Computer Vision & Deep Learning - Generative AI & Visual Synthesis

Exercise 4: Convolutional Neural Networks

Due on 24.05.2024., 10:00

Important notes

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- Moodle: Please use the Moodle platform and post your questions to the forum. They will be answered by us or your fellow students.
- Submission: Put your code and potentially other materials inside a single ZIP file. If you use jupyter notebooks, please always create a PDF file and include it in your ZIP file. The final submission should therefore be a single zip file with a PDF of your code and the original code inside. The ZIP file should contain your surname and your matriculation number (Surname-MatriculationNumber.zip). Submissions that fail to follow the naming convention will not be graded!

```
import os
import numpy as np
from PIL import Image
from tgdm import tgdm
import matplotlib.pyplot as plt
import torch
import torchvision
import torch.nn as nn
import torch.optim as optim
import torch.nn.functional as F
from torchvision import transforms as T
# fix size of images for matplot
plt.rcParams['figure.figsize'] = [10, 5]
RESULTS DIR = 'results'
# Create results directory
if not os.path.exists(RESULTS DIR):
    os.makedirs(RESULTS_DIR)
```

Task 1: Convolutions Revisited

Compute the output of a 2D-convolution operation with 3 different filters of kernel size 2 ×2 defined by the following weight matrices and biases

```
# Define the input tensor
x = torch.tensor([[1., 0., 0., 1., 1., 0.],
                   [0., 1., 0., 0., 0., 1.],
                   [0., 0., 1., 1., 0., 0.]
                   [1., 0., 0., 0., 1., 0.],
                   [1., 0., 0., 0., 0., 1.]]
# Define the filters
w1 = torch.tensor([[1., 0.],
                    [0., 1.]]
w2 = torch.tensor([[1., 0.],
                    [1., 0.]]
w3 = torch.tensor([[1., 1.]],
                    [0., 0.]
# Define the biases
b1 = torch.tensor([-1])
b2 = torch.tensor([-1])
b3 = torch.tensor([-1])
# Perform the convolution operation for each filter
conv w1 = F.conv2d(x.unsqueeze(0).unsqueeze(1),
w1.unsqueeze(0).unsqueeze(0), stride=1, padding=0)
conv w2 = F.conv2d(x.unsqueeze(0).unsqueeze(1),
w2.unsqueeze(0).unsqueeze(0), stride=1, padding=0)
conv w3 = F.conv2d(x.unsqueeze(0).unsqueeze(1),
w3.unsqueeze(0).unsqueeze(0), stride=1, padding=0)
# Subtract the bias and apply the ReLU activation function
output w1 = F.relu(conv w1 - b1)
output w2 = F.relu(conv w2 - b2)
output w3 = F.relu(conv w3 - b3)
# Print the output for each filter
print(output w1.squeeze())
print(output w2.squeeze())
print(output_w3.squeeze())
tensor([[3., 1., 1., 2., 3.],
        [1., 3., 2., 1., 1.],
        [1., 1., 2., 3., 1.],
        [2., 1., 1., 1., 3.]])
tensor([[2., 2., 1., 2., 2.],
        [1., 2., 2., 2., 1.],
        [2., 1., 2., 2., 2.],
```

Manual Calculation

1. Using s = 1 (stride) and p = 0 (padding), calculate by hand the output of the convolution operation followed by the ReLU activation function.

```
x = [1., 0., 0., 1., 1., 0.], [0., 1., 0., 0., 0., 1.], [0., 0., 1., 1., 0., 0.], [1., 0., 0., 0., 1., 0.], [1., 0., 0., 0., 0., 0., 0.]
w1 = [1., 0.], [0., 1.]
w2 = [1., 0.], [1., 0.]
w3 = [1., 1.], [0., 0.]
b1 = [-1]
b2 = [-1]
```

(1) For Filter 1

Step-by-step

Filter RELU(w1 * x - b1):

- 1. 1*1+0*0+0*1+1*1=2subtract bias b1=2-1=1ReLU(1)=1
- 2. 1*0+0*1+0*0+1*0=0subtract bias b1=0-1=-1ReLU(-1)=0
- 3. 1*0+0*0+0*1+1*1=1subtract bias b1 = 1-1=0ReLU(0) = 0

4.
$$1*1+0*0+0*1+1*0=1$$

subtract bias b1 = 1 - 1 = 0

$$ReLU(0) = 0$$

5.
$$1*1+0*0+0*0+1*1=2$$

subtract bias b1 = 2 - 1 = 1

$$ReLU(1) = 1$$

6.
$$1*0+0*1+0*0+1*0=0$$

subtract bias b1 = 0 - 1 = -1

$$ReLU(-1) = 0$$

Result

convolve filter with image -- w1 * x

subtract bias b1 -- output of q1 - b1

$$a1 = [0., -1., 0., 0., -1.], [-1., 0., -1., -1., 0.], [-1., -1., 0., -1., -1.], [0., -1., -1., 0., -1.], [0., -1., -1., 0.]$$

activate feature map -- output of RELU(a1)

$$z1 = [0., 0., 0., 0., 0.], [0., 0., 0., 0., 0.], [0., 0., 0., 0.], [0., 0., 0., 0., 0., 0.], [0., 0., 0., 0.]$$

(2) For Filter 2 Filter RELU(w2 * x - b2)

Result

convolve filter with image -- w3 * x q2 = [1, 0, 1, 0, 0] [0, 1, 0, 0, 0] [0, 0, 1, 1, 0] [1, 0, 0, 0, 0] [1, 0, 0, 0, 0]

substract bias b1 -- output of q3 - b3 a2 = [0, -1, 0, -1, -1] [-1, 0, -1, -1, -1] [-1, -1, 0, 0, -1] [0, -1, -1, -1, -1] [0, -1, -1, -1, -1]

activate feature map -- output of RELU(a3) z2 = [0, 0, 0, 0, 0] [0, 0, 0, 0, 0] [0, 0, 0, 0, 0] [0, 0, 0, 0, 0]

(3) For Filter 3 Filter RELU(w3 * x - b3)

Result convolve filter with image -- w3 * x q3 = [1, 1, 2, 1, 1] [1, 1, 0, 0, 1] [0, 0, 1, 1, 0] [1, 1, 0, 0, 1] [1, 1, 0, 0, 1]

substract bias b1 -- output of q3 - b3 a3 = [0, 0, 1, 0, 0] [0, 0, -1, -1, 0] [-1, -1, 0, 0, -1] [0, 0, -1, -1, 0] [0, 0, -1, -1, 0]

activate feature map -- output of RELU(a3) z3 = [0, 0, 1, 0, 0] [0, 0, 0, 0, 0] [0, 0, 0, 0, 0] [0, 0, 0, 0, 0]

1. Think about and describe how the input image patch, the convolution kernel, and the ReLU activation function interact, and how you can avoid manual computation and directly infer the output for each patch

The fed throuh image patch is a small, localized region of the overall image which represents features of a neighbouring area or local patterns typical in rasterized image data. A convolution operation involves sliding the kernel (a small matrix) over an image patch, element-wise multiplying the kernel's values with the corresponding values in the patch, and summing up the results.

Other options

- (1) Kernel Flipping: In the mathematical definition of convolution, the kernel is actually flipped both horizontally and vertically before it is slid over the input patch.
- (2) Convolution in the Fourier Domain: The convolution operation can also be performed in the frequency domain, rather than the spatial domain. According to the convolution theorem, the convolution of two signals is equivalent to the point-wise multiplication of their Fourier transforms. This can be more efficient than performing the convolution in the spatial domain
 - 1. Now repeat (a) with s = 2. Please only report the final outputs, not the intermediate ones before the activation.

```
z1 = [0, 0, 0] [0, 0, 0]

z2 = [0, 0, 0] [0, 0, 0]

z3 = [0, 1, 0] [0, 0, 0]
```

```
# Perform the convolution operation for each filter
conv w1 = F.conv2d(x.unsqueeze(0).unsqueeze(1),
w1.unsqueeze(0).unsqueeze(0), stride=2, padding=0)
conv w2 = F.conv2d(x.unsqueeze(\frac{0}{2}).unsqueeze(\frac{1}{2}),
w2.unsqueeze(0).unsqueeze(0), stride=2, padding=0)
conv w3 = F.conv2d(x.unsqueeze(0).unsqueeze(1),
w3.unsqueeze(0).unsqueeze(0), stride=2, padding=0)
# Subtract the bias and apply the ReLU activation function
output_w1 = F.relu(conv_w1 - b1)
output_w2 = F.relu(conv_w2 - b2)
output w3 = F.relu(conv w3 - b3)
# Print the output for each filter
print(output w1.squeeze())
print(output w2.squeeze())
print(output_w3.squeeze())
tensor([[3., 1., 3.],
        [1., 2., 1.]
tensor([[2., 1., 2.],
        [2., 2., 2.]])
```

```
tensor([[2., 2., 2.], [1., 3., 1.]])
```

1. What might be the purpose of using s = 2? What other operation can be used instead for the same purpose? Hint: Think about compression.

This is a form of downsampling, also known as subsampling or pooling. The purpose of downsampling is to reduce the spatial size of the input image or feature map, which can help to reduce the number of parameters and computations in the network, and to make the network more invariant to small translations of the input.

Further a larger stride can increase the receptive field of the neurons in the subsequent layers

1. Please derive the general formula to compute the output size of a convolution operation (size of the feature map), depending on the kernel size, the stride, and padding. It is sufficient to derive it for the 1D case.

$$Out = [(W - K + 2P)/S) + 1$$

- W is the input volume $(W_h \times W_w)$
- K is the Kernel size $(K_h \times K_w)$
- P is the padding
- S is the stride

Task 2: Receptive Field

The receptive field of a neuron in a Convolutional Neural Network (CNN) is the size of the image region that is connected to the neuron

- 1. Suppose that we have a CNN with 3 convolutional layers with a kernel size of 3 × 3 and s = 1. Calculate the receptive field of a neuron after the last convolutional layer.
- Conv_Layer 0: $RF = 3 \times 3 -->$ since the input is the original image 3x3
- Conv_Layer 1: RF = (3-1)*1+3=5
- Conv_Layer 2: RF = (5-1)x1+3x3=7

The receptive field is 7×7

- 1. Suppose that we have a CNN with 2 convolutional layers with a kernel size of 4 × 4 and s = 2. Calculate the receptive field of a neuron after the last convolutional layer.
- Conv_Layer 0: $RF = 4 \times 4 -->$ since the input is the original image 4x4
- Conv_Layer 1: RF = (4-1)x2+4x4=10

The receptive field is 10×10

Based on your findings, how is the depth related to the receptive field of a CNN?

The depth of a CNN, or the number of convolutional layers, is directly related to the receptive field of a neuron in the network. As previously observable, the receptive field of a neuron

increases with each convolutional layer, as the neuron's input is a function of the outputs of the previous layer's neurons.

1. When keeping the depth fixed, can you infer any strategies to still increase the receptive field of a CNN?

Increasing the receptive field by simply adding more convolutional layers is only possible to a limited extent. Further, the receptive field grows only slowly from the first to the last layer which might not be desirable if contextual information can be beneficial.

Therefore, one strategy is dilation, where gaps are inserted between the elements of the convolutional kernel. This can effectively increase the receptive field of a neuron without increasing the number of parameters or computations in the network.

Another strategy is the use of stride in the convolutional layers, which can increase the receptive field of a neuron while skipping one or more elements of the input at a time.

In addition, skip connections, where one or more layers in the network are bypassed and the input is connected directly to the output, can also be used to increase the receptive field of a CNN. Skip connections can allow neurons in subsequent layers to access the features and information from the earlier layers and can help alleviate the vanishing gradient problem.

Task 3: Convolutional Neural Networks

In the previous exercise, we already used the PyTorch framework to train a neural network classifier. Now, we want to move on to a more complex dataset than MNIST and use a Convolutional Neural Network for classification.

Task 3.1: Dataset

```
# Load CIFAR10 dataset
dataset = torchvision.datasets.CIFAR10(root='data/', download=True)
# Split the dataset into training and testing sets
train size = int(0.8 * len(dataset))
test size = len(dataset) - train size
train dataset, test dataset = torch.utils.data.random split(dataset,
[train size, test size])
Files already downloaded and verified
# Plot 10 images from each class
fig, axs = plt.subplots(2, 5, figsize=(15, 6))
for i in range(10):
    img, label = dataset[i]
    ax = axs[i // 5, i % 5]
    ax.imshow(img)
    ax.set title(dataset.classes[label])
    ax.axis('off')
```

```
# set title
fig.suptitle('CIFAR10 dataset')

# save the plot
plt.savefig(f'{RESULTS_DIR}/cifar10.png')
```



```
# define data transformations
transform = T.Compose([
    T.Resize((32,32)),
    T.ToTensor(),
    #T.Normalize(mean=[0.5, 0.5, 0.5], std=[0.5, 0.5, 0.5])
1)
# apply data transformations to the training and testing sets
train dataset = torchvision.datasets.CIFAR10(root='data/', train=True,
download=True, transform=transform)
test dataset = torchvision.datasets.CIFAR10(root='data/', train=False,
download=True, transform=transform)
train loader = torch.utils.data.DataLoader(train dataset,
batch size=16, shuffle=True, num workers=2)
test loader = torch.utils.data.DataLoader(test dataset, batch size=16,
shuffle=False, num workers=2)
Files already downloaded and verified
Files already downloaded and verified
```

3.2 Network

```
class Net(nn.Module):
    def __init__(self, normalization=False):
        super(Net, self).__init__()
```

```
self.normalization = normalization
        # convolution block
        self.conv1 = nn.Conv2d(in channels=3, out channels=6,
kernel size=(5, 5), padding=0, stride=1)
        self.norm1 = nn.BatchNorm2d(num features=6)
        self.pool1 = nn.MaxPool2d(kernel_size=(2, 2), stride=2)
        self.conv2 = nn.Conv2d(in channels=6, out channels=16,
kernel_size=(5, 5), padding=0, stride=1)
        self.norm2 = nn.BatchNorm2d(num features=16)
        self.pool2 = nn.MaxPool2d(kernel size=(2, 2), stride=2)
        # fully connected block
        self.fc1 = nn.Linear(16 * 5 * 5, 120)
        self.fc2 = nn.Linear(120, 84)
        self.fc3 = nn.Linear(84, 10)
    def forward(self, x):
        # convolution block
        x = self.conv1(x)
        if self.normalization:
            x = self.norm1(x)
        x = F.relu(x)
        x = self.pool1(x)
        x = self.conv2(x)
        if self.normalization:
            x = self.norm2(x)
        x = F.relu(x)
        x = self.pool2(x)
        # flatten the signal
        x = x.view(x.size(0), -1)
        # fully connected block
        x = F.relu(self.fc1(x))
        x = F.relu(self.fc2(x))
        x = self.fc3(x)
        return x
```

3.3 Training

```
def train(model, criterion, optimizer, train_loader, test_loader,
num_epochs=10, use_gpu=False):

# move the model to the device (CPU or GPU)
if use_gpu and torch.cuda.is_available():
    model = model.cuda()
    criterion = criterion.cuda()

# create empty lists to store accuracy for each epoch
train_acc = []
test_acc = []
```

```
# iterate over epochs
    for epoch in tqdm(range(num epochs), desc='Training',
unit='epoch'):
        # set the model to training mode
        model.train()
        running loss = 0.0
        running correct = 0.0
        for step, [example, label] in enumerate(train loader):
            # move the data to the device (CPU or GPU)
            if use gpu and torch.cuda.is available():
                example = example.cuda()
                label = label.cuda()
            # zero the parameter gradients - prevent accumulation of
gradients
            optimizer.zero grad()
            # forward pass
            prediction = model(example)
            loss = criterion(prediction, label)
            # backward pass
            loss.backward()
            optimizer.step()
            # calculate accuracy on the current training batch
            _, predicted = torch.max(prediction.data, 1)
            running correct += (predicted == label).sum().item()
            running loss += loss.item()
        # calculate training accuracy for the current epoch
        train acc.append(running correct / len(train loader.dataset))
        # validate the model
        model.eval()
        correct = 0
        total = 0
        # set the model to evaluation mode
        for idx, [test example, test label] in
enumerate(tqdm(test loader, desc='Testing', unit='batch')):
            # move the data to the device (CPU or GPU)
            if use gpu and torch.cuda.is available():
                test example = test example.cuda()
                test label = test label.cuda()
```

```
# forward pass
            test prediction = model(test example)
            loss = criterion(test prediction, test label)
            # get the predicted class label
            , predicted = torch.max(test prediction.data, 1)
            total += test_label.size(0)
            correct += (predicted == test label).sum().item()
        # calculate test accuracy for the current epoch
        test acc.append(correct / total)
        # print the loss and accuracy
        print("\n *** Summary: Epoch [{} / {}] Train Accuracy: {}
Test Accuracy: {}***".format(epoch + 1, num_epochs, train_acc[-1],
test acc[-1]))
        torch.save(model.state dict(),
'{}/ConvNet{}.ckpt'.format(RESULTS DIR, epoch + 1))
    return train acc, test acc
# train the model
model = Net(normalization=True)
criterion = nn.CrossEntropyLoss()
optimizer = optim.Adam(model.parameters(), lr=1e-3)
train acc, test acc = train(model, criterion, optimizer, train loader,
test loader, num epochs=5, use gpu=False)
Testing: 100% | 625/625 [00:13<00:00, 46.13batch/s]
Training: 20% | | 1/5 [00:43<02:55, 43.81s/epoch]
*** Summary: Epoch [1 / 5] Train Accuracy: 0.45742 Test Accuracy:
0.5061***
Testing: 100%| 625/625 [00:13<00:00, 45.37batch/s]
Training: 40% | 2/5 [01:27<02:11, 43.81s/epoch]
*** Summary: Epoch [2 / 5] Train Accuracy: 0.55606 Test Accuracy:
0.5423***
Testing: 100%| 625/625 [00:13<00:00, 47.22batch/s]
Training: 60%| 3/5 [02:09<01:25, 42.90s/epoch]
*** Summary: Epoch [3 / 5] Train Accuracy: 0.59518 Test Accuracy:
0.5997***
Testing: 100% | 625/625 [00:13<00:00, 46.78batch/s]
Training: 80% | 4/5 [02:50<00:42, 42.23s/epoch]
```

```
*** Summary: Epoch [4 / 5] Train Accuracy: 0.62242 Test Accuracy: 0.6062***

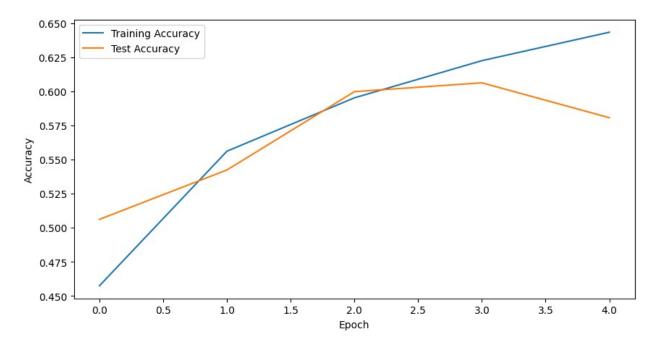
Testing: 100%| 625/625 [00:14<00:00, 42.41batch/s]
Training: 100%| 5/5 [03:35<00:00, 43.06s/epoch]

*** Summary: Epoch [5 / 5] Train Accuracy: 0.64328 Test Accuracy: 0.5806***
```

Plot how the accuracy (i.e. percentage of correctly classified images) of your model evolves for both, the training- and test-set of CIFAR10. Please make sure to add meaningful axis labels for your plot (y-axis as the accuracy and x-axis as the epoch number).

```
def plot_accuracy(train_acc, test_acc):
    # plot accuracy versus epoch number
    plt.plot(train_acc, label='Training Accuracy')
    plt.plot(test_acc, label='Test Accuracy')
    plt.xlabel('Epoch')
    plt.ylabel('Accuracy')
    plt.legend()
    plt.show()

# plot the accuracy
plot_accuracy(train_acc, test_acc)
```



What do you observe in the accuracy plot, in particular if you compare the train and test accuracy? Do you have an explanation for that?

Without the use of normalization or augmentation techniques, the network's performance on the training set is significantly higher than on the test set. This discrepancy suggests that the network may have too many degrees of freedom for the relatively simple CIFAR dataset, leading to high bias or overfitting. In other words, the network may be memorizing the training data rather than learning the underlying patterns that would allow it to generalize to new, unseen data.

To address this issue, regularization techniques such as normalization and data augmentation can be employed. Normalization helps to ensure that the network learns at the same speed everywhere by scaling the input data to a consistent range, while data augmentation increases the variance in the training set, providing the network with more diverse examples to learn from. By using these techniques, we can reduce overfitting and improve the network's ability to generalize to new data.

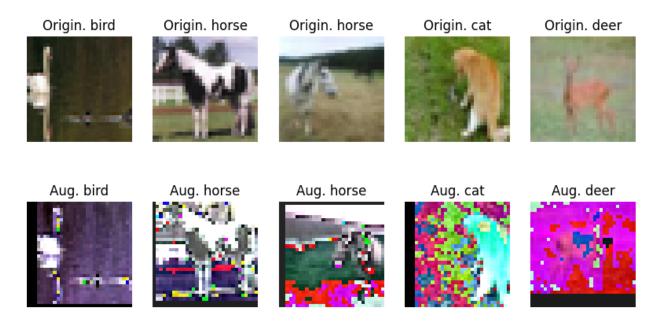
3.4 Augmentation

Please apply random horizontal flipping, random cropping with padding=4, and ColorJitter to the CIFAR10 images and visualize a few pairs of augmented images against their original.

```
# define the transformations
transform aug = T.Compose([
    T.RandomHorizontalFlip(),
    T.RandomCrop(32, padding=4),
    T.ColorJitter(brightness=0.5, contrast=0.5, saturation=0.5,
hue=0.5),
    T.ToTensor().
    T.Normalize(mean=[0.5, 0.5, 0.5], std=[0.5, 0.5, 0.5])
])
# load the dataset and apply the transformations
train dataset aug = torchvision.datasets.CIFAR10(root='data/',
train=True, download=True, transform=transform aug)
test dataset aug = torchvision.datasets.CIFAR10(root='data/',
train=False, download=True, transform=transform aug)
train loader aug = torch.utils.data.DataLoader(train dataset,
batch size=16, shuffle=True, num workers=2)
test loader aug = torch.utils.data.DataLoader(test dataset,
batch size=16, shuffle=False, num workers=2)
Files already downloaded and verified
Files already downloaded and verified
# visualize a few pairs of augmented images against their original
images
fig, axs = plt.subplots(\frac{2}{5}, figsize=(\frac{10}{5}))
for i in range(5):
```

```
# get a random sample from the training set
    index = torch.randint(len(train dataset), size=(1,))
    img_aug, label_aug = train_dataset_aug[index]
    img, label = train dataset[index]
    # convert image back from tensor
    if isinstance(img, torch.Tensor) or isinstance(img_aug,
torch.Tensor):
        img = img.numpy()
        img_aug = img_aug.numpy()
    # convert images
    img_original = T.ToPILImage()(np.transpose(img, (1, 2, 0)))
    img augmented = T.ToPILImage()(np.transpose(img aug, (1, 2, 0)))
    class name = train dataset.classes[label]
    # plot the original image
    ax = axs[0, i]
    ax.imshow(img original)
    ax.set title(f'Origin. {class name}')
    ax.axis('off')
    # plot the augmented image
    ax = axs[1, i]
    ax.imshow(img augmented)
    ax.set title(f'Aug. {class name}')
    ax.axis('off')
# set title
fig.suptitle('CIFAR10 Augmented Images')
# save the plot
plt.savefig(f'{RESULTS DIR}/augmented images.png')
```

CIFAR10 Augmented Images

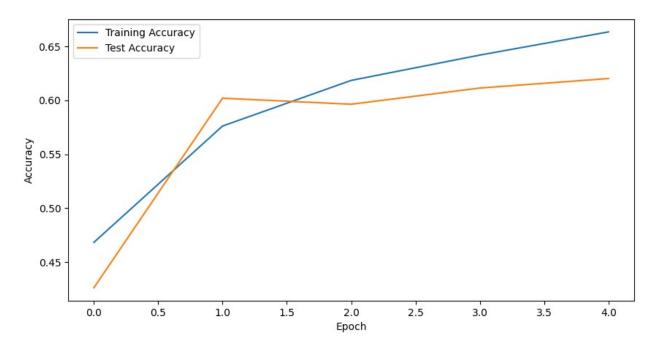


Train your model using the same setup as above, however, with additional normalization (normalize your input images channel-wise using $\mu = 0.5$ and $\sigma = 0.5$) and data augmentation.

```
# train the model
model = Net(normalization=True)
criterion = nn.CrossEntropyLoss()
optimizer = optim.Adam(model.parameters(), lr=1e-3)
train acc, test acc = train(model, criterion, optimizer,
train_loader_aug, test_loader_aug, num_epochs=5, use_gpu=False)
Testing: 100% | 625/625 [00:13<00:00, 46.61batch/s]
Training: 20% | | 1/5 [00:43<02:55, 43.98s/epoch]
*** Summary: Epoch [1 / 5] Train Accuracy: 0.46842 Test Accuracy:
0.4264***
                     | 625/625 [00:13<00:00, 45.31batch/s]
Testing: 100%
                 2/5 [01:26<02:08, 42.95s/epoch]
Training: 40%
*** Summary: Epoch [2 / 5] Train Accuracy: 0.57612 Test Accuracy:
0.6019***
Testing: 100%
                   | 625/625 [00:13<00:00, 46.90batch/s]
Training: 60% | 3/5 [02:08<01:25, 42.80s/epoch]
*** Summary: Epoch [3 / 5] Train Accuracy: 0.61836 Test Accuracy:
0.5963***
```

Plot the accuracy of this model and the one you trained before over time. Make sure to use proper axis labels and a meaningful legend.

```
# plot the accuracy
plot_accuracy(train_acc, test_acc)
```



Based on your results, provide a brief discussion of why the extra normalization and augmentation improves/harms the performance of your model.

Data augmentation and normalization are two crucial techniques for improving the performance of deep learning models. Augmentation involves artificially increasing the size of the training dataset by applying various transformations to the original images, such as rotation, scaling, and flipping. This not only provides the network with more data to learn from, but also increases the variance in the dataset, making the network more robust to changes in the input.

Normalization, on the other hand, involves scaling the pixel values of the input images to a fixed range, typically between 0 and 1. This ensures that the input data is consistent and well-behaved, which in turn allows the network to learn more effectively and converge faster. By bringing the values of the input image into a specific range, the network can better process and analyze the image, leading to improved performance and accuracy.

Moreover, normalization can help regulate the learning process of the network by ensuring that all input features have the same range and that there are no extreme values that could dominate the gradient updates. This can help prevent overfitting, which occurs when the network becomes too specialized to the training data and fails to generalize to new, unseen examples.

In the above experiment, it an improvement in accuracy is observable for the test dataset. This might be due to a significant reduction in overfitting due to the use of data augmentation and normalization. The network was able to learn more effectively and generalize better to new examples, resulting in improved performance and accuracy.

Task 4: Activation and Saliency Maps

In this task, we will try to visualize what CNNs learn by means of the activation map, which is considered to be more informative than just visualizing e.g. the kernel weights.

4.1 Instantiate a pre-traind ResNet18

```
from torchvision.models import resnet18
# 1. instantiate a pre-trained ResNet18
model = resnet18(pretrained=True)
# set the model to evaluation mode
model.eval()
# print the model architecture
print(model)
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 running 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_running_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 running 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 running 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_running 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 running 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 running stats=True)
      (downsample): Sequential(
        (0): Conv2d(64, 128, \text{kernel size}=(1, 1), \text{stride}=(2, 2),
bias=False)
        (1): BatchNorm2d(128, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
    (1): BasicBlock(
      (conv1): Conv2d(128, 128, kernel size=(3, 3), stride=(1, 1),
padding=(1, 1), bias=False)
      (bn1): BatchNorm2d(128, eps=1e-05, momentum=0.1, affine=True,
track running 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 running 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_running_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 running stats=True)
      (downsample): Sequential(
        (0): Conv2d(128, 256, kernel size=(1, 1), stride=(2, 2),
bias=False)
        (1): BatchNorm2d(256, eps=1e-05, momentum=0.1, affine=True,
track running 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 running 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 running 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 running 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 running stats=True)
      (downsample): Sequential(
        (0): Conv2d(256, 512, kernel size=(1, 1), stride=(2, 2),
bias=False)
        (1): BatchNorm2d(512, eps=1e-05, momentum=0.1, affine=True,
track_running_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 running 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 running stats=True)
    )
  (avgpool): AdaptiveAvgPool2d(output size=(1, 1))
  (fc): Linear(in features=512, out features=1000, bias=True)
/Users/janinaalicamattes/miniforge3/envs/pytorch-py11/lib/python3.11/
site-packages/torchvision/models/ utils.py:208: UserWarning: The
parameter 'pretrained' is deprecated since 0.13 and may be removed in
the future, please use 'weights' instead.
  warnings.warn(
/Users/janinaalicamattes/miniforge3/envs/pytorch-py11/lib/python3.11/
site-packages/torchvision/models/_utils.py:223: UserWarning: Arguments
other than a weight enum or `None for 'weights' are deprecated since
0.13 and may be removed in the future. The current behavior is
equivalent to passing `weights=ResNet18 Weights.IMAGENET1K V1`. You
can also use `weights=ResNet18 Weights.DEFAULT` to get the most up-to-
date weights.
 warnings.warn(msq)
# test -- print a convolutional layer information
print(model.layer3[0].downsample[1])
BatchNorm2d(256, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
```

4.2 Intermediate Activation Maps

```
# extract intermediate activations
# e.g., the out after the first conv in the first residual block and
# the out after the last conv in the last residual block
activations = {}
def get activation(name):
    def hook(model, input, output):
        activations[name] = output.detach()
    return hook
# register the hooks
h1 = model.avgpool.register forward hook(get activation('avgpool'))
h2 = model.maxpool.register forward hook(get activation('maxpool'))
h3 =
model.layer3[0].downsample[1].register forward hook(get activation('co
(('am
h4 =
model.layer4[1].conv2.register forward hook(get activation('conv2'))
# perform the forward pass -- getting the outputs
output = model(X)
# print the shape of the intermediate activations
print(activations['avgpool'].shape)
# detach the hooks
h1.remove()
h2.remove()
h3.remove()
h4.remove()
torch.Size([1, 512, 1, 1])
# print shape of all the intermediate activations
for key in activations:
    print(f'{key} shape: {activations[key].shape}')
maxpool shape: torch.Size([1, 64, 56, 56])
comp shape: torch.Size([1, 256, 14, 14])
conv2 shape: torch.Size([1, 512, 7, 7])
avgpool shape: torch.Size([1, 512, 1, 1])
```

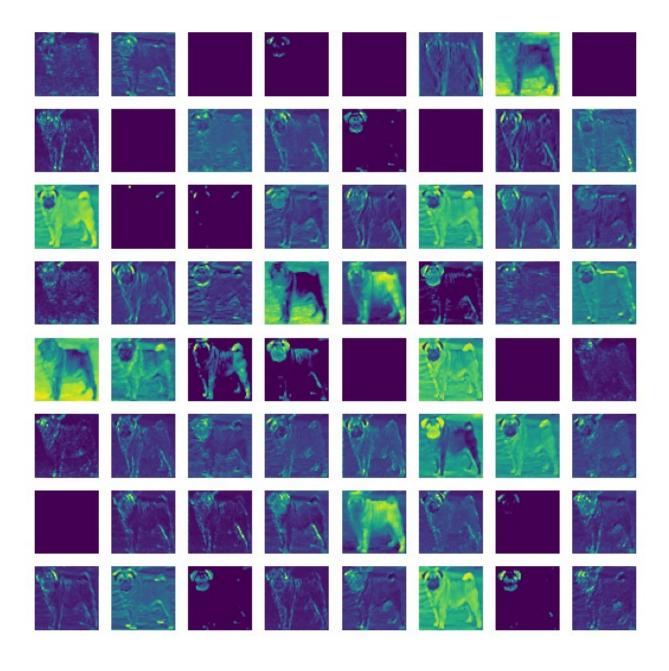
4.3 Visualize the Activation Maps

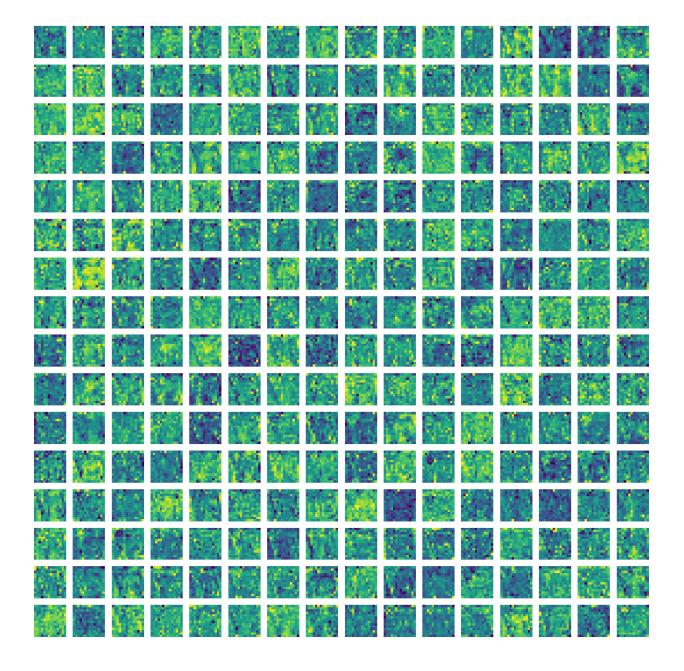
```
import matplotlib.pyplot as plt

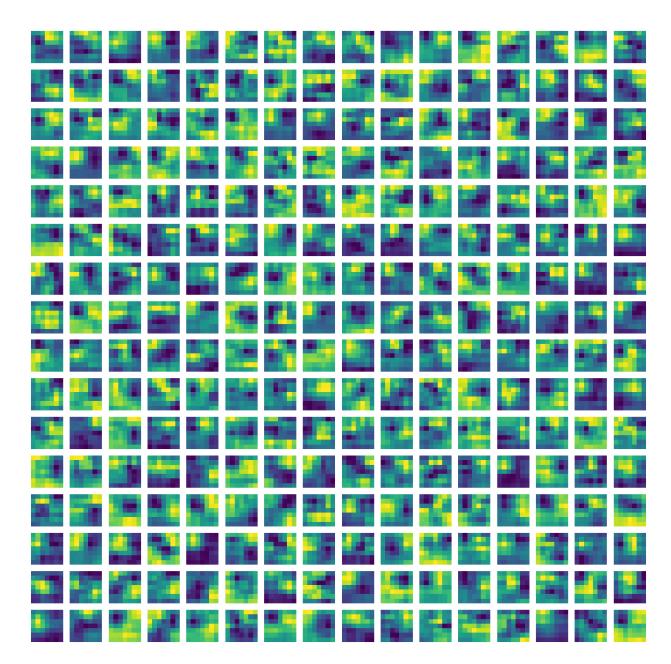
# Extract the activations for each layer
maxpool_act = activations['maxpool'].squeeze(0)
comp_act = activations['comp'].squeeze(0)
conv2_act = activations['conv2'].squeeze(0)
avgpool_act = activations['avgpool'].squeeze(0)
```

```
# Plot the activations for the maxpool layer
fig, axarr = plt.subplots(8, 8, figsize=(8, 8))
fig.suptitle('MaxPool Activation', fontsize=16, y=1.05) # Add title
to the figure
# title to figure
for idx in range(64):
    axarr[idx // 8, idx % 8].imshow(maxpool act[idx, :, :],
cmap='viridis')
    axarr[idx // 8, idx % 8].axis('off')
plt.show()
# Plot the activations for the comp layer
fig, axarr = plt.subplots(16, 16, figsize=(16, 16))
fig.suptitle('Comp Activation', fontsize=16, y=1.05) # Add title to
the figure
for idx in range(256):
    axarr[idx // 16, idx % 16].imshow(comp act[idx, :, :],
cmap='viridis')
    axarr[idx // 16, idx % 16].axis('off')
plt.show()
# Plot the activations for the conv2 layer
fig, axarr = plt.subplots(16, 16, figsize=(16, 16))
fig.suptitle('Conv2 Activation', fontsize=16, y=1.05) # Add title to
the figure
for idx in range(256):
    axarr[idx // 16, idx % 16].imshow(conv2 act[idx, :, :],
cmap='viridis')
    axarr[idx // 16, idx % 16].axis('off')
plt.show()
```

MaxPool Activation



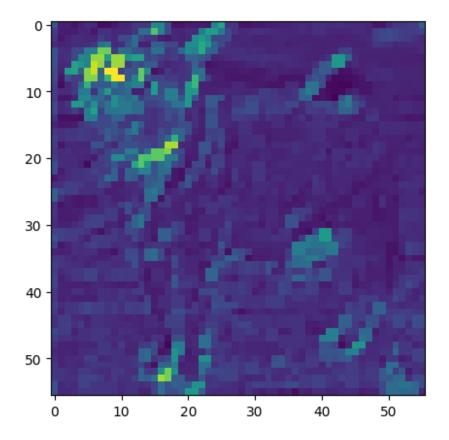


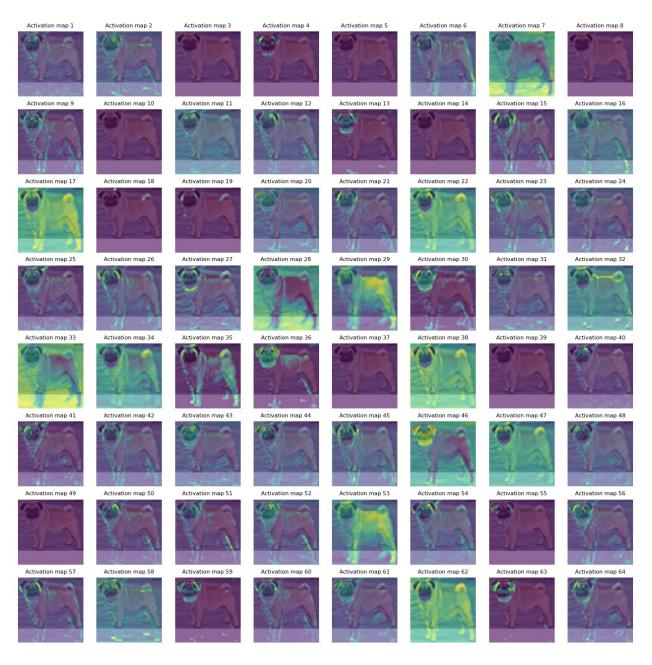


4.4 Plot Image and Activation Map

img = np.array(img)
print(img.shape)

```
(824, 1034, 3)
import numpy as np
import matplotlib.pyplot as plt
from PIL import Image
# Resize the original image to have the same height and width as the
activation maps
resize factor = min(maxpool act.shape[1] / img.shape[1],
maxpool act.shape[2] / img.shape[2])
resized height = int(img.shape[0] * resize factor)
resized width = int(img.shape[1] * resize factor)
if resized height > 0 and resized width > 0:
    resized img = np.array(Image.fromarray(img).resize((resized width,
resized height)))
    print("Error: resized image has a height or width of zero or
less")
    resized img = None
# Convert the activation maps to heatmaps
heatmaps = []
for i in range(maxpool act.shape[0]):
    heatmap = plt.imshow(maxpool act[i], cmap='viridis')
    heatmaps.append(heatmap)
# Overlay the heatmaps on the original image and plot in a grid
if resized img is not None:
    num cols = 8
    num rows = np.ceil(maxpool act.shape[0] /
num cols).astype(np.int32)
    fig, axs = plt.subplots(num rows, num cols, figsize=(num cols * 2,
num rows * 2))
    fig.suptitle('Activation Maps Overlayed on Original Image',
fontsize=16, y=1.05)
    for i in range(maxpool act.shape[0]):
        row = i // num_cols
        col = i % num cols
        ax = axs[row, col]
        ax.imshow(resized img)
        ax.imshow(maxpool act[i], cmap='viridis', alpha=0.6)
        ax.set_title(f'Activation map {i + 1}', fontsize=8)
        ax.axis('off')
    plt.show()
else:
    print("Error: cannot overlay heatmaps on original image")
```





4.5 Saliency Map

Introduced in: *Deep Inside Convolutional Networks: Visualising Image Classification Models and Saliency Maps*

```
# illustrate how gather() works
def gather example():
    N, C = 4, 5
    s = torch.randn(N, C)
    y = torch.LongTensor([1, 2, 1, 3])
    print(s)
    print(y)
    print(s.gather(1, y.view(-1, 1)).squeeze())
gather example()
# load the images
names = ['catdog 243.png', 'catdog 243.png', 'snake 56.png',
'spider 72.png']
# merge names with the path
names = [f'./data/{name}' for name in names]
X = [np.array(Image.open(name).convert('RGB')) for name in names]
y = [int(s.rsplit(' ')[1].rsplit(' .')[0])  for s in names]
# intentionally change the label to a wrong one
v[1] = 285
def preprocess(img, size=224):
    transform = T.Compose([
        T.Resize(size),
        T.ToTensor(),
        T.Lambda(lambda x: x[None]), # equivalent to unsqueeze()
    1)
    return transform(img)
tensor([[ 0.1899, -0.6072, 0.1361, 0.9758, 0.4135],
        [-0.7803, 0.7713, 0.3468, -0.8428, -1.6304],
        [0.4518, -1.5922, -1.0456, 1.1665, -0.3299],
        [1.1794, -0.0566, -0.2523, 0.0968, 1.0805]])
tensor([1, 2, 1, 3])
tensor([-0.6072, 0.3468, -1.5922, 0.0968])
```

a) Complete the saliency map function.

```
def saliency_maps(X, y, model):
    Compute a class saliency map using the model for images X and
labels y.

Input:
    X: Input images; Tensor of shape (N, 3, H, W)
    y: Labels for X; LongTensor of shape (N,)
    model: A pretrained CNN that will be used to compute the
saliency map.
```

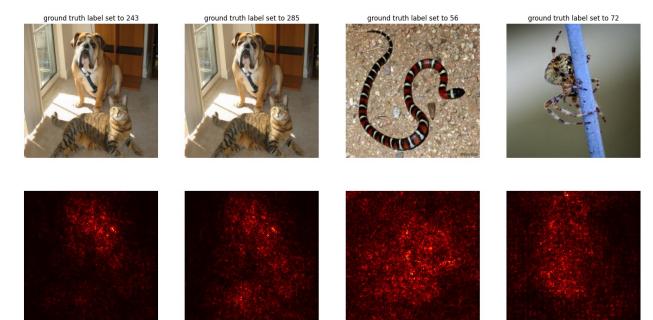
```
Returns:
    - saliency: A Tensor of shape (N, H, W) giving the saliency maps
for the input
   images.
   # don't compute gradients
   for param in model.parameters():
        param.requires grad = False
   # Make sure the model is in "test" mode
   model.eval()
   # Wrap the input tensors in Variables
   X var = X.clone().detach()
   # if not tensor
   if not isinstance(X var, torch.Tensor):
        X var = torch.tensor(X var)
   # calculate gradient of highest score w.r.t input
   X var.requires grad = True
   y var = y.clone().detach()
   saliency = None
   # forward pass
   prediction = model(X var)
   loss = F.cross_entropy(prediction, y var)
   # backward pass
   loss.backward()
   # compute the saliency map
    saliency = X var.grad.data.abs().mean(dim=1)
   # normalize the saliency map
    saliency = (saliency - saliency.min()) / (saliency.max() -
saliency.min())
    return saliency
```

b) Visualize the original images together with the saliency maps.

Visualize the original image with the saliency maps together and provide a short discussion on why different ground truth labels with even the same input image would yield different saliency maps, for instance, the catdog_243.png image with y = 243 (bull mastiff) and y = 285 (Egyptian cat).

```
# Please use the same pre-trained network from the previous task for
this task
model = resnet18(pretrained=True)
```

```
# if you finished the task 3.1 and defined your model, then this
function should run flawlessly
def show saliency maps(X, y):
    # Convert X and y from numpy arrays to Torch Tensors
    X tensor = torch.cat([preprocess(Image.fromarray(x)) for x in X],
dim=0)
    y tensor = torch.LongTensor(y)
    # Compute saliency maps for images in X
    saliency = saliency maps(X tensor, y tensor, model)
    # Convert the saliency map from Torch Tensor to numpy array and
show images
    # and saliency maps together.
    saliency = saliency.numpy()
    N = len(X)
    for i in range(N):
        plt.subplot(2, N, i + 1)
        plt.imshow(X[i])
        plt.axis('off')
        plt.title('ground truth label set to {}'.format(y[i]))
        plt.subplot(2, N, N + i + 1)
        plt.imshow(saliency[i], cmap=plt.cm.hot)
        plt.axis('off')
        plt.gcf().set size inches(20, 10)
    plt.show()
show_saliency_maps(X, y)
```



Discussion

Visualize the original image with the saliency maps together and provide a short discussion on why different ground truth labels with even the same input image would yield different saliency maps, for instance the $catdog_243.png$ image with y = 243 (bull mastiff) and y = 285 (Egyptian cat).

Visualising the original image with the saliency maps can help understand how the model makes its predictions. In the case of the catdog_243.png image, it can be seen that the model focuses on different salient features depending on the ground truth label chosen for the backward pass.

When the ground truth is set to y=243 (Bullmastiff), the saliency map highlights the features of the dog in the input image, such as its muzzle, ears, and coat. This indicates that the model is using these features to distinguish the Bullmastiff from other elements in the image.

Conversely, when the ground truth is set to y=285 (Egyptian cat), the saliency map emphasizes the features of the reclining cat, such as its eyes, whiskers, and fur. This suggests that the model is using these features to distinguish the Egyptian cat from other elements in the image, including the dog.

Interestingly, we can also observe that the saliency map for the Egyptian cat contains some pixels from the dog's fur. This may be due to the fact that the model is using the contrast between the cat's and the dog's fur to distinguish between the two animals. Additionally, the wider spread of the saliency map pixels highlighted for the Egyptian cat could be caused by the fact that the model was actively labeled with the "incorrect" label of Egyptian cat, whereas in the training data, the label was possibly set to Bullmastiff if two objects of different categories appeared in the same image. In this case, the model may be incorporating the salient features of the cat in order to distinguish the expected class.

This experiment further shows how saliency maps can be used to interpret the decisions of a deep learning model.