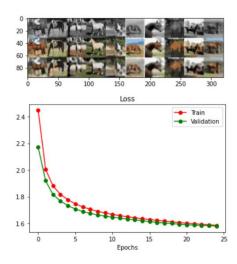
Part A code

```
class PoolUpsampleNet(nn.Module):
   def __init__(self, kernel, num_filters, num_colours, num_in_channels):
       super().__init__()
       # Useful parameters
       padding = kernel // 2
       self.block1 = nn.Sequential(
           nn.Conv2d(num_in_channels, num_filters, kernel, padding = padding),
           nn.MaxPool2d(kernel_size=2, stride=2),
           nn.BatchNorm2d(num_features=num_filters),
           nn.ReLU()
       )
       self.block2 = nn.Sequential(
           nn.Conv2d(num_filters, 2*num_filters, kernel, padding = padding),
           nn.MaxPool2d(kernel size=2, stride=2),
           nn.BatchNorm2d(num_features=2*num_filters),
           nn.ReLU()
       self.block3 = nn.Sequential(
           nn.Conv2d(2*num_filters, num_filters, kernel, padding = padding),
           nn.Upsample(scale_factor=2),
           nn.BatchNorm2d(num_features=num_filters),
           nn.ReLU()
       self.block4 = nn.Sequential(
           nn.Conv2d(num_filters, num_colours, kernel, padding = padding),
           nn.Upsample(scale_factor=2),
           nn.BatchNorm2d(num_features=num_colours),
           nn.ReLU(),
           nn.Conv2d(num_colours, num_colours, kernel, padding=padding)
       )
```

Part A visualization and comments:



The results are not good at all since the output graphs are quite vague and blurred, and the validation accuracy has only 41.2% after 25 epochs.

Part A Q3 six values to report:

To calculate the number of weights, outputs and connections, I follow the formula from the lecture slides, and here I would show several layers' computation methods, and the rest would be pretty much the same. From the images to the first convolutional layer, we have the output dimensions as (assuming kernel size k being odd):

$$\hat{W} = \hat{H} = [(32 - k + 2 \times (k//2))/1 + 1]$$

= $32 - k + 2 \times \frac{k-1}{2} + 1$
= 32

and so we have:

- Number of weights: $k^2 imes \mathrm{NIC} imes \mathrm{NF}$
- ullet Number of outputs: $32^2 imes NF = 1024 imes NF$
- Number of connections: $32^2 \times k^2 \times NF \times NIC = 1024 \times k^2 \times NF \times NIC$ From the first convolutional layer to the first max-pooling layer, since the max-pooling does not have parameters, the number of weights would be 0, which is the same situation when dealing with the Upsampling layer. Since we are assuming the kernel size and stride of the max-pooling layer to be 2, we can calculate the number of outputs and connections similar as above. Following the above procedure, the final 6 figures to report are following:
- When each input dimension (width/height) is not doubled (original input):
 - \circ Number of weights: $k^2 imes NIC imes NF + 4k^2 imes NF^2 + k^2 imes NC imes NF + K^2 imes NC^2$
 - $\circ~$ Number of outputs: 2240 imes NF + 2304 imes NC
 - Number of connections:

$$1024k^2 \times NF \times NIC + 1792 \times NF + 640 \times K^2 \times NF^2 + 256k^2 \times NC \times NF + 1024 \times NC + 1024k^2 \times NC^2 + 1024k^2 \times NC \times NF + 1024$$

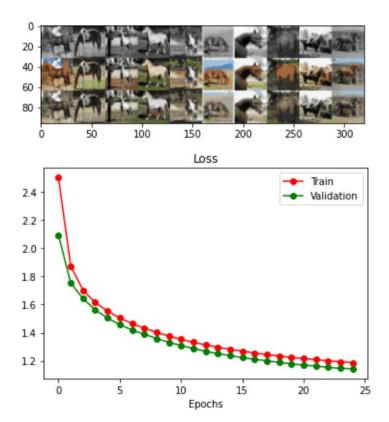
- When each input dimension (width/height) is doubled, we would have the original answers multiplied by 4 since we have two dimensions for an image, and each dimension is being doubled:
 - \circ Number of weights: $4 imes (k^2 imes NIC imes NF + 4k^2 imes NF^2 + k^2 imes NC imes NF + K^2 imes NC^2)$
 - \circ Number of outputs: $4 \times (2240 \times NF + 2304 \times NC)$
 - Number of connections:

$$4 imes (1024k^2 imes NF imes NIC + 1792 imes NF + 640 imes K^2 imes NF^2 + 256k^2 imes NC imes NF + 1024 imes NC + 1024k^2 imes NC^2)$$

Part B Q1 code:

```
self.block1 = nn.Sequential(
   nn.Conv2d(num_in_channels, num_filters, kernel_size=kernel,
             stride = 2, padding = 1),
   nn.BatchNorm2d(num features=num filters),
   nn.ReLU()
)
self.block2 = nn.Sequential(
   nn.Conv2d(num_filters, 2*num_filters, kernel_size=kernel,
             stride = 2, padding = 1),
   nn.BatchNorm2d(num features=2*num filters),
   nn.ReLU()
)
self.block3 = nn.Sequential(
   nn.ConvTranspose2d(2*num filters, num filters, kernel size=kernel,
                      stride = 2, padding = 1, output_padding = 1,
                      dilation=1),
   nn.BatchNorm2d(num features=num filters),
   nn.ReLU()
)
self.block4 = nn.Sequential(
   nn.ConvTranspose2d(num filters, num colours, kernel size=kernel,
                      stride = 2, padding = 1, output padding = 1,
                      dilation=1),
   nn.BatchNorm2d(num features=num colours),
   nn.ReLU(),
   nn.Conv2d(num colours, num colours, kernel size=kernel, padding = 1)
```

Part B Q2 visualization:



The validation accuracy is better than the previous model.

Part B Q3-5:

- ▼ Questions 3 5
 - Q3: The results are better than what in part A, as we can notice that the validation accuracy has come to 55.3% and the graphs are
 clearer with more accurate color pixels right now. The validation loss for the ConvTransposeNet is 1.1426, which is lower than 1.5788,
 which is the validation loss for the PoolUpsampleNet. The reason for this case might be that the Upsample layer has no trainable
 parameters, but ConvTranspose2d layer has trainable parameters, which can capture the gradient information from loss and so it can
 further update the model.
 - Q4: To make sure the model is in the same shape after the first two nn.Conv2d layers, for kernel size being 4, we need padding = 1, and for kernel size being 5, we need padding = 2 since torch.nn calculate the output size using round-down operation in math. The formula for calculating the padding for kernel size being 5 would be:

$$\lfloor \frac{32+2 \times padding - 5}{2} + 1 \rfloor = 16$$

and the formula to calculate the padding when kernel size is 4 is similar except replacing 5 into 4 in the above formula. For the $nn.\ ConvTranspose2d$ layers, we need $2 \times padding - output_padding = 2$ when kernel size is 4, and $2 \times padding - output_padding = 3$ when kernel size is 5. The formula for kernel size being 5 would be:

$$(8-1) \times 2 - 2 \times padding + 1 \times (5-1) + output_padding + 1 = 16$$

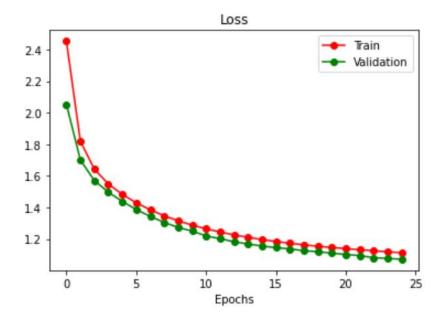
and it is similar when kernel size is 4 with replacing of 5 into 4 in the above formula.

Q5: As the batch size increases, the training/validation loss increases as well, and the validation accuracy decreases, which means the
quality of the output images are decreasing.

Part C Q1 code:

```
self.block1 = nn.Sequential(
   nn.Conv2d(num_in_channels, num_filters, kernel_size=kernel,
             stride = 2, padding = 1),
   nn.BatchNorm2d(num features=num filters),
   nn.ReLU()
)
self.block2 = nn.Sequential(
   nn.Conv2d(num_filters, 2*num_filters, kernel_size=kernel,
             stride = 2, padding = 1),
   nn.BatchNorm2d(num_features=2*num_filters),
   nn.ReLU()
)
self.block3 = nn.Sequential(
   nn.ConvTranspose2d(2*num_filters, num_filters, kernel_size=kernel,
                      stride = 2, padding = 1, output_padding = 1,
                      dilation=1),
   nn.BatchNorm2d(num features=num filters),
   nn.ReLU()
)
self.block4 = nn.Sequential(
   nn.ConvTranspose2d(2*num filters, num colours, kernel size=kernel,
                      stride = 2, padding = 1, output_padding = 1,
                      dilation=1),
   nn.BatchNorm2d(num features=num colours),
   nn.ReLU()
self.conv5 = nn.Conv2d(num_in_channels + num_colours, num_colours,
                      kernel_size=kernel, padding = 1)
```

Part C Q2 curve:



Part C Q3:

Question 3

The result is better than the previous model qualitatively since now we can see the graphs much clearer with more accurate color pixels than the previous model, and the validation loss and accuracy are both improved by the skip connections. The reasons for skip connections improving our CNN models are:

- 1. The *skip connections* can provide the relevant locality information where it is needed by when doing the *Upsampling* or *ConvTranspose2d*, we find the last layer before the image is shrinked, where the image still had the same size and simply add it pixel-wise to the upsampled image so that the now upsampled feature map has both the locality information lost in shrinking the image, say max-pooling or strided convolution layers, but also the dominant info after shrinking the image. This allows for much better detail in the task like prediction color pixels where we need both local information around the predicted pixel and also the global information in the image with previous size.
- 2. The *skip connections* will have more trainable parameters so that we can capture more information from the loss in the backward propagation to update our model better.

Part D.1 code:

Part D.2 code:

```
# --- YOUR CODE GOES HERE ---
BCEcls = nn.BCEWithLogitsLoss(pos_weight=torch.tensor([h['cls_pw']], device=device))
# -----
BCEobj = nn.BCEWithLogitsLoss(pos_weight=torch.tensor([h['obj_pw']], device=device))
```

Part D.2 graphs:

