle-creds (http://bit.ly/kaggle-creds)
Your Kaggle username: ijachiijachi
Your Kaggle Key: ......
Downloading covid-pneumonia-normal-chest-xray-images.zip to .\covid-pneumon
ia-normal-chest-xray-images

100%|
| 277M/277M [03:34<00:00, 1.36MB/s]</pre>
```

```
In [ ]: ► C:\Users\Hello\ADVANCED DATASCIENCE IBM CAPSTONE\covid-pneumonia-normal-chest
```

Preparing Training and Test Sets

```
In [3]:

▶ | class_names = ['covid', 'normal', 'pneumonia']
            root_dir = 'covid-pneumonia-normal-chest-xray-images'
            source_dirs = ['COVID', 'NORMAL', 'PNEUMONIA']
            if os.path.isdir(os.path.join(root_dir, source_dirs[1])):
                os.mkdir(os.path.join(root_dir, 'test'))
                for i, d in enumerate(source_dirs):
                    os.rename(os.path.join(root_dir, d), os.path.join(root_dir, class_nam
                for c in class_names:
                    os.mkdir(os.path.join(root_dir, 'test', c))
                for c in class_names:
                    images = [x for x in os.listdir(os.path.join(root_dir, c)) if x.lower
                    selected images = random.sample(images, 30)
                    for image in selected_images:
                        source_path = os.path.join(root_dir, c, image)
                        target_path = os.path.join(root_dir, 'test', c, image)
                        shutil.move(source_path, target_path)
```

Creating Custom Dataset

```
In [4]:

▶ class ChestXRayDataset(torch.utils.data.Dataset):
                def __init__(self, image_dirs, transform):
                     def get_images(class_name):
                         images = [x \text{ for } x \text{ in os.listdir(image dirs[class name]) if } x[-3:]
                         print(f'Found {len(images)} {class_name} examples')
                         return images
                     self.images = {}
                     self.class_names = ['covid', 'normal', 'pneumonia']
                     for class_name in self.class_names:
                         self.images[class_name] = get_images(class_name)
                     self.image dirs = image dirs
                     self.transform = transform
                def __len__(self):
                     return sum([len(self.images[class_name]) for class_name in self.class
                def __getitem__(self, index):
                     class name = random.choice(self.class names)
                     index = index % len(self.images[class_name])
                     image_name = self.images[class_name][index]
                     image_path = os.path.join(self.image_dirs[class_name], image_name)
                     image = Image.open(image path).convert('RGB')
                     return self.transform(image), self.class_names.index(class_name)
```

Image Transformations

Prepare DataLoader

```
In [7]:
          covid': 'C:/Users/Hello/ADVANCED DATASCIENCE IBM CAPSTONE/covid-pneumoni'
                 'normal': 'C:/Users/Hello/ADVANCED DATASCIENCE IBM CAPSTONE/covid-pneumon
                 'pneumonia': 'C:/Users/Hello/ADVANCED DATASCIENCE IBM CAPSTONE/covid-pneu
             }
             train_dataset = ChestXRayDataset(train_dirs, train_transform)
             Found 1596 covid examples
             Found 1772 normal examples
             Found 1770 pneumonia examples
 In [9]:
          test_dirs = {
                 'covid': 'C:/Users/Hello/ADVANCED DATASCIENCE IBM CAPSTONE/covid-pneumoni
                 'normal': 'C:/Users/Hello/ADVANCED DATASCIENCE IBM CAPSTONE/covid-pneumon
                 'pneumonia': 'C:/Users/Hello/ADVANCED DATASCIENCE IBM CAPSTONE/covid-pneu
             }
             test_dataset = ChestXRayDataset(test_dirs, test_transform)
             Found 30 covid examples
             Found 30 normal examples
             Found 30 pneumonia examples
In [10]:
         ▶ batch size = 6
             dl train = torch.utils.data.DataLoader(train dataset, batch size=batch size,
             dl_test = torch.utils.data.DataLoader(test_dataset, batch_size=batch_size, sf
             print('Number of training batches', len(dl train))
             print('Number of test batches', len(dl_test))
```

Number of training batches 857 Number of test batches 15

Data Visualization

```
In [11]:
             class_names = train_dataset.class_names
             def show images(images, labels, preds):
                 plt.figure(figsize=(8, 4))
                 for i, image in enumerate(images):
                     plt.subplot(1, 6, i + 1, xticks=[], yticks=[])
                     image = image.numpy().transpose((1, 2, 0))
                     mean = np.array([0.485, 0.456, 0.406])
                     std = np.array([0.229, 0.224, 0.225])
                     image = image * std + mean
                     image = np.clip(image, 0., 1.)
                     plt.imshow(image)
                     col = 'green'
                     if preds[i] != labels[i]:
                         col = 'red'
                     plt.xlabel(f'{class_names[int(labels[i].numpy())]}')
                     plt.ylabel(f'{class_names[int(preds[i].numpy())]}', color=col)
                 plt.tight_layout()
                 plt.show()
             images, labels = next(iter(dl_train))
In [12]:
             show_images(images, labels, labels)
                                          pneumonia
                                                                   pneumonia
In [13]:
             images, labels = next(iter(dl test))
             show_images(images, labels, labels)
```

Creating the Model

covid

pneumonia

covid

```
    | resnet18 = torchvision.models.resnet18(pretrained=True)

In [14]:
             print(resnet18)
             ResNet(
               (conv1): Conv2d(3, 64, kernel_size=(7, 7), stride=(2, 2), padding=(3, 3),
             bias=False)
               (bn1): BatchNorm2d(64, eps=1e-05, momentum=0.1, affine=True, track_runnin
             g stats=True)
               (relu): ReLU(inplace=True)
               (maxpool): MaxPool2d(kernel_size=3, stride=2, padding=1, dilation=1, ceil
             mode=False)
               (layer1): Sequential(
                 (0): BasicBlock(
                    (conv1): Conv2d(64, 64, kernel_size=(3, 3), stride=(1, 1), padding=
             (1, 1), bias=False)
                   (bn1): BatchNorm2d(64, eps=1e-05, momentum=0.1, affine=True, track_ru
             nning_stats=True)
                    (relu): ReLU(inplace=True)
                    (conv2): Conv2d(64, 64, kernel size=(3, 3), stride=(1, 1), padding=
             (1, 1), bias=False)
                   (bn2): BatchNorm2d(64, eps=1e-05, momentum=0.1, affine=True, track ru
             nning_stats=True)
                 (1): BasicBlock(
                   (conv1): Conv2d(64, 64, kernel_size=(3, 3), stride=(1, 1), padding=
             (1, 1), bias=False)
                   (bn1): BatchNorm2d(64, eps=1e-05, momentum=0.1, affine=True, track ru
             nning stats=True)
                   (relu): ReLU(inplace=True)
                   (conv2): Conv2d(64, 64, kernel size=(3, 3), stride=(1, 1), padding=
             (1, 1), bias=False)
                   (bn2): BatchNorm2d(64, eps=1e-05, momentum=0.1, affine=True, track ru
             nning_stats=True)
               (layer2): Sequential(
                 (0): BasicBlock(
                    (conv1): Conv2d(64, 128, kernel size=(3, 3), stride=(2, 2), padding=
             (1, 1), bias=False)
                   (bn1): BatchNorm2d(128, eps=1e-05, momentum=0.1, affine=True, track r
             unning_stats=True)
                   (relu): ReLU(inplace=True)
                    (conv2): Conv2d(128, 128, kernel_size=(3, 3), stride=(1, 1), padding=
             (1, 1), bias=False)
                   (bn2): BatchNorm2d(128, eps=1e-05, momentum=0.1, affine=True, track_r
             unning stats=True)
                   (downsample): Sequential(
                      (0): Conv2d(64, 128, kernel_size=(1, 1), stride=(2, 2), bias=False)
                     (1): BatchNorm2d(128, eps=1e-05, momentum=0.1, affine=True, track r
             unning_stats=True)
                   )
                 (1): BasicBlock(
                   (conv1): Conv2d(128, 128, kernel_size=(3, 3), stride=(1, 1), padding=
             (1, 1), bias=False)
```

IBM.

