coursework

February 21, 2024

1 Coursework: Self-supervised learning

In this coursework, you will explore the popular self-supervised contrastive learning approach Sim-CLR.

You will be asked to implement some of the key components of SimCLR, including a suitable data augmentation strategy (for generating positive pairs), the SimCLR loss function, and the SimCLR training step. Additionally, you will be using transfer learning strategies for evaluating the performance of different pre-trained models for a downstream classification task.

The coursework is divided into three-parts: - **Part A:** Implementation of a suitable dataset for contrastive model training; - **Part B:** Implementation of the SimCLR loss and training step; - **Part C:** Implementation of transfer learning strategies (linear probing and finetuning) for model evaluation.

Important: Read the text descriptions carefully and look out for hints and comments indicating a specific 'TASK'. Make sure to add sufficient documentation to your code.

Submission: You are asked to submit two versions of your notebook: 1. You should submit the raw notebook in .ipynb format with *all outputs cleared*. Please name your file coursework.ipynb. 2. Additionally, you will be asked to submit an exported version of your notebook in .pdf format, with *all outputs included*. We will primarily use this version for marking, but we will use the raw notebook to check for correct implementations. Please name this file coursework_export.pdf.

1.1 Your details

Please add your details below. You can work in groups up to two.

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1.2 Setup

```
[1]: # On Google Colab uncomment the following line to install PyTorch Lightning and the MedMNIST dataset
#!pip install lightning medmnist
```

```
[1]: import os import numpy as np import torch
```

1.3 Part A: Implement a dataset suitable for contrastive learning.

We will be using the MedMNIST Pneumonia dataset, which is a medical imaging inspired dataset but with the characteristics of MNIST. This allows efficient experimentation due to the small image size. The dataset contains real chest X-ray images but downsampled to 28 x 28 pixels, with binary labels indicating the presence of Pneumonia (which is an inflammation of the lungs).

1.3.1 Task A-1: Complete the dataset implementation.

You are asked to implement a dataset class SimCLRPneumoniaMNISTDataset suitable for training a self-supervised model with a contrastive objective. For each sample, your dataset class should return two 'views' of the corresponding image, forming the positive pairs for contrastive learning. It is up to you to design suitable augmentation pipeline for generating these views. Please provide a short description in plain language of what your data augmentation pipeline is meant to do.

To get you started, we have provided the skeleton of the dataset class in the cell below. Once you have implemented your dataset class, you are asked to run the provided visualisation code to visualise one batch of your training dataloader.

Note: You can use the same data augmentation pipeline for training, validation, and testing.

```
self.size = 28
       self.size_flag = ""
       self.root = './data/coursework/'
       self.info = INFO[self.flag]
       self.download()
      npz_file = np.load(os.path.join(self.root, "pneumoniamnist.npz"))
      self.split = split
       # Load all the images
       assert self.split in ['train','val','test']
      self.imgs = npz_file[f'{self.split}_images']
      self.labels = npz_file[f'{self.split}_labels']
       # TASK: Define here your data augmentation pipeline
       # Add a short description in plain language.
       ########EXPLANATION: ##############
       # As part of the data augmentation pipeline, we apply a series of
       #transformations to the dataset. For each image in the dataset, the
       #augmentation pipeline will be applied twice to generate two
       # distinct views of the same image to be used for contrastive learning.
       #We decided to apply color jttering effect to the images with a low
       #probability (i.e. 20%) which essentially adjusts the brightness and
\hookrightarrow the
       \#contrast of the image. We also applied a rather mild random resized
#to introduce scale and aspect ratio variability into the dataset.
       #Since we are working with medical images where the details are crucial,
       # we are less "aggressive" in cropping so that most of the image_
\hookrightarrow content
       #is preserved. We also apply random rotation within a small range
       #(+/- 10 degrees)
       # to encourage the model to become invariant to slight changes in
       # orientation.
       # We believe this transformation is particularly relevant to this.
\rightarrow dataset,
       #since the orientation might vary slightly due to patient positioning or
        #imaging set up at inference time, so by including this transformation
       #in the data augmentation pipeline
       # we hope that the model will become more robust in recognising \Box
       #of pneumonia by rexognisine the features of pneumonia across various
       #orientations.
       # We didn't apply gaussian blurring/ random noise because we find that
```

```
#the images are already heavily downsampled and we didn't want
       # to alter the signal further. Since we are working with grayscale
       #images, we did not apply any colour-related transformations since they
       #are not relevant.
       # Finally, the last tranformation is applied such that the images are
       #converted from PIL objects to tensors to be able to develop the deep_{\sqcup}
→ learning model
       # which operates on tensors.
      self.augmentation_pipeline = transforms.Compose([
           transforms.ToPILImage(),
           transforms.RandomApply(transforms=[transforms.
→ColorJitter(brightness=0.2, contrast=0.2)], p=0.1),
           transforms.RandomResizedCrop(size=(28, 28), scale=(0.9, 0.9)),
            # #relatively mild crop, preserving most of the integrity of the
⇒image
           transforms.RandomRotation(degrees=(10)),
           transforms.ToTensor()
      ])
  def len (self):
      return self.imgs.shape[0]
  def __getitem__(self, index):
       # TASK: Fill in the blanks such that you return two tensors
       # of shape [1, 28, 28], img_view1 and img_view2, representing two
       #augmented view of the images.
      img= self.imgs[index]
       img_view1, img_view2= self.augmentation_pipeline(img), self.
→augmentation_pipeline(img)
      return img view1, img view2
```

We use a LightningDataModule for handling your PneumoniaMNIST dataset. You do not need to make any modifications to the code below.

```
[12]: class SimCLRPneumoniaMNISTDataModule(LightningDataModule):
    def __init__(self, batch_size: int = 8):
        super().__init__()
        self.batch_size = batch_size
        self.train_set = SimCLRPneumoniaMNISTDataset(split='train')
        self.val_set = SimCLRPneumoniaMNISTDataset(split='val')
        self.test_set = SimCLRPneumoniaMNISTDataset(split='test')

def train_dataloader(self):
```

```
return DataLoader(dataset=self.train_set, batch_size=self.batch_size,ushuffle=True)

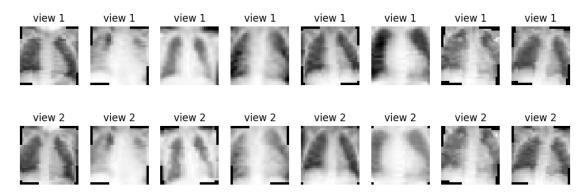
def val_dataloader(self):
    return DataLoader(dataset=self.val_set, batch_size=self.batch_size,ushuffle=False)

def test_dataloader(self):
    return DataLoader(dataset=self.test_set, batch_size=self.batch_size,ushuffle=False)
```

Check dataset implementation. Run the below cell to visualise a batch of your training dataloader.

```
[15]: # DO NOT MODIFY THIS CELL! IT IS FOR CHECKING THE IMPLEMENTATION ONLY.
      # Initialise data module
      datamodule = SimCLRPneumoniaMNISTDataModule()
      # Get train dataloader
      train dataloader = datamodule.train dataloader()
      # Get first batch
      batch = next(iter(train_dataloader))
      # Visualise the images
      view1, view2 = batch
      f, ax = plt.subplots(2, 8, figsize=(12,4))
      for i in range(8):
        ax[0,i].imshow(view1[i, 0], cmap='gray')
        ax[1,i].imshow(view2[i, 0], cmap='gray')
        ax[0,i].set title('view 1')
        ax[1,i].set_title('view 2')
        ax[0, i].axis("off")
        ax[1, i].axis("off")
```

Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz



1.4 Part B: Implement the SimCLR loss and training step.

In this part, we ask you to: 1. Implement the SimCLR loss function, as per the equation in the lecture notes (and the original paper). 2. Once you have implemented the loss, implement the training step function in the provided LightningModule.

1.4.1 Task B-1: SimCLR loss function.

For the implementation of the SimCLR loss, you should follow the 'recipe' from the lecture slides. We provide a code skeleton to get you started. Fill in all the blanks.

Hint: In PyTorch, to compute scalar products (also called dot products) between many elements efficiently, note that for two batches of d-dimensional feature vectors v1 and v2 of size [N, d] (with N being the batch size) computing the matrix multiplication torch.mm(v1, v2.t()) returns a matrix S of size [N, N] where each element S[i, j] is the scalar product of $v1_i$ and $v2_j$.

```
[6]: def simclr_loss(embedding_view1, embedding_view2, tau = 1.0):
       This funtion implements the SimCLR loss function as described in the original \Box
      \hookrightarrow paper.
       See lecture notes for formulas.
       It takes as input the embeddings from both views and returns the loss value,
      ⇔for that batch.
       Args:
         embedding_view1: torch tensor of shape [batch_size, embedding_dimension]
         embedding view2: torch tensor of shape [batch size, embedding dimension]
       Returns:
         loss: torch.tensor of shape 1
       device = embedding_view1.device
       # Step 1: normalise the embeddings
       similarity = torch.nn.CosineSimilarity(dim=1)
       out = similarity(embedding_view1, embedding_view2)
       embedding_view1 = F.normalize(embedding_view1, dim=1)
       embedding view2 = F.normalize(embedding view2, dim=1)
       # Step 2: gather all embeddings into one big vector of size [2*N ,,,
      → feature dim]
       z_all_views = torch.cat((embedding_view1, embedding_view2), dim=0)
       # Step 3: compute all possible similarities, should be a matrix of size [2 *_1
      \hookrightarrow N, 2 * N]
```

```
# all\_similarities[i,j] will be the similarity between z\_all\_views[i] and
\hookrightarrow z_all_views[j].
# Use the hint.
all_similarities = torch.mm(z_all_views, z_all_views.t())
all_similarities = torch.exp(all_similarities / tau)
# Step 4: Here we want to return a mask of size[2 * N, 2* N] for which
\hookrightarrow mask[i,j] = 1 if
\# z_{all\_views[i]} and z_{all\_views[j]} form a positive pair.
# There should be exactely 2 * N non-zeros elements in this matrix.
mask = torch.zeros(2 * embedding_view1.size(0), 2 * embedding_view1.size(0),__
⇒dtype=torch.bool, device=device)
for i in range(embedding_view1.size(0)):
  mask[i, i + embedding_view1.size(0)] = True
  mask[i + embedding_view1.size(0), i] = True
# Step 5: self-mask. For computing the denominator term in the loss function,
# we need to sum over all possible similarities except the self-similarity.
# Create a mask of shape [2*N, 2*N] that is 1 for all valid pairs and 0 for
\rightarrow all self-pairs (i = j).
self_mask = ~torch.eye(2 * embedding_view1.size(0), dtype=torch.bool,__
→device=device)
# Step 6: Computing all numerators for the loss function.
# Should be vector of size [2 * N],
# where element is exp(sim(i, j) / t) for each positive pair (i, j).
# Re-use the computed quantities above.
numerators = all_similarities[mask]
# Step 7: Computing all denominators for the loss function.
# Should be a vector of size [2 * N].
# Where each element should be the sum of \exp(\sin(i,k)/\tan) for all k = i.
denominators = all_similarities.masked_fill(~self_mask, 0)
denominators = denominators.sum(dim=1)
# Step 8: Return the final loss values, using the previously computing
umerators and denominators.
return -torch.log(numerators / denominators).mean()
```

Check SimCLR loss function. To check your implementation, please run the following tests. Note that we will also use other tests on different inputs to test your code.

```
[7]: # DO NOT MODIFY THIS CELL! IT IS FOR CHECKING THE IMPLEMENTATION ONLY.

seed_everything(33)
```

```
Seed set to 33
```

```
Expected loss: 1.7517999410629272, Computed loss: 1.751814365386963
Expected loss: 1.6375999450683594, Computed loss: 1.6376436948776245
Expected loss: 4.193999767303467, Computed loss: 4.194351673126221
Expected loss: 4.1753997802734375, Computed loss: 4.17537260055542
Passed all tests successfully!
```

1.4.2 Task B-2: SimCLR training step.

In this next task you are asked to complete the blanks in the provided LightningModule.

We provide the implementation of an image encoder (the CNN backbone that will act as feature extractor). No changes are needed for this part.

```
[8]: class ImageEncoder(torch.nn.Module):
         def __init__(self) -> None:
             super().__init__()
             self.net = models.resnet50(weights=None)
             del self.net.fc
             self.net.conv1 = torch.nn.Conv2d(1, 64, kernel_size=7, stride=2,__
      →padding=3, bias=False)
         def forward(self, x: torch.Tensor) -> torch.Tensor:
             x = self.net.conv1(x)
             x = self.net.bn1(x)
             x = self.net.relu(x)
             x0 = self.net.maxpool(x)
             x1 = self.net.layer1(x0)
             x2 = self.net.layer2(x1)
             x3 = self.net.layer3(x2)
             x4 = self.net.layer4(x3)
             x4 = self.net.avgpool(x4)
             x4 = torch.flatten(x4, 1)
             return x4
```

Next, you will need to complete the implementation of the SimCLR model. In order to make the training step work correctly, you will need to implement the process_batch function.

```
[9]: class SimCLRModel(LightningModule):
         def __init__(self, learning_rate: float = 0.001):
             super().__init__()
             self.learning_rate = learning_rate
             self.encoder = ImageEncoder()
             self.projector = torch.nn.Sequential(
                 torch.nn.Linear(2048, 1024),
                 torch.nn.ReLU(),
                 torch.nn.Linear(1024, 128),
             )
         def configure_optimizers(self):
             optimizer = torch.optim.Adam(self.parameters(), lr=self.learning rate)
             return optimizer
         def process_batch(self, batch):
             # TASK: Implement the process_batch function
             embedding_view1 = self.encoder(batch[0])
             embedding_view2 = self.encoder(batch[1])
             embedding_view1 = self.projector(embedding_view1)
             embedding_view2 = self.projector(embedding_view2)
             loss = simclr loss(embedding view1, embedding view2)
             return loss
         def training_step(self, batch, batch_idx):
             loss = self.process_batch(batch)
             self.log('train_loss', loss, prog_bar=True)
             if batch_idx == 0:
                 grid = torchvision.utils.make_grid(torch.cat((batch[0][0:4, ...],_
      ⇒batch[1][0:4, ...]), dim=0), nrow=4, normalize=True)
                 self.logger.experiment.add_image('train_images', grid, self.
      ⇔global_step)
             return loss
         def validation_step(self, batch, batch_idx):
             loss = self.process_batch(batch)
             self.log('val_loss', loss, prog_bar=True)
```

Check SimCLR training step. Here you can test that your code runs fine by training the model for 5 epochs using the cell below.

Report the training and validation loss at the end of 5 epochs.

```
[10]: # DO NOT MODIFY THIS CELL! IT IS FOR CHECKING THE IMPLEMENTATION ONLY.
      seed_everything(33, workers=True)
      data = SimCLRPneumoniaMNISTDataModule(batch_size=32)
      model = SimCLRModel()
      trainer = Trainer(
         max_epochs=5,
         accelerator='auto',
         devices=1,
         logger=TensorBoardLogger(save_dir='./lightning_logs/coursework/',u

¬name='simclr'),
          callbacks=[ModelCheckpoint(monitor='val_loss', mode='min'),__
       →TQDMProgressBar(refresh_rate=10)],
      trainer.fit(model=model, datamodule=data)
     Seed set to 33
     Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz
     Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz
     Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz
     GPU available: False, used: False
     TPU available: False, using: 0 TPU cores
     IPU available: False, using: 0 IPUs
     HPU available: False, using: 0 HPUs
                             | Params
       Name
               | Type
     0 | encoder | ImageEncoder | 23.5 M
     1 | projector | Sequential | 2.2 M
     25.7 M
              Trainable params
               Non-trainable params
     25.7 M
               Total params
     102.925 Total estimated model params size (MB)
                                 | 0/? [00:00<?, ?it/s]
     Sanity Checking: |
     /Users/kively/VSCodeProjects/Imperial_Projects/Machine_Learning_Imaging/.venv/li
     b/python3.12/site-
     packages/pytorch_lightning/trainer/connectors/data_connector.py:441: The
     'val dataloader' does not have many workers which may be a bottleneck. Consider
     increasing the value of the `num_workers` argument` to `num_workers=15` in the
     `DataLoader` to improve performance.
```

1.5 Part C: Linear probing and model finetuning.

In this part, you are given two different image encoders that were pre-trained with different datasets and training strategies. The objective for this task is to assess the performance of these two encoders in a downstream classification task. This this end, you are asked to implement evaluation routines seen in the lecture: linear probing and model finetuning. The downstream task is the prediction of Pneumonia in the (small) chest X-ray images from the PneumoniaMNIST dataset.

This part can be broken down into the following tasks: 1. Adapt your PneunomiaMNIST dataset for the image classification task. 2. Implement a classification model with a linear layer attached to a pre-trained image encoder. 3. For both pre-trained encoders: - a) Train the classifier on top of the frozen encoder (linear probing) - b) Finetune the entire model (including the encoder). 4. Evaluate all models on the test set, and provide a brief summary (no more than 300 words) with an analysis of your findings.

1.5.1 Task C-1: Adapt your PneunomiaMNIST dataset for the image classification task.

We can base our implementation largely on the SimCLRPneumoniaMNISTDataset and adapt it to make it suitable for image classification. Think about a suitable data augmentation pipeline. Check previous tutorials for inspiration.

```
self.info = INFO[self.flag]
      self.download()
      npz_file = np.load(os.path.join(self.root, "pneumoniamnist.npz"))
      self.split = split
      # Load all the images
      assert self.split in ['train','val','test']
      self.imgs = npz file[f'{self.split} images']
      self.labels = npz_file[f'{self.split}_labels']
      self.do_augment = augmentation
      # TASK: Define here your data augmentation pipeline suitable for
\hookrightarrow classification.
      # Check previous tutorials for inspiration.
       ########FXPI.ANATTON:#########
      #As part of the photometric data augmentation transformations, we apply
      #color jitter to adjust the brightness and contrast
      #with probablity 50%, hoping that the model will become more robust to
      #lighting variations.
      #As part of the geometric data augmentation, we introduce small _{\sqcup}
\rightarrowrotations
      #and slight scaling. These tranformations are useful
      #for simulating variations in patient positioning and/or the distance
      #between the X-ray source and the patient which might help
      #the model at test time. Nearest interpolation is applied to prevent any
       # unwanted smoothness in the images, as we ned to maintain
      #fine details in medical images.
      # photometric data augmentation
      self.photometric_augment = transforms.Compose([
          transforms.RandomApply(transforms=[transforms.
→ColorJitter(brightness=0.2, contrast=0.2)], p=0.5),
      1)
      # geometric data augmentation
      self.geometric_augment = transforms.Compose([
          transforms.RandomApply(transforms=[transforms.
→RandomAffine(degrees=5, scale=(0.9, 1.1), interpolation=transforms.
→InterpolationMode.NEAREST)], p=0.5),
      1)
```

```
def __len__(self):
      return self.imgs.shape[0]
  def __getitem__(self, index):
      # TASK: Implement the __getitem__ function to return the image and its_
⇔class label.
      image = torch.from_numpy(self.imgs[index]).unsqueeze(0)
      image = image.type(torch.FloatTensor)
      # label = torch.from_numpy(self.labels[index])
      label = self.labels[index]
      if self.do_augment:
          image = self.photometric_augment(image.type(torch.ByteTensor)).
→type(torch.FloatTensor)
          image = self.geometric_augment(image)
      # normalize image intensities to [0,1] to stabilise training
      image /= 255
      return image, label
```

Again, we use a LightningDataModule for handling your PneumoniaMNIST dataset. No changes needed for this part.

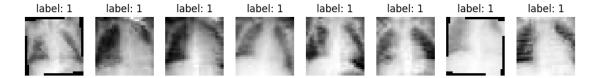
```
[12]: class PneumoniaMNISTDataModule(LightningDataModule):
          def __init__(self, batch_size: int = 32):
              super().__init__()
              self.batch size = batch size
              self.train_set = PneumoniaMNISTDataset(split='train', augmentation=True)
              self.val set = PneumoniaMNISTDataset(split='val', augmentation=False)
              self.test_set = PneumoniaMNISTDataset(split='test', augmentation=False)
          def train_dataloader(self):
              return DataLoader(dataset=self.train_set, batch_size=self.batch_size,__
       ⇔shuffle=True)
          def val_dataloader(self):
              return DataLoader(dataset=self.val_set, batch_size=self.batch_size,_u
       shuffle=False)
          def test_dataloader(self):
              return DataLoader(dataset=self.test_set, batch_size=self.batch_size,_
       ⇔shuffle=False)
```

Check dataset implementation. Run the below cell to visualise a batch of your training dataloader.

```
# DO NOT MODIFY THIS CELL! IT IS FOR CHECKING THE IMPLEMENTATION ONLY.

# Initialise data module
datamodule = PneumoniaMNISTDataModule()
# Get train dataloader
train_dataloader = datamodule.train_dataloader()
# Get first batch
batch = next(iter(train_dataloader))
# Visualise the images
images, labels = batch
f, ax = plt.subplots(1, 8, figsize=(12,4))
for i in range(8):
    ax[i].imshow(images[i, 0], cmap='gray')
    ax[i].set_title('label: ' + str(labels[i].item()))
    ax[i].axis("off")
```

Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz
Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz
Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz



1.5.2 Task C-2: Implement a classification model with a linear layer attached to a pre-trained image encoder.

We first download the weights of the two pre-trained image encoders. One of them has been trained with the self-supervised SimCLR objective on a large publicly available chest X-ray dataset (different from PneunomiaMNIST). The other encoder is a standard ImageNet backbone that has been trained with a supervised classification objective on the ImageNet dataset.

```
[14]: # ! wget https://www.doc.ic.ac.uk/~bglocker/teaching/mli/coursework.zip # ! unzip coursework.zip
```

We provide the function for loading the encoders. No changes needed here.

```
[15]: def load_encoder_from_checkpoint(checkpoint_path):
    ckpt = torch.load(checkpoint_path, map_location='cpu')
    simclr_module = SimCLRModel()
    print(simclr_module.load_state_dict(state_dict=ckpt))
    return simclr_module.encoder.eval()

imagenet_model = './data/coursework/model_imagenet.ckpt'
```

```
chestxray_model = './data/coursework/model_chestxray.ckpt'
```

Now, implement a classification model as a LightningModule for image classification using a pretrained image encoder.

The model should have a flag in the init function freeze_encoder that if set to true freezes all the weights in the encoder (used for linear probing), and if set to false all weights are trainable (used for model finetuning).

Hint: Check out previous tutorials for inspiration on how to implement a classification model as LightningModule. For the coursework, we recommend using the Area Under the Receiver Operating Characteristic Curve (ROC-AUC) performance metric (instead of accuracy). ROC-AUC is measure of the overall discriminative power of a classification model. You can use the readily available implementation in torchmetrics. You should log the ROC-AUC similar to how we logged accuracy in previous tutorials.

```
[18]: # TASK: Implement the ImageClassifier class
      # Check previous tutorials for insipration how to implement an `ImageClassifier`
      class ImageClassifier(LightningModule):
          def __init__(self, pretrained_encoder: torch.nn.Module, freeze_encoder:u
       ⇒bool = True, output_dim: int = 2, learning_rate: float = 0.001):
              super().__init__()
              self.learning_rate = learning_rate
              self.output_dim = output_dim
              self.freeze_encoder = freeze_encoder
              self.encoder = pretrained_encoder
              if self.freeze_encoder:
                  for param in self.encoder.parameters():
                      param.requires_grad = False
              self.classifier = torch.nn.Linear(2048, output_dim)
          def configure_optimizers(self):
              optimizer = torch.optim.Adam(self.parameters(), lr=self.learning rate)
              return optimizer
          def forward(self, x):
              x = self.encoder(x)
              x = self.classifier(x)
              return x
          def process_batch(self, batch):
              x, y = batch
              logits = self(x)
              loss = F.cross_entropy(logits, y.squeeze())
              probs = torch.softmax(logits, dim=1)
              aur = auroc(probs, y.squeeze(), task="multiclass", num_classes=2)
```

```
def training_step(self, batch, batch_idx):
    loss, aur = self.process_batch(batch)
    self.log('train_loss', loss, prog_bar=True)
    self.log('train_auroc', aur, prog_bar=True)
    return loss

def validation_step(self, batch, batch_idx):
    loss, aur = self.process_batch(batch)
    self.log('val_loss', loss, prog_bar=True)
    self.log('val_auroc', aur, prog_bar=True)

def test_step(self, batch, batch_idx):
    loss, aur = self.process_batch(batch)
    self.log('test_loss', loss, prog_bar=True)
    self.log('test_loss', loss, prog_bar=True)
    self.log('test_auroc', aur, prog_bar=True)
```

1.5.3 Task C-3a: Implement training and testing for linear probing.

Train two classification models using linear probing, one for each of the two provided image encoders. Evaluate on both the validation and test sets.

Note: Training for 25 epochs should be sufficient.

```
[19]: seed_everything(33, workers=True)
      data = PneumoniaMNISTDataModule(batch size=32)
      # TASK: Implement the linear probing training and testing routines.
      # Model 1: Linear probing with the ImageNet pre-trained encoder
      model =
       →ImageClassifier(pretrained_encoder=load_encoder_from_checkpoint(imagenet_model),

¬freeze_encoder=True)
      trainer = Trainer(
          max_epochs=25,
          accelerator='auto',
          devices=1,
          log_every_n_steps=5,
          logger=TensorBoardLogger(save_dir='./lightning_logs/coursework/',_
       ⇔name='linear_probe_imagenet'),
          callbacks=[ModelCheckpoint(monitor='val_loss', mode='min'),__
       →TQDMProgressBar(refresh rate=10)]
      trainer.fit(model=model, datamodule=data)
```

```
trainer.validate(model=model, datamodule=data, ckpt_path=trainer.
  ⇔checkpoint_callback.best_model_path)
trainer.test(model=model, datamodule=data, ckpt_path=trainer.
  →checkpoint_callback.best_model_path)
# Model 2: Linear probing with the SimCLR pre-trained encoder
  -، ImageClassifier(pretrained_encoder=load_encoder_from_checkpoint(chestxray_model),

¬freeze_encoder=True)
trainer = Trainer(
    max_epochs=25,
    accelerator='auto',
    devices=1,
    log_every_n_steps=5,
    logger=TensorBoardLogger(save_dir='./lightning_logs/coursework/',_
 ⇔name='linear_probe_chestxray'),
    callbacks=[ModelCheckpoint(monitor='val_loss', mode='min'),_
 →TQDMProgressBar(refresh_rate=10)]
)
trainer.fit(model=model, datamodule=data)
trainer.validate(model=model, datamodule=data, ckpt_path=trainer.

¬checkpoint_callback.best_model_path)
trainer.test(model=model, datamodule=data, ckpt_path=trainer.
  →checkpoint_callback.best_model_path)
Seed set to 33
Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz
Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz
Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz
GPU available: True (mps), used: True
TPU available: False, using: 0 TPU cores
IPU available: False, using: 0 IPUs
HPU available: False, using: 0 HPUs
           | Type
                          | Params
0 | encoder | ImageEncoder | 23.5 M
1 | classifier | Linear | 4.1 K
4.1 K
         Trainable params
23.5 M
         Non-trainable params
23.5 M Total params
94.023
         Total estimated model params size (MB)
<All keys matched successfully>
Sanity Checking: | 0/? [00:00<?, ?it/s]
```

```
Training: |
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Validation: |
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Validation: |
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`Trainer.fit` stopped: `max_epochs=25` reached.
```

Restoring states from the checkpoint path at ./lightning_logs/coursework/linear_ probe_imagenet/version_8/checkpoints/epoch=7-step=1184.ckpt

Loaded model weights from the checkpoint at ./lightning_logs/coursework/linear_p robe_imagenet/version_8/checkpoints/epoch=7-step=1184.ckpt

Validation: | | 0/? [00:00<?, ?it/s] Restoring states from the checkpoint path at ./lightning_logs/coursework/linear_probe_imagenet/version_8/checkpoints/epoch=7-step=1184.ckpt
Loaded model weights from the checkpoint at ./lightning_logs/coursework/linear_probe_imagenet/version_8/checkpoints/epoch=7-step=1184.ckpt

Validate metric	DataLoader O				
val_auroc val_loss	0.9082255959510803 0.35509124398231506				
Testing: 0/? [[00:00 , ?it/s]</td				
Test metric	DataLoader 0				
test_auroc test_loss	0.7919484376907349 0.5793928503990173				
GPU available: True (mps), used: True TPU available: False, using: 0 TPU cores IPU available: False, using: 0 IPUs HPU available: False, using: 0 HPUs Name					
0 encoder ImageEncoder 23.5 M 1 classifier Linear 4.1 K					
4.1 K Trainable params 23.5 M Non-trainable params 23.5 M Total params 94.023 Total estimated model params size (MB)					
<all keys="" matched="" successfully=""></all>					
Sanity Checking: 0/? [00:00 , ?it/s]</td					
Training: 0/? [00:00 , ?it/s]</td					
Validation: 0/? [00:00 , ?it/s]</td					
Validation: 0/? [00:00 , ?it/s]</td					
Validation: 0/? [00:00 , ?it/s]</td					

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```

Restoring states from the checkpoint path at ./lightning_logs/coursework/linear_probe_chestxray/version_7/checkpoints/epoch=12-step=1924.ckpt
Loaded model weights from the checkpoint at ./lightning_logs/coursework/linear_probe_chestxray/version_7/checkpoints/epoch=12-step=1924.ckpt

Validation: | 0/? [00:00<?, ?it/s]

Restoring states from the checkpoint path at ./lightning_logs/coursework/linear_probe_chestxray/version_7/checkpoints/epoch=12-step=1924.ckpt
Loaded model weights from the checkpoint at ./lightning_logs/coursework/linear_probe_chestxray/version_7/checkpoints/epoch=12-step=1924.ckpt

Validate metric DataLoader 0

[`]Trainer.fit` stopped: `max_epochs=25` reached.

1.5.4 Task C-3b: Implement training and testing for model finetuning.

Repeat the experiments, but this time using model finetuning instead of linear probing. Evaluate on both the validation and test sets.

```
[98]: seed_everything(33, workers=True)
      data = PneumoniaMNISTDataModule(batch_size=32)
      # TASK: Implement the model finetuning training and testing routines.
      # Model 1: Finetuning with the ImageNet pre-trained encoder
       →ImageClassifier(pretrained_encoder=load_encoder_from_checkpoint(imagenet_model), __

¬freeze_encoder=False)
      trainer = Trainer(
          max_epochs=25,
          accelerator='auto',
          devices=1,
          log_every_n_steps=5,
          logger=TensorBoardLogger(save_dir='./lightning_logs/coursework/',_
       ⇔name='finetune_imagenet'),
          callbacks=[ModelCheckpoint(monitor='val_loss', mode='min'),_
       →TQDMProgressBar(refresh_rate=10)]
      trainer.fit(model=model, datamodule=data)
      trainer.validate(model=model, datamodule=data, ckpt_path=trainer.
       checkpoint_callback.best_model_path)
```

```
trainer.test(model=model, datamodule=data, ckpt_path=trainer.
 ⇔checkpoint_callback.best_model_path)
# Model 2: Finetuning with the SimCLR pre-trained encoder
model =
 → ImageClassifier(pretrained encoder=load encoder from checkpoint(chestxray model),
 ⇔freeze encoder=False)
trainer = Trainer(
    \max_{epochs=25},
    accelerator='auto',
    devices=1,
    log_every_n_steps=5,
    logger=TensorBoardLogger(save_dir='./lightning_logs/coursework/',u
 →name='finetune_chestxray'),
    callbacks=[ModelCheckpoint(monitor='val_loss', mode='min'),__
 →TQDMProgressBar(refresh_rate=10)]
trainer.fit(model=model, datamodule=data)
trainer.validate(model=model, datamodule=data, ckpt_path=trainer.
 Green checkpoint_callback.best_model_path)
trainer.test(model=model, datamodule=data, ckpt_path=trainer.
  →checkpoint_callback.best_model_path)
Seed set to 33
Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz
Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz
Using downloaded and verified file: ./data/coursework/pneumoniamnist.npz
GPU available: True (mps), used: True
TPU available: False, using: 0 TPU cores
IPU available: False, using: 0 IPUs
HPU available: False, using: 0 HPUs
Missing logger folder: ./lightning_logs/coursework/finetune_imagenet
  | Name
              | Type
                              | Params
0 | encoder | ImageEncoder | 23.5 M
1 | classifier | Linear
23.5 M
         Trainable params
         Non-trainable params
23.5 M
         Total params
94.023
         Total estimated model params size (MB)
<All keys matched successfully>
Sanity Checking: | 0/? [00:00<?, ?it/s]
```

```
Training: |
                     | 0/? [00:00<?, ?it/s]
                        | 0/? [00:00<?, ?it/s]
Validation: |
Validation: |
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```

Restoring states from the checkpoint path at ./lightning_logs/coursework/finetune_imagenet/version_0/checkpoints/epoch=6-step=1036.ckpt

Loaded model weights from the checkpoint at ./lightning_logs/coursework/finetune _imagenet/version_0/checkpoints/epoch=6-step=1036.ckpt

Validation: | 0/? [00:00<?, ?it/s]

Restoring states from the checkpoint path at ./lightning_logs/coursework/finetune_imagenet/version_0/checkpoints/epoch=6-step=1036.ckpt

[`]Trainer.fit` stopped: `max_epochs=25` reached.

Validate metric DataLoader 0

val_auroc 0.9960390329360962 val_loss 0.07328185439109802

Loaded model weights from the checkpoint at ./lightning_logs/coursework/finetune _imagenet/version_0/checkpoints/epoch=6-step=1036.ckpt

Testing: | 0/? [00:00<?, ?it/s]

Test metric DataLoader 0

test_auroc 0.9606674313545227 test_loss 0.32537344098091125

GPU available: True (mps), used: True TPU available: False, using: 0 TPU cores

IPU available: False, using: 0 IPUs HPU available: False, using: 0 HPUs

Missing logger folder: ./lightning_logs/coursework/finetune_chestxray

	Name		Туре		Params
	encoder classifier		ImageEncoder Linear		23.5 M 4.1 K

23.5 M Trainable params
0 Non-trainable params

23.5 M Total params

94.023 Total estimated model params size (MB)

<All keys matched successfully>

Sanity Checking: | 0/? [00:00<?, ?it/s]

Training: | 0/? [00:00<?, ?it/s]
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```

Restoring states from the checkpoint path at ./lightning_logs/coursework/finetune_chestxray/version_0/checkpoints/epoch=21-step=3256.ckpt

Loaded model weights from the checkpoint at ./lightning_logs/coursework/finetune_chestxray/version_0/checkpoints/epoch=21-step=3256.ckpt

DataLoader 0

Validation: | 0/? [00:00<?, ?it/s]

Validate metric

val_auroc	0.9958899021148682
val loss	0.07362958043813705

[`]Trainer.fit` stopped: `max_epochs=25` reached.

Restoring states from the checkpoint path at ./lightning_logs/coursework/finetune_chestxray/version_0/checkpoints/epoch=21-step=3256.ckpt
Loaded model weights from the checkpoint at ./lightning_logs/coursework/finetune_chestxray/version_0/checkpoints/epoch=21-step=3256.ckpt

Testing: | 0/? [00:00<?, ?it/s]

Test metric DataLoader 0

test_auroc 0.9633634686470032 test_loss 0.46492835879325867

[98]: [{'test loss': 0.46492835879325867, 'test auroc': 0.9633634686470032}]

1.5.5 Task C-4: Your evaluation report.

Provide a brief summary (no more than 300 words) with an analysis of your findings. Try explaining the observed performance.

The comparison between two image encoders—SimCLR, trained on chest X-rays, and an ImageNettrained encoder—on the PneumoniaMNIST dataset highlights the impact of domain-specific pretraining and contrastive learning. The SimCLR encoder outperforms its ImageNet counterpart (AUROC of 0.89 vs. 0.79 in linear probing reported on the test set) due to its training on a relevant domain, capturing features better suited for medical tasks like pneumonia detection.

Contrastive learning's role in the SimCLR encoder's success underscores the value of learning discriminative features through augmentation strategies. It is worth noting that upon fine-tuning both encoders on the PneumoniaMNIST dataset, the initial performance gap narrows, with both models achieving around 0.96 AUROC. This convergence suggests that fine-tuning effectively mitigates the domain-specific advantage by allowing both models to adjust to the task at hand (i.e. pneumonia detection in chest X-ray).

The choice of AUROC as a performance metric over accuracy is particularly justified in this medical imaging context where the cost of false negatives is critical. AUROC provides a more "refined" measure of model performance, especially useful in datasets with class imbalance, offering a more reliable assessment than accuracy. Accuracy is known to be "susceptible" to predicting the majority class, whereas AUROC is better suited for medical tasks where there is often class imbalance and failing to detect a condition when it is present, might have severe consequences.

In conclusion, the initial advantage of the SimCLR encoder showcases the benefits of domain-specific pre-training and contrastive learning in medical imaging. However, the closely matched performance post-fine-tuning emphasizes the potential of both models to adapt to the task at hand.

1.6 Logging

```
[]: %load_ext tensorboard %tensorboard --logdir './lightning_logs/coursework/'
```