# Masked Auto-Encoder

We will explore the popular self-supervised masked auto-encoder approach MAE.

The work is divided in the following parts:

- Part A: Create a dataset and a data module to handle the PneumoniaMNIST dataset.
- Part B: Implement MAE utility functions.
- Part C: Implement and train a full MAE model.
- Part D: Inspect the trained model.

# Setup

```
In [24]: # On Google Colab uncomment the following line to install PyTorch Lightning
#! pip install lightning medmnist timm
```

```
In [25]: import os
         import numpy as np
         import torch
         import torch.nn as nn
         import torch.nn.functional as F
         import torchvision
         import matplotlib.pyplot as plt
         from torch.utils.data import DataLoader
         from torchvision import models
         from torchvision import transforms
         from pytorch lightning import LightningModule, LightningDataModule, Trainer,
         from pytorch lightning.loggers import TensorBoardLogger
         from pytorch lightning.callbacks import ModelCheckpoint, TQDMProgressBar
         from torchmetrics.functional import auroc
         from PIL import Image
         from medmnist.info import INFO
         from medmnist.dataset import MedMNIST
```

# **Part A:** Create a dataset and a data module to handle the PneumoniaMNIST dataset.

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 here downsampled to 28 x 28 pixels, with binary labels indicating the presence of Pneumonia (which is an inflammation of the lungs).

#### A-1. Complete the dataset implementation.

We implement a dataset class **PneumoniaMNISTDataset** suitable for training a classification model. For each sample, your dataset class should return one image and the corresponding label. We won't use the labels during training but for simplicity we will return them for model inspection purposes (part D).

In terms of augmentation, we want to follow what has been done in the original MAE paper, that is **use random cropping (70%-100%) and horizontal flipping only** (see paragraph Data augmentation, page 6 of the paper for further details).

```
In [26]: class PneumoniaMNISTDataset(MedMNIST):
             def __init__(self, split = 'train', augmentation: bool = False):
                 ''' Dataset class for Pneumonia MNST.
                 The provided init function will automatically download the necessary
                 files at the first class initialistion.
                 :param split: 'train', 'val' or 'test', select subset
                 1.1.1
                 self.flag = "pneumoniamnist"
                 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']
                 self.do augment = augmentation
                 # Transformations
                 self.transform = transforms.Compose([
                     transforms.ToPILImage(), # Convert from numpy to PIL
                     transforms.RandomResizedCrop(size=self.size, scale=(0.7, 1.0)),
                     transforms.RandomHorizontalFlip(p=0.5), # Randomly flip the imag
                     transforms.ToTensor() # Convert from PIL to tensor
                 ]) if self.do augment else transforms.ToTensor() # If there is no a
             def len (self):
                 return self.imqs.shape[0]
             def getitem (self, index):
                 img = self.imgs[index] # Get the image from index
                 label = self.labels[index] # Get the label from index
```

```
img = self.transform(img) # Apply the transformation to the image
return img, label
```

We use a LightningDataModule for handling your PneumoniaMNIST dataset. No changes needed for this part.

```
class PneumoniaMNISTDataModule(LightningDataModule):
    def __init__(self, batch_size: int = 32):
        super().__init__()
        self.batch_size = batch_size
        self.train_set = PneumoniaMNISTDataset(split='train', augmentation=I
        self.val_set = PneumoniaMNISTDataset(split='val', augmentation=False
        self.test_set = PneumoniaMNISTDataset(split='test', augmentation=Fal

    def train_dataloader(self):
        return DataLoader(dataset=self.train_set, batch_size=self.batch_size)

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

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

#### **Check** dataset implementation.

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

```
In [28]: # 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
                                                  label: 1
          label: 1
                    label: 0
                              label: 1
                                        label: 1
                                                            label: 1
                                                                       label: 1
                                                                                 label: 1
```

# **Part B**: Implement MAE utility functions.

Masked Auto-Encoders are based on a Vision Transformer (ViT) architecture. Importantly, the ViT architecture operates on a patch-level, not on the image-level. Hence, to feed the image into the ViT based encoder first we need to divide the images in small patches (typically 16x16 pixels).

In this part, we ask you to write three utility functions:

- patchify: takes in a batch of images (N, C, H, W) where N is the batch size, and returns a batch of patches of size (N, L, D) where L is the number of patches fitting in one image and D = patch\_size\*\* 2\*C.
- unpatchify: inverts the above operation, takes in a batch of patches of size (N, L, D) and returns the corresponding a batch of images (N, C, H, W).
- random\_masking: Randomly masks out patches during training to create a self-supervised training task of patch prediction.

#### Task B-1: Implement patchify

```
In [29]: def patchify(imgs, patch size):
                 patch_size: (patch_h, patch_w)
                 ### ADD YOUR CODE HERE
                 N, C, H, W = imgs.shape
                 patch h, patch w = patch size
                 # Check if H and W are divisible by patch size
                 assert H % patch h == 0, "H is not divisible by patch h"
                 assert W % patch w == 0, "W is not divisible by patch w"
                 # Calculate number of patches
                 L = (H // patch h) * (W // patch w)
                 # Extract patches
                 patches = imgs.unfold(2, patch h, patch h).unfold(3, patch w, patch
                 # Calculate D
                 D = patch h * patch w * C
                 # Reshape patches to (N, L, D)
                 patches = patches.reshape(N, L, D)
                 return patches
```

Let's test our implementation on the first batch of the validation set.

```
In [30]: # Load a batch of validation images
datamodule = PneumoniaMNISTDataModule()
dataloader = datamodule.val_dataloader()
```

```
batch = next(iter(dataloader))
    images, labels = batch

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

In [31]: images.shape

Out[31]: torch.Size([32, 1, 28, 28])

In [32]: # Assuming a patch size of (4,4) test your patchify function
    # and test that the shape of the outputs corresponds at what is expected patch_size = (4,4)
    patches = patchify(images, (4, 4))

In [33]: patches.shape

Out[33]: torch.Size([32, 49, 16])
```

#### Visualisation of patchify output

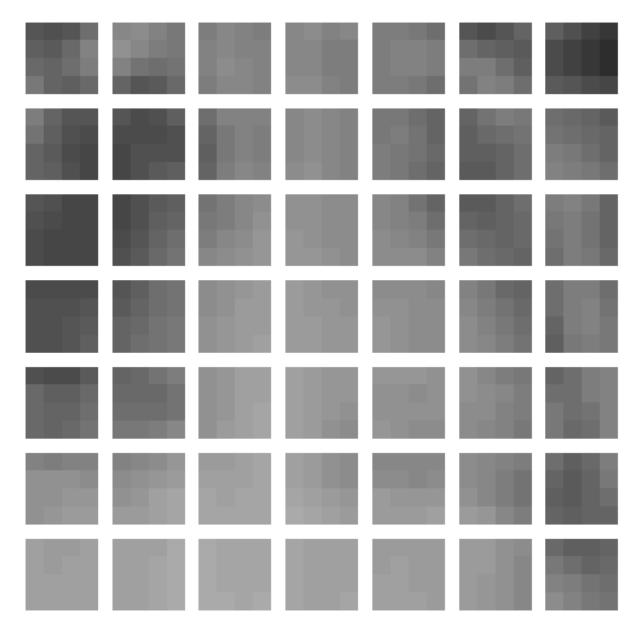
Next, we want to check our output visually. In the next cell, plot all the patches of the first image in the batch as a grid of subplots where subplot(i,j) shows patch(i,j) at the right position in the original image. You should be able to recognise the original image.

```
In [34]: # Plot the patches of the first image

patches1 = patches[0]
plot_dim = int(np.sqrt(patches1.shape[0]))

_, ax = plt.subplots(plot_dim, plot_dim, figsize=(8,8))

for i in range(plot_dim):
    for j in range(plot_dim):
        # Fix the color range to [0, 1]
        ax[i, j].imshow(patches1[i*plot_dim+j].reshape(patch_size[0], patch_ #ax[i, j].imshow(patches1[i*plot_dim+j].reshape(patch_size[0], patch_ ax[i, j].axis('off')
```



Compare the ouput with the original image

```
In [35]: plt.imshow(images[0][0], cmap='gray')
plt.axis('off')

Out[35]: (-0.5, 27.5, 27.5, -0.5)
```



### Task B-2: Implement unpatchify

Next, you are asked to create the reverse function able to take in a batch of patches and return the corresponding batch of images.

```
In [36]: def unpatchify(patches, patch_size, image_size, number_of_channels=1):
    ### TODO
    ### Write a function that takes a batch of patches (N, L, D) where D = patches and returns the batch of images (N, C, H, W)
    ### ADD YOUR CODE HERE

N, L, _ = patches.shape
    patch_h, patch_w = patch_size
    img_h, img_w = image_size

# Check if L matches the expected number of patches
    assert L == (img_h // patch_h) * (img_w // patch_w), "Number of patches

# Reshape patches to (N, H // patch_h, W // patch_w, patch_h, patch_w, C, patches = patches.view(N, img_h // patch_h, img_w // patch_w, patch_h, patch_h
```

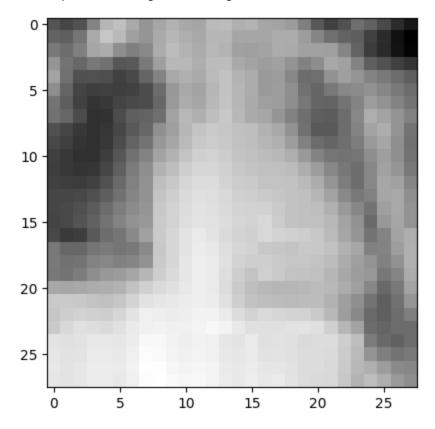
Check that after unpatchifying the patches obtained in the last cells, we get back to the original image batch.

```
In [37]: assert (unpatchify(patches, (4,4), (28,28)) == images).all()

### TODO plot the first image after applying patchify and unpatchify
### ADD YOUR CODE HERE

# Plot the original image
images = unpatchify(patches, (4, 4), (28, 28))
plt.imshow(images[0][0], cmap='gray')
```

Out[37]: <matplotlib.image.AxesImage at 0x72359df59bd0>



Task B-3: Implement random\_masking

Next we need to write the function that will randomly mask out some of the patches for the encoder. We want to follow the approach described in the paper:

Simple implementation. Our MAE pre-training can be implemented efficiently, and importantly, does not require any specialized sparse operations. First we generate a token for every input patch (by linear projection with an added positional embedding). Next we randomly shuffle the list of tokens and remove the last portion of the list, based on the masking ratio. This process produces a small subset of tokens for the encoder and is equivalent to sampling patches without replacement. After encoding, we append a list of mask tokens to the list of encoded patches, and unshuffle this full list (inverting the random shuffle operation) to align all tokens with their targets. The decoder is applied to this full list (with positional embeddings added). As noted, no sparse operations are needed. This simple implementation introduces negligible overhead as the shuffling and unshuffling operations are fast.

We implement the random\_masking function.

This function takes the original patched batch of size (N, L, D) as input and returns:

- (a) patches kept : the sequence of non-masked tokens
- (b) mask: a binary mask indicating which grid position are masked for every image in the batch
- (c) ids\_restore: list of indices indicating how to revert the patch shuffling operation used to create the mask.

```
In [38]:

def random_masking(patches, mask_ratio):
    """

Args:
    patches: original patched batch of size (N, L, D)
    mask_ratio: float between 0 and 1, the proportion of patches to

Returns:
    patches_kept: tensor (N, L_kept, D) the sequence of non-masked pask: tensor (N, L) binary mask indicating which positions are mask: tensor (N, L) list of indices indicating how to un-
"""

N, L, D = patches.shape # batch, length, dim

# Step 1: create noise in [0, 1]
### ADD YOUR CODE HERE
```

```
noise = torch.rand(N, L)
# Step 2: sort noise for each sample
### ADD YOUR CODE HERE
noise, indices = noise.sort(dim=1)
# Step 3: store list of indices to revert shuffling operation later
### ADD YOUR CODE HERE
ids restore = torch.argsort(indices, dim=1)
# Step 4: used shuffled list to keep only a subset of patches
### ADD YOUR CODE HERE
# Calculate the number of patches to keep
L \text{ kept} = \text{int}(L * (1 - \text{mask ratio}))
# Select the indices of the patches to keep (the first L kept after
indices keep = indices[:, :L kept]
# Extract the patches to keep using gather
patches kept = torch.gather(patches, dim=1, index=indices keep.unsqu
# Step 5 : generate the binary mask
### ADD YOUR CODE HERE
# First, in the mixed order: the first L kept are 0 (unmasked) and t
mask shuffled = torch.ones(N, L, device=patches.device)
mask shuffled[:, :L kept] = 0
# Revert the permutation so that the mask corresponds to the original
mask = torch.gather(mask shuffled, dim=1, index=ids restore)
return patches kept, mask, ids restore
```

```
In [39]: patches_kept, mask, ids_restore = random_masking(patches, 0.75)
```

Check the shapes of our outputs. Are there as expected?

```
In [40]: patches_kept.shape, mask.shape, ids_restore.shape
```

```
Out[40]: (torch.Size([32, 12, 16]), torch.Size([32, 49]), torch.Size([32, 49]))
```

The original input is a batch of images represented as patches with a shape of (32, 49, 16), where 32 is the batch size, 49 represents the number of patches per image (which fits a  $28\times28$  image when divided into a  $7\times7$  grid), and 16 is patch\_size\*\*2\*C. When applying a mask ratio of 0.75, only 25% of the patches are retained. Since 25% of 49 is approximately 12.25, the function truncates this value to 12 patches per image, resulting in a tensor for patches\_kept with a shape of (32, 12, 16).

Additionally, the mask tensor is created to indicate which of the 49 patches in each image are masked (represented by 1s) or not (0s), and hence it retains the original patch count,

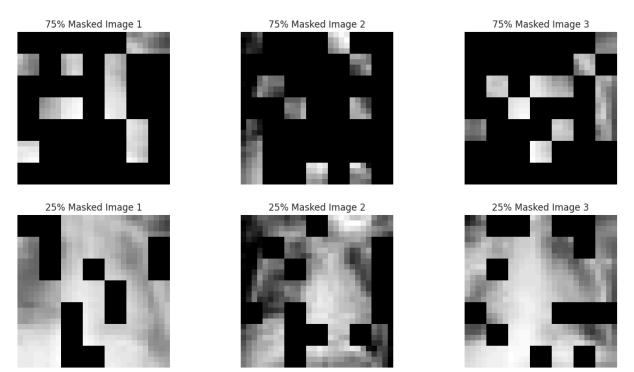
yielding a shape of (32, 49). The ids\_restore tensor, which is used to restore the original order of the patches, also has one entry per original patch, leading to the same shape of (32, 49).

Therefore, all the output shapes are as expected.

#### Visualisation of random masking

In this cell, we ask you to use the previously implemented functions <code>patchify</code>, <code>unpatchify</code> and <code>random\_masking</code> to visualise the first three images in the validation batch at a masking ratio of 75% and 25%. Create a 2 x 3 subplots grids, the first row should be masked at 75%, the second one at 25%

```
In [63]: patch size = (4,4)
         images, = next(iter(datamodule.val dataloader()))
         batch size = images.shape[0]
         f, ax = plt.subplots(2, 3, figsize=(15, 8))
         # Masking ratio 75%
         for i in range(3):
             # Select the i-th image and add the batch dimension
             img = images[i].unsqueeze(0) # Shape: (1, C, 28, 28)
             # Divide the image into patches
             patches = patchify(img, patch size) # (1, total patches, D)
             # Apply random masking
             patches kept, mask, ids restore = random masking(patches, 0.75)
             patches restored = torch.zeros((1, patches.shape[1], patches.shape[2]),
             patches restored[:, :patches kept.shape[1], :] = patches kept
             patches restored = patches restored[:, ids restore[0]]
             # Reconstruct the image from the patches
             reconstructed = unpatchify(patches restored, patch size, (28,28))
             # For grayscale images, we extract the only channel (reconstructed[0][0]
             ax[0, i].imshow(reconstructed[0][0].cpu().numpy(), cmap='gray')
             ax[0, i].set_title(f'75% Masked Image {i+1}')
             ax[0, i].axis('off')
         # Masking ratio 25%
         for i in range(3):
             img = images[i].unsqueeze(0) # (1, C, 28, 28)
             patches = patchify(img, patch size)
             patches kept, mask, ids restore = random masking(patches, 0.25)
             patches restored = torch.zeros((1, patches.shape[1], patches.shape[2]),
             patches restored[:, :patches kept.shape[1], :] = patches kept
             patches restored = patches restored[:, ids restore[0]]
             reconstructed = unpatchify(patches restored, patch size, (28,28))
             ax[1, i].imshow(reconstructed[0][0].cpu().numpy(), cmap='gray')
             ax[1, i].set title(f'25% Masked Image {i+1}')
             ax[1, i].axis('off')
         plt.show()
```



# Part C: Implement and train a full MAE model.

We will use the previously defined utility functions along with some helper code that we provide to implement the full training pipeline of Masked Auto-Encoder.

In the following, we provide code for creating the positional embeddings for the ViT. You do not need to implement anything here, just run this cell.

```
In [42]: from functools import partial
         import torch
         import torch.nn as nn
         from timm.models.vision transformer import PatchEmbed, Block
         import numpy as np
         def get 2d sincos pos embed(embed dim, grid size, cls token=False):
             grid size: int of the grid height and width
             return:
             pos_embed: [grid_size*grid_size, embed dim] or [1+grid size*grid size, ε
             if isinstance(grid size, int):
                 grid size = (grid size, grid size)
             grid h = np.arange(grid size[0], dtype=np.float32)
             grid_w = np.arange(grid_size[1], dtype=np.float32)
             grid = np.meshgrid(grid w, grid h) # here w goes first
             grid = np.stack(grid, axis=0)
             grid = grid.reshape([2, 1, grid size[0], grid size[1]])
```

```
# use half of dimensions to encode grid h
   emb h = get 1d sincos pos embed from grid(embed dim // 2, grid[0]) # (F
   emb w = get 1d sincos pos embed from grid(embed dim // 2, grid[1]) # (F)
   pos embed = np.concatenate([emb h, emb w], axis=1) # (H*W, D)
   if cls token:
        pos embed = np.concatenate([np.zeros([1, embed dim]), pos embed], ax
    return pos embed
def get 1d sincos pos embed from grid(embed dim, pos):
   embed dim: output dimension for each position
   pos: a list of positions to be encoded: size (M,)
   out: (M, D)
   0.000
   assert embed dim % 2 == 0
   omega = np.arange(embed dim // 2, dtype=np.float32)
   omega /= embed dim / 2.0
   omega = 1.0 / 10000**omega # (D/2,)
   pos = pos.reshape(-1) # (M,)
   out = np.einsum("m,d->md", pos, omega) # (M, D/2), outer product
   emb \sin = np.sin(out) # (M, D/2)
   emb cos = np.cos(out) # (M, D/2)
   emb = np.concatenate([emb sin, emb cos], axis=1) # (M, D)
    return emb
```

### Task C-1: MAE model implementation

```
In [43]: class MaskedAutoencoderViT(LightningModule):
             def init (
                 self,
                 img size=224,
                 patch size=16,
                 in chans=3,
                 embed dim=1024,
                 depth=24,
                 num heads=16,
                 decoder embed dim=512,
                 decoder depth=8,
                 decoder num heads=16,
                 mlp ratio=4.0,
             ):
                 super(). init ()
                 # MAE encoder definition
                 self.embed dim = embed dim
                 self.in chans = in chans
                 self.patch embed = PatchEmbed(img size, patch size, in chans, embed
```

```
num patches = self.patch embed.num patches
print(self.patch embed.grid size)
self.cls token = nn.Parameter(torch.zeros(1, 1, embed dim))
self.pos embed = nn.Parameter(
    torch.zeros(1, num patches + 1, embed dim), requires grad=False
self.blocks = nn.ModuleList(
        Block(
            embed dim,
            num heads,
            mlp ratio,
            qkv bias=True,
            norm layer=nn.LayerNorm,
        for i in range(depth)
    1
self.norm = nn.LayerNorm(embed dim)
# MAE decoder definition
self.decoder embed = nn.Linear(embed dim, decoder embed dim, bias=Tr
self.mask token = nn.Parameter(torch.zeros(1, 1, decoder embed dim))
self.decoder pos embed = nn.Parameter(
    torch.zeros(1, num patches + 1, decoder embed dim), requires gra
self.decoder blocks = nn.ModuleList(
        Block(
            decoder embed dim,
            decoder num heads,
            mlp ratio,
            gkv bias=True,
            norm layer=nn.LayerNorm,
        for i in range(decoder depth)
    ]
self.decoder norm = nn.LayerNorm(decoder embed dim)
self.decoder pred = nn.Linear(
    decoder embed dim, patch size**2 * in chans, bias=True
# Positional embeddings
pos embed = get 2d sincos pos embed(
    embed dim=self.pos embed.shape[-1],
    grid size=self.patch embed.grid size,
    cls token=True,
self.pos embed.data.copy (torch.from numpy(pos embed).float().unsque
decoder pos embed = get 2d sincos pos embed(
    self.decoder pos embed.shape[-1],
```

```
grid size=self.patch embed.grid size,
        cls token=True,
    self.decoder pos embed.data.copy (
        torch.from numpy(decoder pos embed).float().unsqueeze(0)
def patchify(self, imgs):
    imgs: (N, C, H, W)
    x: (N, L, D)
    ### TODO: Use the previously defined function
    ### ADD YOUR CODE HERE
    N, C, H, W = imgs.shape
    patch h, patch w = patch size
    # Check if H and W are divisible by patch size
    assert H % patch_h == 0, "H is not divisible by patch h"
    assert W % patch w == 0, "W is not divisible by patch w"
    # Calculate number of patches
    L = (H // patch h) * (W // patch w)
    # Extract patches
    patches = imgs.unfold(2, patch h, patch h).unfold(3, patch w, patch
    # Calculate D
    D = patch h * patch w * C
    # Reshape patches to (N, L, D)
    patches = patches.reshape(N, L, D)
    return patches
def configure optimizers(self):
    ### TODO: configure the optimiser to be Adam with learning rate 1e-4
    ### ADD YOUR CODE HERE
    return torch.optim.Adam(self.parameters(), lr=1e-4)
def unpatchify(self, x):
    0.00
    x: (N, L, D)
    imgs: (N, C, H, W)
    ### TODO: Use the previously defined function
    ### ADD YOUR CODE HERE
    N, L, \underline{\phantom{a}} = x.shape
    number of channels = self.in chans
    patch h, patch w = self.patch embed.patch size
    img h, img w = self.patch embed.img size
    # Check if L matches the expected number of patches
```

```
assert L == (img_h // patch_h) * (img_w // patch_w), "Number of patc
    # Reshape patches to (N, H // patch h, W // patch w, patch h, patch
    patches = x.view(N, img h // patch h, img w // patch w, patch h, pat
    # Permute and reshape to get the final images (N, C, H, W)
    imgs = patches.permute(0, 5, 1, 3, 2, 4).contiguous().view(N, number
    return imas
def random masking(self, x, mask ratio):
    #def random masking(patches, mask ratio):
    0.00
    Perform per-sample random masking by per-sample shuffling.
    Per-sample shuffling is done by argsort random noise.
    x: [N, L, D], sequence
    ### TODO: Use the previously defined function
    ### ADD YOUR CODE HERE
    x = x.to(self.pos embed.device)
    N, L, D = x.shape # batch, length, dim
    # Step 1: create noise in [0, 1]
    noise = torch.rand(N, L).to(x.device)
    # Step 2: sort noise for each sample
    noise, indices = noise.sort(dim=1)
    # Step 3: store list of indices to revert shuffling operation later
    ids restore = torch.argsort(indices, dim=1)
    # Step 4: used shuffled list to keep only a subset of patches
    # Calculate the number of patches to keep
    L \text{ kept} = \text{int}(L * (1 - \text{mask ratio}))
    # Select the indices of the patches to keep (the first L kept after
    indices keep = indices[:, :L kept]
    # Extract the patches to keep using gather
    patches kept = torch.gather(x, dim=1, index=indices keep.unsqueeze(-
    # Step 5 : generate the binary mask
    # First, in the mixed order: the first L kept are 0 (unmasked) and t
    mask shuffled = torch.ones(N, L, device=x.device)
    mask shuffled[:, :L kept] = 0
    # Revert the permutation so that the mask corresponds to the original
    mask = torch.gather(mask shuffled, dim=1, index=ids restore)
    return patches kept, mask, ids restore
def forward encoder(self, x, mask ratio):
```

```
Forward function for the encoding part.
    # embed patches (use self.patch embed)
    x = self.patch embed(x)
    # add pos embed w/o cls token
    x = x + self.pos embed[:, 1:, :]
    # masking: length -> length * mask ratio
    x, mask, ids restore = self.random masking(x, mask ratio)
    # append cls token
    cls token = self.cls token + self.pos embed[:, :1, :]
    cls tokens = cls token.expand(x.shape[0], -1, -1)
    x = torch.cat((cls tokens, x), dim=1)
    # apply Transformer blocks
    for blk in self.blocks:
        x = blk(x)
    x = self.norm(x)
    return x, mask, ids restore
def forward decoder(self, x, ids restore):
    Forward function for the decoding part.
    # embed tokens
    x = self.decoder embed(x)
    # append mask tokens to sequence
    mask tokens = self.mask token.repeat(
        x.shape[0], ids restore.shape[1] + 1 - x.shape[1], 1
    x = torch.cat([x[:, 1:, :], mask tokens], dim=1) # no cls token
    x = torch.gather(
       x , dim=1, index=ids restore.unsqueeze(-1).repeat(1, 1, x.shape[
    ) # unshuffle
    x = torch.cat([x[:, :1, :], x], dim=1) # append cls token
    # add pos embed
    x = x + self.decoder pos embed
    # apply Transformer blocks
    for blk in self.decoder blocks:
        x = blk(x)
    x = self.decoder norm(x)
    # predictor projection
    x = self.decoder pred(x)
    # remove cls token
    x = x[:, 1:, :]
    return x
```

```
def compute loss(self, target patches, pred patches, mask):
    This function returns the MAE loss value for a given batch.
    Should be MSE loss over masked patches
    Aras:
      target patches: [N, L, D] ground truth patches
      pred patches: [N, L, D] predicted patches
      mask: [N, L] binary mask indicating which patches are masked
    ### TODO
    ### ADD YOUR CODE HERE
    mask = mask.unsqueeze(2) # (N, L, 1) Add third dimension to mask
    masked pred patches = pred patches * mask # Masked predicted patche
    masked target patches = target patches * mask # Masked target patch
    loss = F.mse loss(masked pred patches, masked target patches) # Loss
    return loss
def forward(self, imgs, mask ratio=0.75):
    Forward function
    Aras:
      imgs: batch of [N, C, H, W] images
      mask ratio: masking ratio to use for the encoder
    Returns:
      predicted patches [N, L, D], where D = patch size[0]*patch size[1]
     mask [N, L]
    0.00
    ### TODO
    ### ADD YOUR CODE HERE
    # Forward pass through the encoder
    x, mask, ids restore = self.forward encoder(imgs, mask ratio)
    # Forward pass through the decoder
    predicted patches = self.forward decoder(x, ids restore)
    return predicted patches, mask
def training step(self, batch, batch idx):
    images = batch[0]
    predicted patches, mask = self(images)
    target patches = self.patchify(images)
    loss = self.compute loss(target patches, predicted patches, mask)
    self.log('loss train', loss, prog bar=True)
    if batch idx == 0:
        images output = self.unpatchify(predicted patches * mask.unsquee
        grid = torchvision.utils.make grid(images[0:4], nrow=4, normaliz
        self.logger.experiment.add image('train images input', grid, sel
        grid = torchvision.utils.make grid(images output[0:4], nrow=4, r
        self.logger.experiment.add image('train images output', grid, se
        grid = torchvision.utils.make grid(self.unpatchify(target patche
        self.logger.experiment.add image('train patches target', grid, s
```

```
grid predicted = torchvision.utils.make grid(self.unpatchify(pre
        self.logger.experiment.add image('train patches predicted', grid
    return loss
def validation step(self, batch, batch idx):
    images = batch[0]
    predicted patches, mask = self(batch[0])
    target patches = self.patchify(images)
    loss = self.compute loss(target patches, predicted patches, mask)
    self.log('loss val', loss, prog bar=True)
def get class embeddings(self, images):
    Return the class embeddings extracted from the encoder
    for each image in the batch.
   This function is meant to be used at inference, we do not mask
    any patches.
    embeddings, , = self.forward encoder(images, mask ratio=0)
    return embeddings[:, 0, :]
def predict step(self, batch, batch idx):
    images, labels = batch[0], batch[1]
    return {'embeddings': self.get class embeddings(images), 'labels': l
```

Next, we define a tiny toy VIT architecture for you to use in this coursework. This is much smaller than standard VIT architectures but will allow you to train your MAE rapidly on a single GPU. Note that we use again a patch size of 4 given the small resolution of the input images.

```
In [44]: def mae_vit_toy_patch4_dec256d4b():
    """

    Creates a toy ViT with patch size 4.
    """

    model = MaskedAutoencoderViT(
        in_chans=1,
        img_size=28,
        patch_size=4,
        embed_dim=384,
        depth=6,
        num_heads=6,
        decoder_embed_dim=256,
        decoder_depth=4,
        decoder_num_heads=8,
        mlp_ratio=4,
    )
    return model
```

#### Task C-2: MAE training

#### Tensorboard logging

Load tensorboard, we should be able to monitor training and validation loss as well as your reconstructed training images.

```
In [45]: %reload_ext tensorboard
%tensorboard --logdir './lightning_logs/coursework/'
```

# **Index of** /

Name	Size	<b>Date Modified</b>			
bin/		5/9/25, 8:14:21 PM			
bin.usr-is-merged/		2/26/24, 12:58:31 PM			
boot/		5/7/25, 9:13:06 AM			
cdrom/		8/27/24, 5:22:46 PM			
dev/		5/10/25, 2:34:25 PM			
etc/		5/9/25, 8:14:41 PM			
home/		12/28/24, 4:03:13 PM			
lib/		5/9/25, 8:14:19 PM			
lib.usr-is-merged/		4/8/24, 3:37:57 PM			
lib32/		2/9/25, 10:15:30 AM			
lib64/		2/9/25, 10:15:30 AM			
lost+found/		12/28/24, 3:57:36 PM			
media/		4/16/25, 8:00:02 PM			
mnt/		8/27/24, 4:37:13 PM			
opt/		4/11/25, 7:37:58 AM			
proc/		5/10/25, 2:34:01 PM			
root/		4/16/25, 8:46:19 AM			
run/		5/10/25, 3:00:43 PM			
sbin/		5/7/25, 9:12:08 AM			
sbin.usr-is-merged/		3/31/24, 10:00:13 AM			
snap/		5/5/25, 9:08:54 PM			
srv/		8/27/24, 4:37:13 PM			
sys/		5/10/25, 2:34:01 PM			
tmp/		5/10/25, 5:14:21 PM			
usr/		1/17/25, 10:24:56 AM			
var/		1/17/25, 4:45:34 PM			
swap.img	4.0 GB	4/7/25, 6:43:36 AM			

```
In [46]: seed everything(33, workers=True)
         data = PneumoniaMNISTDataModule(batch size=32)
         device = torch.device("cuda" if torch.cuda.is available() else "cpu")
         print(device)
         model = mae vit toy patch4 dec256d4b()
         trainer = Trainer(
            max epochs=60, # Increased a bit because of the jump at the end in the l
            accelerator='auto',
            devices=1,
            logger=TensorBoardLogger(save dir='./lightning logs/coursework/', name='
         trainer.fit(model=model, datamodule=data)
        Seed set to 33
       GPU available: True (cuda), used: True
       TPU available: False, using: 0 TPU cores
       HPU available: False, using: 0 HPUs
       LOCAL RANK: 0 - CUDA VISIBLE DEVICES: [0]
                                  | Params | Mode
         l Name
                          | Type
       0 | patch_embed | PatchEmbed | 6.5 K | train
1 | blocks | ModuleList | 10.6 M | train
       2 | norm | LayerNorm | 768 | train
       3 | decoder_embed | Linear | 98.6 K | train
       4 | decoder blocks | ModuleList | 3.2 M | train
        5 | decoder_norm | LayerNorm | 512 | train
       6 | decoder pred | Linear | 4.1 K | train
         | other params | n/a | 32.6 K | n/a
        _____
       13.9 M
                Trainable params
       32.0 K Non-trainable params
       13.9 M Total params55.796 Total estimated model params size (MB)
       219
                 Modules in train mode
               Modules in eval mode
       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
       cuda
        (7, 7)
       Sanity Checking: | 0/? [00:00<?, ?it/s]
```

/home/eder/miniconda3/envs/mlimaging/lib/python3.10/site-packages/pytorch\_lightning/trainer/connectors/data\_connector.py:425: 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=11` in the `DataLoader` to improve performance.

/home/eder/miniconda3/envs/mlimaging/lib/python3.10/site-packages/pytorch\_lightning/trainer/connectors/data\_connector.py:425: The 'train\_dataloader' does not have many workers which may be a bottleneck. Consider increasing the value of the `num\_workers` argument` to `num\_workers=11` in the `DataLoader` to improve performance.

```
| 0/? [00:00<?, ?it/s]
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Validation: |
`Trainer.fit` stopped: `max epochs=60` reached.
```

# Part D: Inspect the trained model.

In this last part, we analyse the feature embeddings (or representations) obtained from your trained model with t-SNE. Let's see if our model learned anything useful!

# **Task D-1:** Inspect and compare the learned feature representations of your trained model.

We compare the feature embeddings of the trained model to embeddings obtained with a randomly initialised (untrained) model. We create some scatter plot visualisations and describe the findings with a few sentences.

```
In [47]: from sklearn.manifold import TSNE
   import seaborn as sns
   from sklearn import decomposition
   import pandas as pd
```

Let's get the representations from our trained model:

```
In [48]:
    class MaskedAutoencoderViTEmbeddings(MaskedAutoencoderViT):
        def __init__(
            self,
            img_size=224,
            patch_size=16,
            in_chans=3,
            embed_dim=1024,
            depth=24,
            num_heads=16,
            decoder_embed_dim=512,
            decoder_depth=8,
            decoder_num_heads=16,
            mlp_ratio=4.0,
        ):
            super().__init__(img_size, patch_size, in_chans, embed_dim, depth, results)
```

```
self.embeddings = [] # list where we still store the embeddings
             def get embedding(self, x, mask ratio=0.75):
                 x, , = self.forward encoder(x, mask ratio)
                 return x.view(x.size(0), -1)
             def on test start(self):
                 self.embeddings = [] # clear the list of embeddings at the start of
             def test step(self, batch, batch idx):
                 imgs, = batch
                 emb = self.get embedding(imgs)
                 self.embeddings.append(emb)
In [50]: model dir = './lightning logs/coursework/mae test/version 0/checkpoints/epoc
         model modified = MaskedAutoencoderViTEmbeddings load from checkpoint(model c
                                                                               decoder
         trainer.test(model=model modified, datamodule=data)
         embeddings = torch.cat(model modified.embeddings, dim=0).cpu().numpy()
         print(embeddings.shape)
        LOCAL RANK: 0 - CUDA VISIBLE DEVICES: [0]
        (7, 7)
        /home/eder/miniconda3/envs/mlimaging/lib/python3.10/site-packages/pytorch li
        ghtning/trainer/connectors/data connector.py:425: The 'test dataloader' does
        not have many workers which may be a bottleneck. Consider increasing the val
        ue of the `num workers` argument` to `num workers=11` in the `DataLoader` to
        improve performance.
        Testing: |
                            | 0/? [00:00<?, ?it/s]
        (624, 4992)
In [51]: # Create a dataframe with the class labels
         labels = np.array([data.test set[i][1] for i in range(0,len(data.test set))]
         df = pd.DataFrame(labels, columns=['class label'])
         # Perform PCA on the embeddings
         pca = decomposition.PCA(n components=0.95, whiten=False)
         embeddings pca = pca.fit transform(embeddings)
         print("Embedding shape after PCA: ", embeddings pca.shape)
         # Add the PCA components to the dataframe
         df['features - PCA 1'] = embeddings pca[:,0]
         df['features - PCA 2'] = embeddings pca[:,1]
         df['features - PCA 3'] = embeddings pca[:,2]
         df['features - PCA 4'] = embeddings pca[:,3]
         # Perform t-SNE on the embeddings
         embeddings tsne = TSNE(n components=2, init='random', learning rate='auto').
         print("Embedding shape after PCA and TSNE: ", embeddings tsne.shape)
         # Add the t-SNE components to the dataframe
         df['features - t-SNE 1'] = embeddings tsne[:,0]
         df['features - t-SNE 2'] = embeddings tsne[:,1]
```

```
df.head() # showing the first five entries in the dataframe
```

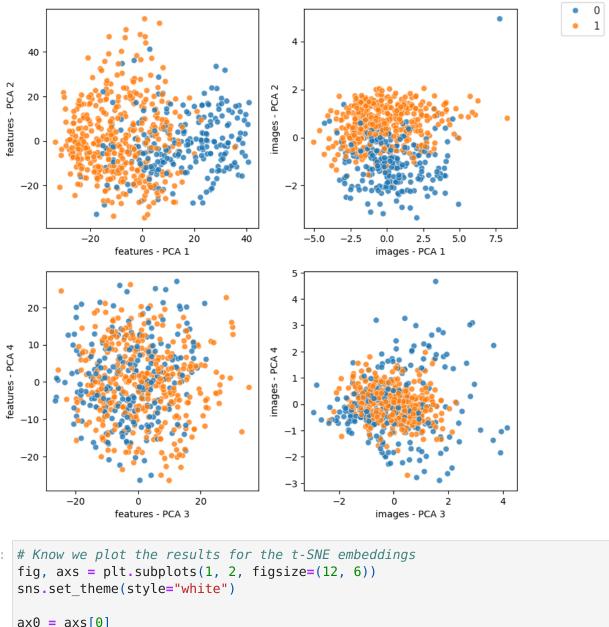
Embedding shape after PCA: (624, 201) Embedding shape after PCA and TSNE: (624, 2)

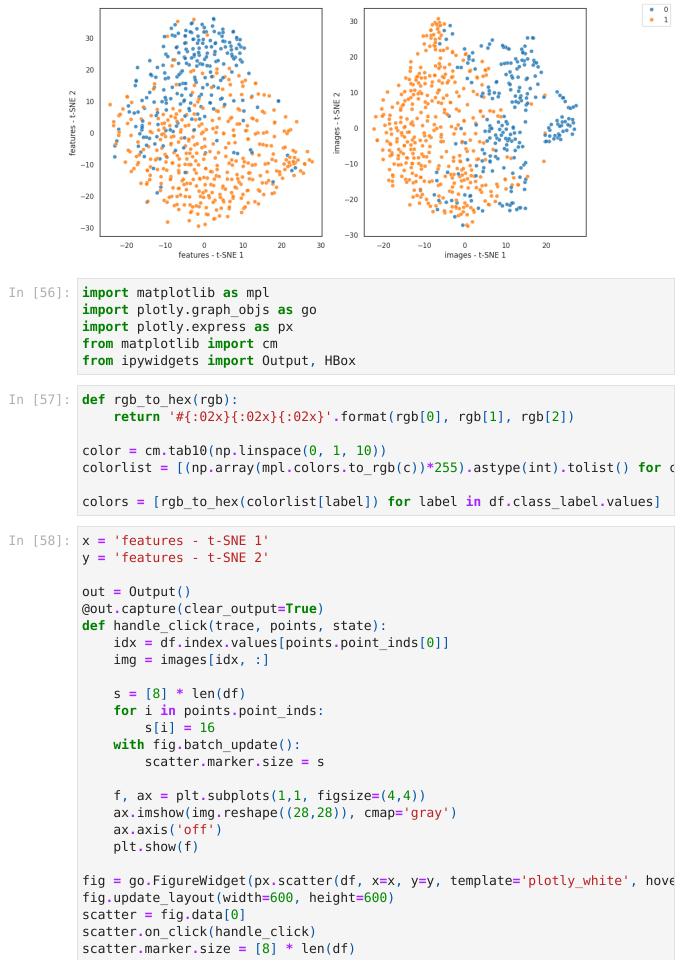
Out[51]: features features features features features features class label PCA 1 PCA 2 PCA 3 PCA 4 t-SNE 1 t-SNE 2 0 3.908165 6.035489 4.070599 -1.341054 6.714702 4.483046 1 0 -13.897316 -11.915040 -12.110461 21.332968 -12.071618 -18.935572 1 2 1 7.879046 -15.328770 -3.921061 12.499488 -10.515244 5.817060 3 28.587406 -2.637968 1.720091 -4.516327 2.685032 24.806839 20.228897 4 1 -16.083618 -1.692487 

```
In [52]: # Convert the images into a numpy array and reshape them
         images = np.array([data.test set[i][0] for i in range(0,len(data.test_set))]
         images = images.reshape(images.shape[0], -1) # linearize the 28x28 Pneumonia
         print(images.shape)
         # Perform PCA on the images
         pca = decomposition.PCA(n components=0.95, whiten=False)
         images pca = pca.fit transform(images)
         # Add the PCA components to the dataframe
         df['images - PCA 1'] = images pca[:,0]
         df['images - PCA 2'] = images pca[:,1]
         df['images - PCA 3'] = images pca[:,2]
         df['images - PCA 4'] = images pca[:,3]
         print("Image shape after PCA: ", images pca.shape)
         # Perform t-SNE on the images
         images tsne = TSNE(n components=2, init='random', learning rate='auto').fit
         # Add the t-SNE components to the dataframe
         df['images - t-SNE 1'] = images tsne[:,0]
         df['images - t-SNE 2'] = images_tsne[:,1]
         print("Image shape after PCA and TSNE: ", images tsne.shape)
         df.head() # showing the first five entries in the dataframe
```

```
(624, 784)
Image shape after PCA: (624, 64)
Image shape after PCA and TSNE: (624, 2)
```

```
Out[52]:
                        features -
                                  features -
                                             features -
                                                       features -
                                                                  features -
                                                                            features -
                                                                                       im.
            class label
                           PCA 1
                                      PCA 2
                                                PCA 3
                                                          PCA 4
                                                                    t-SNE 1
                                                                              t-SNE 2
                         3.908165
                                   6.035489
                                              4.070599
          0
                    1
                                                       -1.341054
                                                                  6.714702
                                                                             4.483046
                                                                                      1.1
          1
                    0 -13.897316 -11.915040 -12.110461 21.332968 -12.071618 -18.935572 -1.34
                         7.879046 -15.328770
          2
                                             -3.921061 12.499488 -10.515244
                                                                             5.817060 -1.20
                       28.587406
                                  -2.637968
                                              1.720091
                                                       -4.516327
                                                                  2.685032
                                                                            24.806839
                                                                                       0.8
          3
          4
                    1 -16.083618
                                  -1.692487
                                             20.228897 4.549816
                                                                10.174796 -23.838036 -0.6
In [53]: # Define the plotting parameters
         alpha = 0.8
          style = 'o'
         markersize = 40
          color palette = 'tab10'
          kind = 'scatter'
In [54]: fig, axs = plt.subplots(2, 2, figsize=(8, 8))
          sns.set theme(style="white")
          # First plot: we leave the legend active to extract it later
          ax0 = axs[0, 0]
          sns.scatterplot(ax=axs[0, 0], data=df, x='features - PCA 1', y='features - F
                          hue='class label', alpha=alpha, marker=style, s=markersize,
                          palette=color palette)
          # Extract the handles and labels
          handles, labels = ax0.get legend handles labels()
          # Remove the legend from the first subplot
          if ax0.get legend() is not None:
              ax0.get legend().remove()
          # Second plot
          sns.scatterplot(ax=axs[0, 1], data=df, x='images - PCA 1', y='images - PCA 2
                          hue='class label', alpha=alpha, marker=style, s=markersize,
                          palette=color palette, legend=False)
          # Third plot
          sns.scatterplot(ax=axs[1, 0], data=df, x='features - PCA 3', y='features - F
                          hue='class label', alpha=alpha, marker=style, s=markersize,
                          palette=color palette, legend=False)
          # Fourth plot
          sns.scatterplot(ax=axs[1, 1], data=df, x='images - PCA 3', y='images - PCA 4
                          hue='class label', alpha=alpha, marker=style, s=markersize,
                          palette=color palette, legend=False)
          # Create the global legend
          fig.legend(handles, labels, loc='upper right', bbox to anchor=(1.15, 1))
          plt.tight layout()
          plt.show()
```





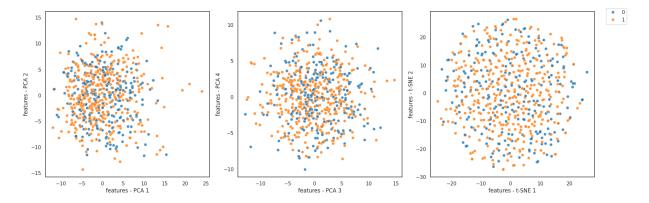
```
scatter.marker.color = colors
         HBox([fig, out])
Out[58]: HBox(children=(FigureWidget({
              'data': [{'customdata': array([[1],
         Let's compare with the representation of an untrained model
In [59]: # Create the untrained model
         untrained model = MaskedAutoencoderViTEmbeddings(in chans=1, img size=28, pa
                                                                               decoder
         trainer.test(model=untrained model, datamodule=data)
         embeddings = torch.cat(untrained model.embeddings, dim=0).cpu().numpy()
         print(embeddings.shape)
        LOCAL RANK: 0 - CUDA VISIBLE DEVICES: [0]
        (7, 7)
        /home/eder/miniconda3/envs/mlimaging/lib/python3.10/site-packages/pytorch li
        ghtning/trainer/connectors/data connector.py:425: PossibleUserWarning:
        The 'test dataloader' does not have many workers which may be a bottleneck.
        Consider increasing the value of the `num workers` argument` to `num workers
        =11` in the `DataLoader` to improve performance.
                            | 0/? [00:00<?, ?it/s]
        Testing: |
        (624, 4992)
In [60]: # Create a dataframe with the class labels for the untrained model
         labels = np.array([data.test set[i][1] for i in range(0,len(data.test set))]
         df = pd.DataFrame(labels, columns=['class label'])
         # Perform PCA on the embeddings
         pca = decomposition.PCA(n components=0.95, whiten=False)
         embeddings pca = pca.fit transform(embeddings)
         print(embeddings pca.shape)
         # Add the PCA components to the dataframe
         df['features - PCA 1'] = embeddings pca[:,0]
         df['features - PCA 2'] = embeddings pca[:,1]
         df['features - PCA 3'] = embeddings pca[:,2]
         df['features - PCA 4'] = embeddings pca[:,3]
         # Perform t-SNE on the embeddings
         embeddings tsne = TSNE(n components=2, init='random', learning rate='auto').
         print(embeddings tsne.shape)
         # Add the t-SNE components to the dataframe
         df['features - t-SNE 1'] = embeddings tsne[:,0]
         df['features - t-SNE 2'] = embeddings tsne[:,1]
```

```
df.head() # showing the first five entries in the dataframe
(624, 62)
(624, 2)
```

Out[60]:

:	class_label	features - PCA 1	features - PCA 2	features - PCA 3	features - PCA 4	features - t-SNE 1	features - t-SNE 2
(	1	0.085608	-2.314194	3.707670	-2.359124	1.803094	-1.547044
	0	0.716355	5.428017	3.517769	4.926601	-5.493159	13.958353
2	2 1	-1.473587	4.459114	3.500272	-0.861529	-18.851530	4.424058
3	0	0.858930	-8.287944	-3.508290	4.701985	10.046527	-13.687622
4	<b>,</b> 1	-1.767120	-5.695096	-8.738058	3.342774	6.433622	-9.634429

```
In [61]: fig, axs = plt.subplots(1, 3, figsize=(18, 6))
         sns.set theme(style="white")
         # Firts plot
         ax0 = axs[0]
         sns.scatterplot(ax=ax0, data=df, x='features - PCA 1', y='features - PCA 2',
                         hue='class_label', alpha=alpha, marker=style, s=markersize,
                         palette=color palette)
         handles, labels = ax0.get legend handles labels()
         # Remove the legend from the first subplot
         if ax0.get legend() is not None:
             ax0.get legend().remove()
         # Sedond plot
         sns.scatterplot(ax=axs[1], data=df, x='features - PCA 3', y='features - PCA
                         hue='class label', alpha=alpha, marker=style, s=markersize,
                         palette=color palette, legend=False)
         # Third plot
         sns.scatterplot(ax=axs[2], data=df, x='features - t-SNE 1', y='features - t-
                         hue='class label', alpha=alpha, marker=style, s=markersize,
                         palette=color palette, legend=False)
         # General legend
         fig.legend(handles, labels, loc='upper right', bbox to anchor=(1.05, 1))
         plt.tight layout()
         plt.show()
```



#### **Analysis**

The scatter plots of the first two PCA components reveal a significantly better separation of class labels compared to the third and fourth components. A similar trend appears when PCA is applied directly to the raw images—the first two dimensions yield much clearer class separation, suggesting that these dimensions are the most informative.

Furthermore, when t-SNE is applied to the PCA outputs, the visual discrimination of the labels is further enhanced. In the case of the embeddings, t-SNE produces a more distinct separation between the two groups, although some misclassified instances still persist. A comparable improvement is evident when t-SNE is applied to the PCA-reduced images, with the classes becoming more distinguishable even though a few samples remain ambiguous. This outcome is intuitive since PCA alone may discard certain dimensions that contain relevant information, whereas t-SNE, by capturing the nonlinear structure in the data, offers a more holistic visualization.

In stark contrast, the visualizations derived from the embeddings of an untrained model—both via PCA and t-SNE—show the classes to be completely intermixed. This stark difference underscores the critical role of training in enabling the model to learn discriminative features.

The interactive visualizations also reveal that while some misclassifications are understandable due to the inherent ambiguity of certain samples, others are less intuitive. Although the trained model exhibits a marked improvement over its untrained counterpart, the presence of misclassified instances in the t-SNE plots indicates that there is still room for refinement.

Rather than simply increasing the number of epochs—which would probably further reduce the training loss as we see in the validation plot—a better improvement could involve incorporating a contrastive learning objective into the training process. By integrating a contrastive loss (or even a triplet loss), the model would be encouraged to pull together similar samples while pushing apart dissimilar ones in the latent space, potentially resulting in a more robust and discriminative feature representation. This approach could lead to a more finely tuned model that better separates the classes in its embeddings.