```
(bn1): BatchNorm2d(128, eps=1e-05, momentum=0.1, affine=True, track r
unning_stats=True)
      (relu): ReLU(inplace=True)
      (conv2): Conv2d(128, 128, kernel_size=(3, 3), stride=(1, 1), padding=
(1, 1), bias=False)
      (bn2): BatchNorm2d(128, eps=1e-05, momentum=0.1, affine=True, track_r
unning stats=True)
  )
  (layer3): Sequential(
    (0): BasicBlock(
      (conv1): Conv2d(128, 256, kernel_size=(3, 3), stride=(2, 2), padding=
(1, 1), bias=False)
      (bn1): BatchNorm2d(256, eps=1e-05, momentum=0.1, affine=True, track_r
unning_stats=True)
      (relu): ReLU(inplace=True)
      (conv2): Conv2d(256, 256, kernel_size=(3, 3), stride=(1, 1), padding=
(1, 1), bias=False)
      (bn2): BatchNorm2d(256, eps=1e-05, momentum=0.1, affine=True, track r
unning_stats=True)
      (downsample): Sequential(
        (0): Conv2d(128, 256, kernel_size=(1, 1), stride=(2, 2), bias=Fals
e)
        (1): BatchNorm2d(256, eps=1e-05, momentum=0.1, affine=True, track r
unning_stats=True)
    (1): BasicBlock(
      (conv1): Conv2d(256, 256, kernel_size=(3, 3), stride=(1, 1), padding=
(1, 1), bias=False)
      (bn1): BatchNorm2d(256, eps=1e-05, momentum=0.1, affine=True, track r
unning stats=True)
      (relu): ReLU(inplace=True)
      (conv2): Conv2d(256, 256, kernel_size=(3, 3), stride=(1, 1), padding=
(1, 1), bias=False)
      (bn2): BatchNorm2d(256, eps=1e-05, momentum=0.1, affine=True, track r
unning_stats=True)
    )
  )
  (layer4): Sequential(
    (0): BasicBlock(
      (conv1): Conv2d(256, 512, kernel size=(3, 3), stride=(2, 2), padding=
(1, 1), bias=False)
      (bn1): BatchNorm2d(512, eps=1e-05, momentum=0.1, affine=True, track r
unning_stats=True)
      (relu): ReLU(inplace=True)
      (conv2): Conv2d(512, 512, kernel size=(3, 3), stride=(1, 1), padding=
(1, 1), bias=False)
      (bn2): BatchNorm2d(512, eps=1e-05, momentum=0.1, affine=True, track_r
unning stats=True)
      (downsample): Sequential(
        (0): Conv2d(256, 512, kernel_size=(1, 1), stride=(2, 2), bias=Fals
e)
        (1): BatchNorm2d(512, eps=1e-05, momentum=0.1, affine=True, track r
unning_stats=True)
```

```
(1): BasicBlock(
                   (conv1): Conv2d(512, 512, kernel_size=(3, 3), stride=(1, 1), padding=
             (1, 1), bias=False)
                   (bn1): BatchNorm2d(512, eps=1e-05, momentum=0.1, affine=True, track_r
             unning stats=True)
                   (relu): ReLU(inplace=True)
                   (conv2): Conv2d(512, 512, kernel_size=(3, 3), stride=(1, 1), padding=
             (1, 1), bias=False)
                   (bn2): BatchNorm2d(512, eps=1e-05, momentum=0.1, affine=True, track_r
             unning_stats=True)
               )
               (avgpool): AdaptiveAvgPool2d(output_size=(1, 1))
               (fc): Linear(in_features=512, out_features=1000, bias=True)
             )
In [15]:
         #This step in the is cell is necessary because the resnet18 model was trained
             #on the imagenet dataset which has 1000 classes. Since we have just 3 classes
             #we have to change the last fully connected layers to have three output featu
             resnet18.fc = torch.nn.Linear(in_features=512, out_features=3)
             loss_fn = torch.nn.CrossEntropyLoss()
             optimizer = torch.optim.Adam(resnet18.parameters(), 1r=3e-5)
In [16]:

  | def show_preds():

                 resnet18.eval()
                 images, labels = next(iter(dl test))
                 outputs = resnet18(images)
                 _, preds = torch.max(outputs, 1)
                 show_images(images, labels, preds)
In [18]:
          # At this point the model has not been trained yet so we can see that the pre
             show preds()
                             pneumonia
                                          pneumonia
                                                         covid
                                                                     covid
                                                                                  covid
```

Training the Model

```
In [19]:

    def train(epochs):

                 print('Starting training..')
                 for e in range(0, epochs):
                     print('='*20)
                     print(f'Starting epoch {e + 1}/{epochs}')
                     print('='*20)
                     train_loss = 0.
                     val_loss = 0.
                     resnet18.train() # set model to training phase
                     for train_step, (images, labels) in enumerate(dl_train):
                         optimizer.zero_grad()
                         outputs = resnet18(images)
                         loss = loss_fn(outputs, labels)
                         loss.backward()
                         optimizer.step()
                         train_loss += loss.item()
                         if train_step % 20 == 0:
                             print('Evaluating at step', train_step)
                             accuracy = 0
                             resnet18.eval() # set model to eval phase
                             for val step, (images, labels) in enumerate(dl test):
                                  outputs = resnet18(images)
                                  loss = loss fn(outputs, labels)
                                  val loss += loss.item()
                                  _, preds = torch.max(outputs, 1)
                                  accuracy += sum((preds == labels).numpy())
                             val loss /= (val step + 1)
                             accuracy = accuracy/len(test dataset)
                             print(f'Validation Loss: {val_loss:.4f}, Accuracy: {accuracy:
                             show preds()
                             resnet18.train()
                             if accuracy >= 0.95:
                                  print('Performance condition satisfied, stopping..')
                                 return
                     train_loss /= (train_step + 1)
                     print(f'Training Loss: {train_loss:.4f}')
                 print('Training complete..')
```

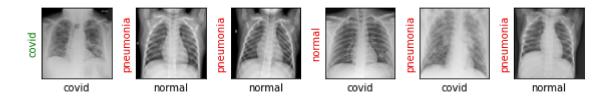
12/22/21, 1-36 PM A D S C IBM - J N

In [20]: ► | %%time

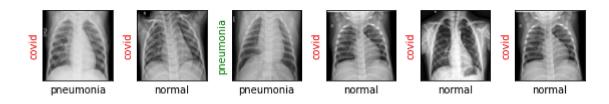
train(epochs=1)

Evaluating at step 0

Validation Loss: 1.0589, Accuracy: 0.3889



Evaluating at step 20 Validation Loss: 0.7467, Accuracy: 0.7333



Evaluating at step 40 Validation Loss: 0.4543, Accuracy: 0.9111



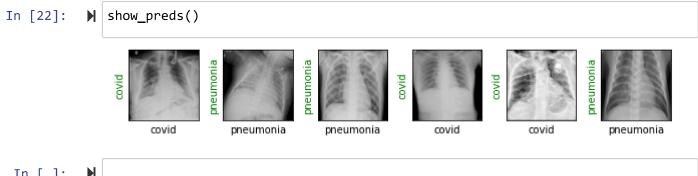
Evaluating at step 60

Validation Loss: 0.1844, Accuracy: 0.9778

Performance condition satisfied, stopping..

Wall time: 3min 24s

Final Results



In []: