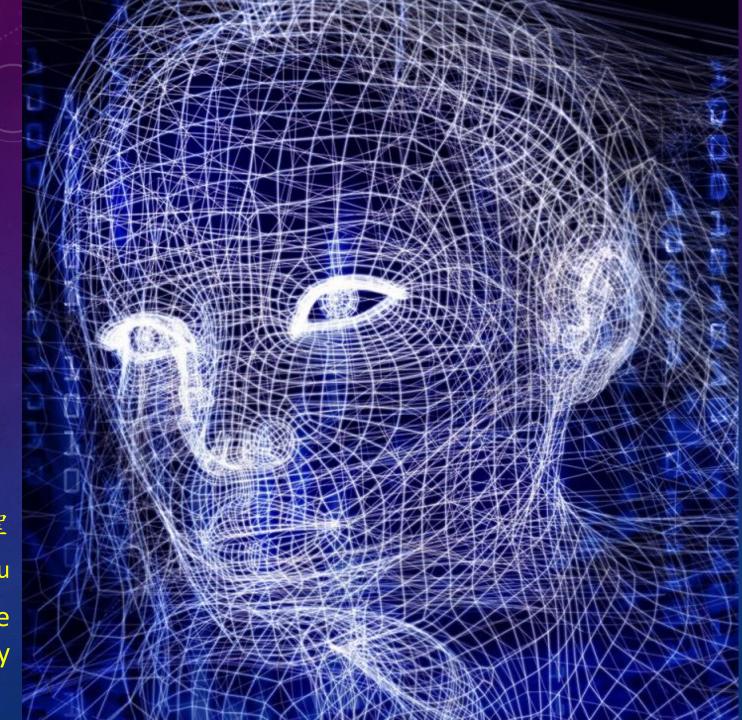


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Overview

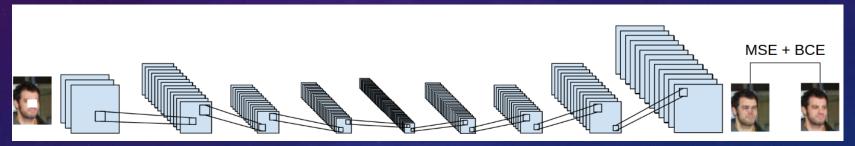
- 1. Fractionally Strided Convolution
- 2. Introduction to Autoencoder
- 3. Variational AutoEncoder (VAE)

(a.k.a. Deconvolution) Fractionally strided convolutions

(a.k.a. Deconvolutions or Transposed Convolutions)

The need for transposed convolutions generally arises from the desire to use a transformation going in the opposite direction of a normal convolution.

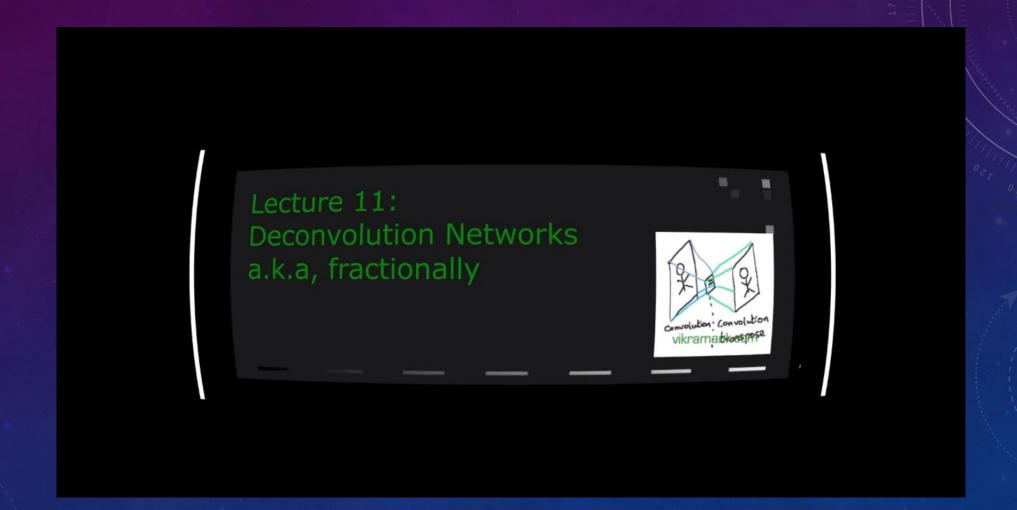
- From something that has the shape of the output of some convolution to something that has the shape of its input while maintaining a connectivity pattern that is compatible with said convolution.
 - (1) Transformation as the decoding layer of a convolutional autoencoder, e.g., Image synthesis.



(2) Project feature maps to a higher-dimension space, e.g., Feature visualization

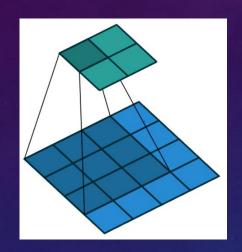


Lecture 11: Deconvolutional Networks



Convolution as a Matrix Operation

Take for example the convolution represented on the left. If the input and output were to be unrolled into vectors from left to right, top to bottom, the convolution could be represented as a sparse matrix C where the non-zero elements are the elements $w_{i,j}$ of the kernel (with i and j being the row and column of the kernel respectively):

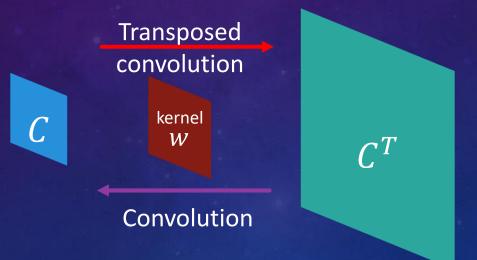


↓ *kernel*

$/w_{0,0}$	$w_{0,1}$	$w_{0,2}$	0	$w_{1,0}$	$w_{1,1}$	$w_{1,2}$	0	$w_{2,0}$	$w_{2,1}$	$w_{2,2}$	0	0	0	0	0
0	$w_{0,0}$	$w_{0,1}$	$w_{0,2}$	0	$w_{1,0}$	$w_{1,1}$	$w_{1,2}$	0	$w_{2,0}$	$w_{2,1}$	$w_{2,2}$	0	0	0	0
0	0	0	0	$w_{0,0}$	$w_{0,1}$	$w_{0,2}$	0	$w_{1,0}$	$w_{1,1}$	$w_{1,2}$	0	$w_{2,0}$	$w_{2,1}$	$w_{2,2}$	0
0 /	0	0	0	0	$w_{0,0}$	$w_{0,1}$	$w_{0,2}$	0	$w_{1,0}$	$w_{1,1}$	$w_{1,2}$	0	$w_{2,0}$	$w_{2,1}$	$w_{2,2}$

This linear operation takes the input matrix flattened as a 16-dimensional vector and produces a 4-dimensional vector that is later reshaped as the 2×2 output matrix.

- Since a kernel \boldsymbol{w} defines a convolution, but whether it's a normal convolution or a transposed convolution is determined by how the forward and backward passes are computed.
- We'll proceed somewhat backwards with respect to the convolution arithmetic, deriving the
 properties of each transposed convolution by referring to the direct convolution with which it
 shares the kernel, and defining the equivalent direct convolution.



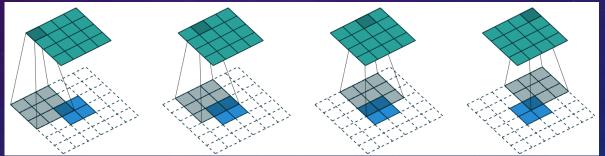
 C^T : stands for transposed C

Note that: It is always possible to emulate a transposed convolution with a normal convolution. The disadvantage is that it usually involves adding many columns and rows of zeros to the input, resulting in a *much less efficient implementation*.

The transposed convolution can be considered as the operation that allows to recover the *shape* of this initial feature map.

No zero padding, unit strides, transposed

To understand the logic behind zero padding is to consider the connectivity pattern of the transposed convolution and use it to guide the design of the equivalent convolution. It is possible to determine similar observations for the other elements of the image, giving rise to the following relationship:



The transpose of convolving a 3x3 kernel over a 4x4 input using unit strides

(i.e.,
$$i = 4$$
, $k = 3$, $s = 1$ and $p = 0$)

It is equivalent to convolving a 3×3 kernel over a 2×2 input (blue) padded with a 2×2 border of zeros using unit strides.

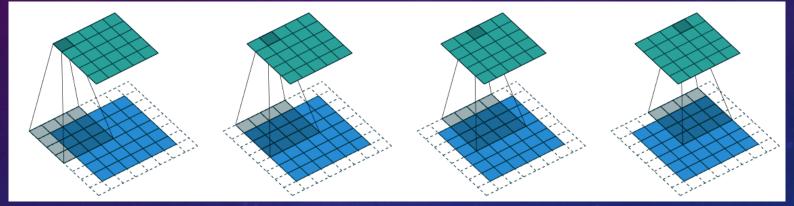
(i.e.,
$$i' = 2$$
, $k' = k$, $s' = 1$ and $p' = 2$)

Relationship 8. A convolution described by s = 1, p = 0 and k has an associated transposed convolution Described by k' = k, s' = s and p' = k - 1 and its output size is

$$o' = i' + (k-1)$$

Zero padding, unit strides, transposed

Knowing that the transpose of a non-padded convolution is equivalent to convolving a zero padded input, it would be reasonable to suppose that the trans-pose of a zero padded convolution is equivalent to convolving an input padded with less zeros.



The transpose of convolving a 4×4 kernel over a 5×5 input (green) padded with a 2×2 border of zeros using unit strides (i.e., i = 5, k = 4, s = 1 and p = 2)

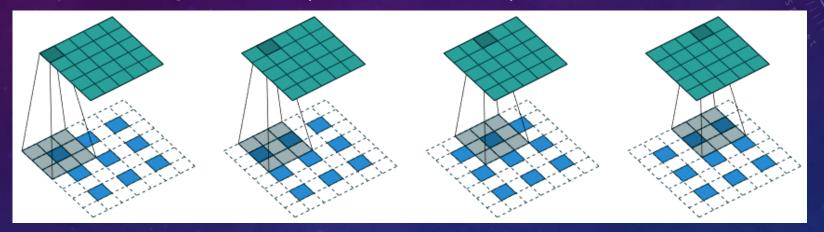
It is equivalent to convolving a 4×4 kernel over a 6×6 input padded with a 1×1 border of zeros using unit strides (i.e., i'=6, k'=k, s'=1 and p'=1)

Relationship 9. A convolution described by s=1, k, and p has an associated transposed convolution described by k'=k, s'=s and p'=k-p-1 and its output size is

$$o' = i' + (k-1) - 2p$$

Zero padding, non-unit strides, transposed

When the convolution's input size i is such that i+2p-k is a multiple of s, the analysis can extended to the zero padded case by combining **Relationship 9** and **Relationship 12**:



The transpose of convolving a 3x3 kernel over a 5x5 input padded with a 1x1 border of zeros using 2x2 strides (i.e., i = 5, k = 3, s = 2 and p = 1)

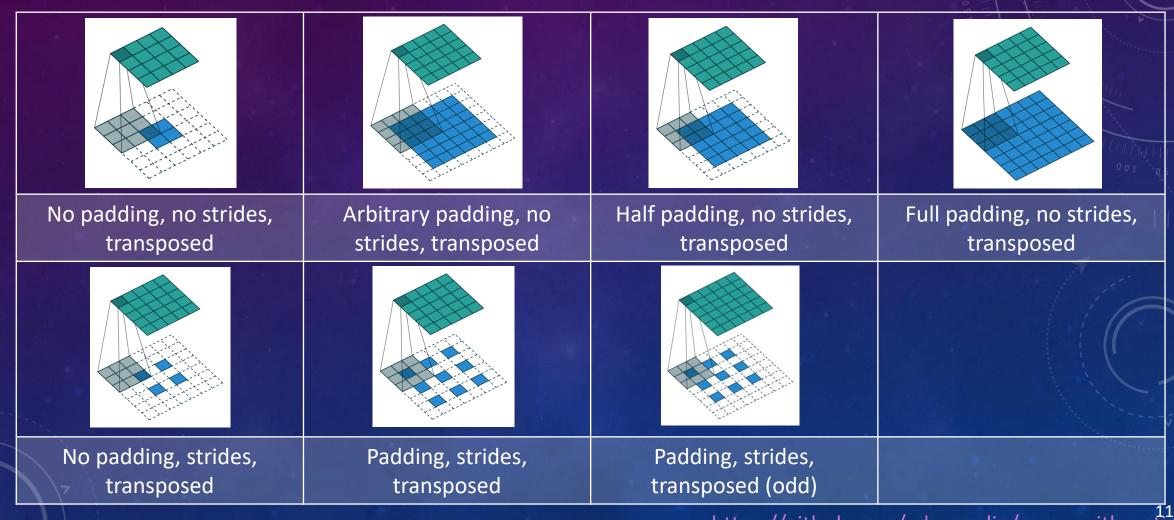
It is equivalent to convolving a 3x3 kernel over a 3x3 input (with 1 zero inserted between inputs) padded with a 1x1 border of zeros using unit strides (i.e., i'=3, $\tilde{i}'=5$, k'=k, s'=1 and p'=1)

Relationship 13. A convolution described k, s and p whose input size i is such that i + 2p - k is mutiple of s has an associated transposed convolution described by i', k' = k, s' = 1 and p' = k - p - 1, where i' is the Size of the streched input obtained by adding s - 1 zeros between each input unit, and tis ouput size is

$$o' = s(i'-1) + k - 2p$$

Transposed Convolution Animations

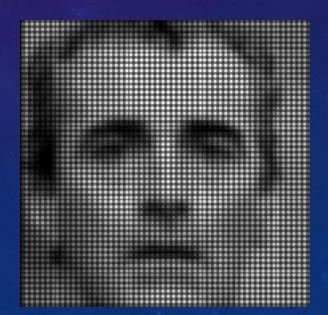
N.B.: Blue maps are inputs, and cyan maps are outputs.



Example 3-1: Convolution and Fractionally Strided Convolution

- Please download the "3-1_Convolution and Deconvolution.zip" from the Moodle.
- Upload the "sample.jpg" to the Google Colab.
- Run the codes and get the output images through the convolution and fractionally strided convolution.





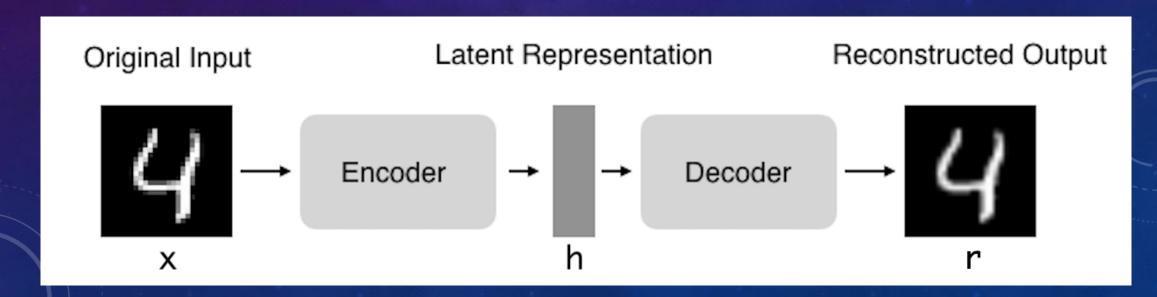
Exercise 3-1: Convolution and Fractionally Strided Convolution

- Please download the "3-1_Convolution and Deconvolution.zip" from the Moodle.
- Upload the "sample.jpg" to the Google Colab.
- Follow the example code and design a convolution layer and a fractionally strided convolution layer.
- Run the codes and get the output images through the designed convolution and fractionally strided convolution.

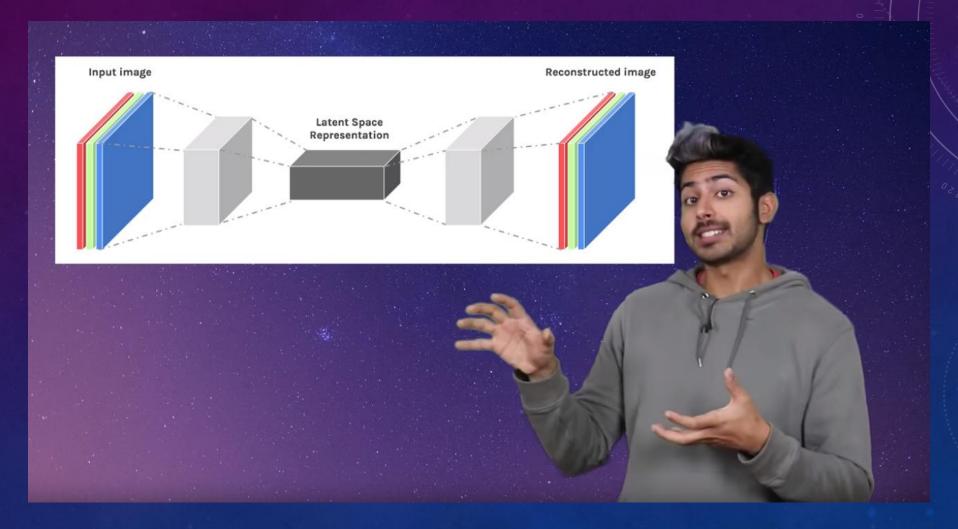
Please write down your results and codes in MS Word, then upload to the Moodle.

Autoencoders (AE)

- Autoencoders (AE) are neural networks that aims to copy their inputs to their outputs. They
 work by compressing the input into a latent-space representation, and then reconstructing the
 output from this representation.
 - Encoder: Compresses the input into a latent-space representation. It can be represented by an encoding function h = f(x).
 - Decoder: Reconstruct the input from the latent space representation. It can be represented by a decoding function r = g(h).

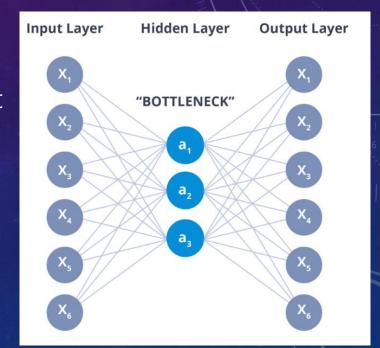


Autoencoder Explained



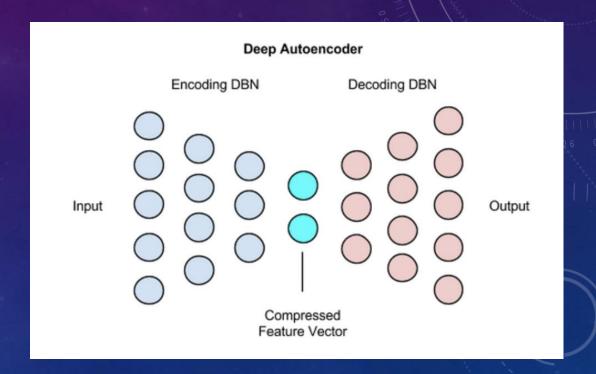
Bottleneck

- The layer between the encoder and decoder, ie. the code is also known as **Bottleneck**. This is a well-designed approach to decide which aspects of observed data are relevant information and what aspects can be discarded.
- It does this by balancing two criteria:
 - Compactness of representation, measured as the compressibility.
 - It retains some behaviourally relevant variables from the input.



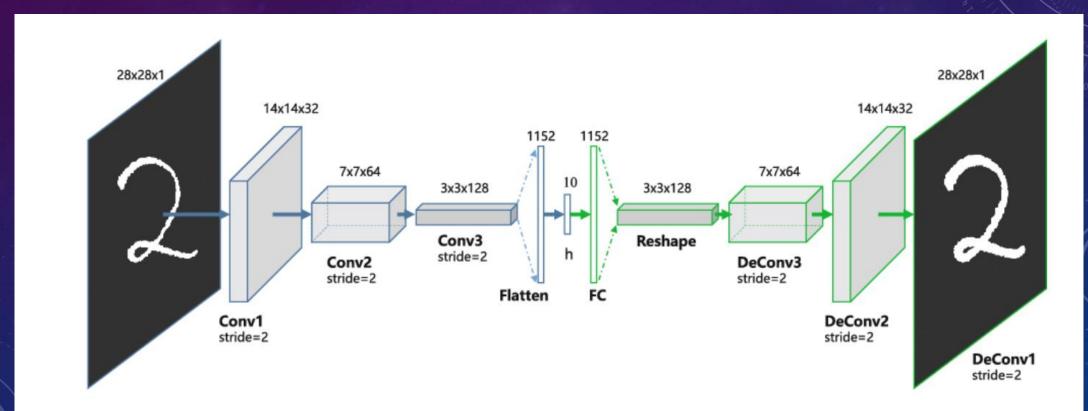
Deep Autoencoders

- The extension of the simple Autoencoder is the Deep Autoencoder. Deeper layers of the Deep Autoencoder tend to learn even higher-order features.
- A deep autoencoder is composed of two, symmetrical deep-belief networks-
 - First four or five shallow layers representing the encoding half of the net.
 - The second set of four or five layers that make up the decoding half.



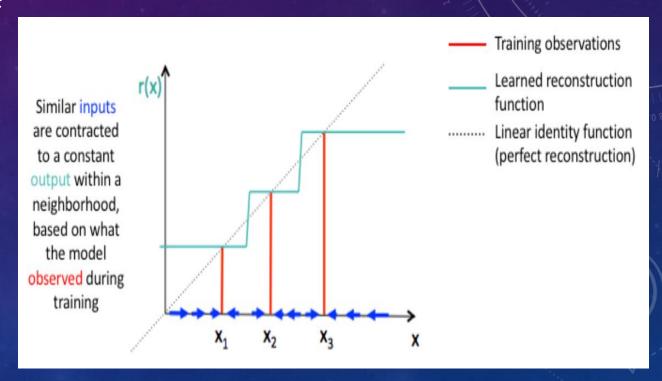
Convolution Autoencoders

Convolutional Autoencoders use the convolution operator to exploit this observation.
 They learn to encode the input in a set of simple signals and then try to reconstruct the input from them, modify the geometry or the reflectance of the image.



Contractive Autoencoders

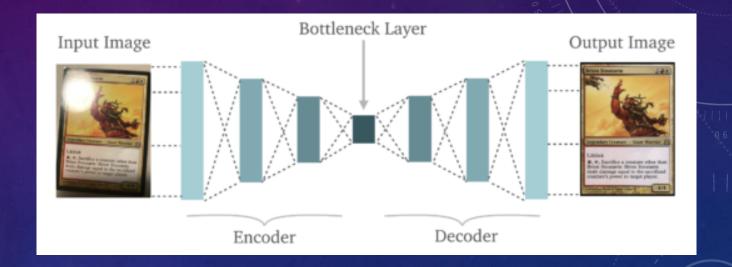
- A contractive autoencoder is an unsupervised deep learning technique that helps a neural network encode unlabeled training data.
- This is accomplished by constructing a loss term which penalizes large derivatives of our hidden layer activations with respect to the input training examples, essentially penalizing instances where a small change in the input leads to a large change in the encoding space.



Applications of Autoencoders

Feature variation

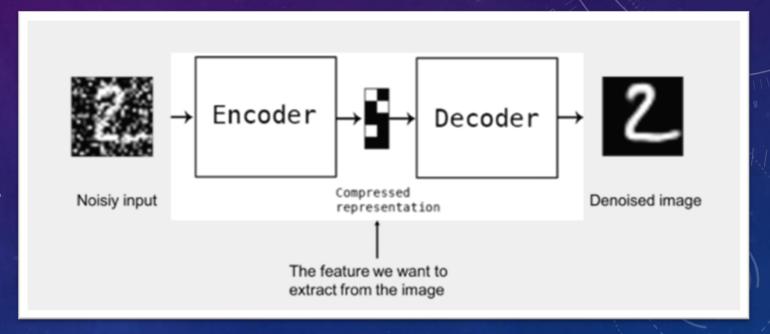
It extracts only the required features of an image and generates the output by removing any noise or unnecessary interruption.



Applications of Autoencoders

Denoising Image

The input seen by the autoencoder is not the raw input but a stochastically corrupted version. A denoising autoencoder is thus trained to reconstruct the original input from the noisy version.



Exercise 3.2: Autoencoder

- Please download the "exercise4.1_Autoencoder.ipynb" on Moodle.
 - Train the autoencoder and compare the images that reconstruct from the different epoch.
- Change the encoder and decoder to the below architecture and compare the difference. Please copy your results and code and paste to a MS Word, then upload to Moodle.



Define the hyper-parameter and load the training data

```
num_epochs = 10
batch_size = 32
learning_rate = 1e-3

Define the hyperparameter

img_transform = transforms.Compose([transforms.ToTensor()])

Download the Mnist dataset to the folder './data' dataset = torchvision.datasets.FashionMNIST(root= ./data', train=True, download=True, transform=img_transform)
dataloader = torch.utils.data.DataLoader(dataset, batch_size=batch_size, shuffle=False)
```

Define the model architecture

```
class autoencoder(nn.Module):
  def __init__(self):
    super(autoencoder, self).__ init__()
    self.encoder = nn.Sequential(
       nn.Linear(28 * 28, 128),
       nn.ReLU(True),
       nn.Linear(128, 64).
       nn.ReLU(True),
       nn.Linear(64, 12),
       nn.ReLU(True),
       nn.Linear(12, 3))
    self.decoder = nn.Sequential(
       nn.Linear(3, 12),
       nn.ReLU(True),
       nn.Linear(12, 64),
       nn.ReLU(True),
       nn.Linear(64, 128),
       nn.ReLU(True), nn.Linear(128, 28 * 28), nn.Tanh())
```

```
def forward(self, x):
    x = self.encoder(x)
    x = self.decoder(x)
    return x
```

```
autoencoder(
 (encoder): Sequential(
   (0): Linear(in features=784, out features=128, bias=True)
   (1): ReLU(inplace=True)
   (2): Linear(in_features=128, out_features=64, bias=True)
   (3): ReLU(inplace=True)
   (4): Linear(in features=64, out features=12, bias=True)
   (5): ReLU(inplace=True)
    (6): Linear(in features=12, out features=3, bias=True)
 (decoder): Sequential(
   (0): Linear(in_features=3, out_features=12, bias=True)
   (1): ReLU(inplace=True)
   (2): Linear(in features=12, out features=64, bias=True)
   (3): ReLU(inplace=True)
   (4): Linear(in features=64, out features=128, bias=True)
   (5): ReLU(inplace=True)
   (6): Linear(in features=128, out features=784, bias=True)
   (7): Tanh()
```

Define the model architecture

```
class autoencoder(nn.Module):
  def __init__(self):
    super(autoencoder, self).__init__()
    self.encoder = nn.Sequential(
       nn.Linear(28 * 28, 128),
       nn.ReLU(True),
       nn.Linear(128, 64),
       nn.ReLU(True),
       nn.Linear(64, 12),
       nn.ReLU(True),
       nn.Linear(12, 3))
```

Define the encoder

```
Define the decoder
```

```
self.decoder = nn.Sequential(
       nn.Linear(3, 12),
       nn.ReLU(True),
       nn.Linear(12, 64),
       nn.ReLU(True),
       nn.Linear(64, 128),
       nn.ReLU(True),
       nn.Linear(128, 28 * 28),
nn.Tanh())
def forward(self, x):
    x = self.encoder(x)
    x = self.decoder(x)
    return x
```

Define the model architecture

```
autoencoder(
  (encoder): Sequential(
    (0): Linear(in features=784, out features=128, bias=True)
    (1): ReLU(inplace=True)
    (2): Linear(in features=128, out features=64, bias=True)
    (3): ReLU(inplace=True)
    (4): Linear(in_features=64, out_features=12, bias=True)
    (5): ReLU(inplace=True)
    (6): Linear(in features=12, out features=3, bias=True)
  (decoder): Sequential(
    (0): Linear(in features=3, out features=12, bias=True)
    (1): ReLU(inplace=True)
    (2): Linear(in_features=12, out_features=64, bias=True)
    (3): ReLU(inplace=True)
    (4): Linear(in features=64, out features=128, bias=True)
    (5): ReLU(inplace=True)
    (6): Linear(in features=128, out features=784, bias=True)
    (7): Tanh()
```

Use the Mean Square Error as the loss function

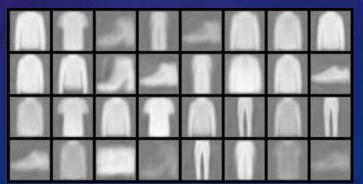
```
criterion = nn.MSELoss()
optimizer = torch.optim.Adam(model.parameters(), lr=learning_rate, weight_decay=1e-5)
```

```
if epoch % 1 == 0:
    img = to_img(img.cpu().data)
    save_image(img, './AE_img/input_{}.png'.format(epoch))
    pic = to_img(output.cpu().data)
    save_image(pic, './AE_img/output_{}.png'.format(epoch))
    Save the input images
    Save the reconstruction images
```

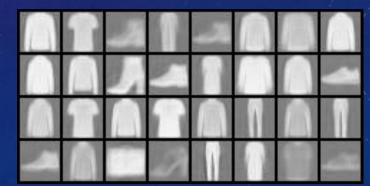




Epoch 1 : reconstruction images



Epoch 10: reconstruction images



Lecture 13 | Generative Models



https://youtu.be/5WoltGTWV54 [1:17:40]

Lecture 13 | Generative Models

- OUTLINE:
- 1:12 Supervised learning
- 5:33 Generative models
- 11:00 PixelRNN and PixelCNN
- 19:36 Traditional and variational autoencoders (VAEs)
- 50:00 Generative adversarial networks (GANs)

Variational Autoencoders

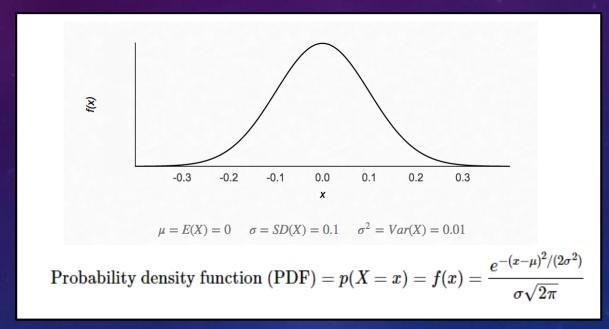


https://youtu.be/9zKuYvjFFS8 [15:05]

Variational Autoencoders

- Variational autoencoders use gaussian models to generate images.
- There are some backgrounds we need to know:
 - Gaussian distribution
 - Autoencoders
 - KL-divergence

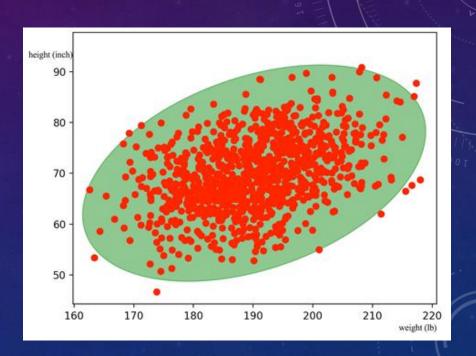
• In the following diagram, we assume the probability of X equal to a certain value x, p(X=x), follows a gaussian distribution:



• We can sample data using the PDF above. We use the following notation for sample data using a gaussian distribution with mean μ and standard deviation σ .

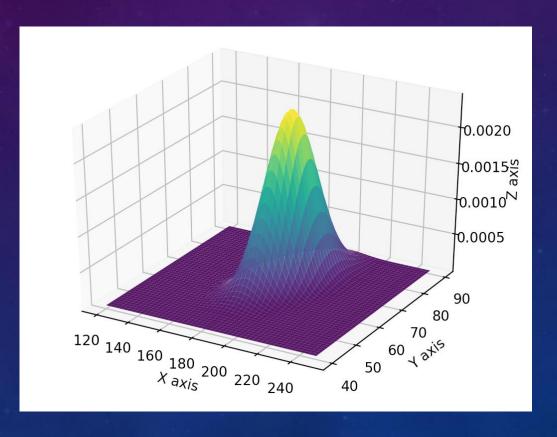
$$x \sim N(\mu, \sigma)$$

- In the example above, mean: μ =0, standard deviation: σ =0.1:
 - In many real world examples, the data sample distribution follows a gaussian distribution.
- Now, we generalize it with multiple variables. For example, we want to model the relationship between the body height and the body weight for San Francisco residents.
- We collect the information from 1000 adult residents and plot the data below with each red dot represents 1 person:



• We can plot the corresponding probability density function in 3D:

PDF = probability(height = h, weight = w)



We can model such a probability density function using a gaussian distribution function. The PDF with p variables is:

$$z = \begin{pmatrix} X_1 \\ \vdots \\ X_p \end{pmatrix}, f(x) = \frac{1}{(2\pi)^{p/2} |\sum|^{1/2}} \exp\{-\frac{1}{2}(x-\mu)^T \sum^{-1} (x-\mu)\}$$

with covariance matrix Σ :

$$\sum = egin{pmatrix} E[(x_1 - \mu_1)(x_1 - \mu_1)] & E[(x_1 - \mu_1)(x_2 - \mu_2)] & \dots & E[(x_1 - \mu_1)(x_p - \mu_p)] \\ E[(x_2 - \mu_2)(x_1 - \mu_1)] & E[(x_2 - \mu_2)(x_2 - \mu_2)] & \dots & E[(x_2 - \mu_2)(x_p - \mu_p)] \\ & dots & & dots & \ddots & dots \\ E[(x_p - \mu_p)(x_1 - \mu_1)] & E[(x_p - \mu_p)(x_2 - \mu_2)] & \dots & E[(x_n - \mu_p)(x_p - \mu_p)] \end{pmatrix}$$

Gaussian Distribution

• The notation for sampling *x* is:

$$x \sim \mathcal{N}\Big(\mu, \sum\Big) \ x = egin{pmatrix} x_1 \ dots \ x_p \end{pmatrix} \sim \mathcal{N}\Big(\mu, \sum\Big) = \mathcal{N}\left(egin{pmatrix} \mu_1 \ dots \ \mu_p \end{pmatrix}, \sum
ight)$$

Let's go back to our weight and height example to illustrate it

$$x = \begin{pmatrix} weight \\ height \end{pmatrix}$$

• From the data, we compute the mean weight is 190 lb and mean height is 70 inches:

$$\mu = \binom{190}{70}$$

Gaussian Distribution

• For the covariance matrix Σ , here, we illustrate how to compute one of the element E_{21}

$$E_{21} = E[(x_2 - \mu_2)(x_1 - \mu_1)] = E[(x_{height} - 70)(x_{weight} - 190)]$$

 which E is the expected value. Let say we have only 2 datapoints (200 lb, 80 inches) and (180 lb, 60 inches)

$$E_{21} = E[(x_{height} - 70)(x_{weight} - 190)]$$

$$= \frac{1}{2}((80 - 70) \times (200 - 190) + (60 - 70) \times (180 - 190))$$

• After computing all 1000 data, here is the value of Σ :

$$\sum = {100 \ 25 \choose 25 \ 50}, \qquad x \sim N\left({190 \choose 70}, {100 \choose 50}\right)$$

Gaussian Distribution

• E_{21} measures the co-relationship between variables x_2 and x_1 . Positive values means both are positively related. With not surprise, E_{21} is positive because weight increases with height. If two variables are independent of each other, it should be 0 like:

$$\sum = \begin{pmatrix} 100 & 0 \\ 0 & 50 \end{pmatrix}$$

We will simplify the gaussian distribution notation here as:

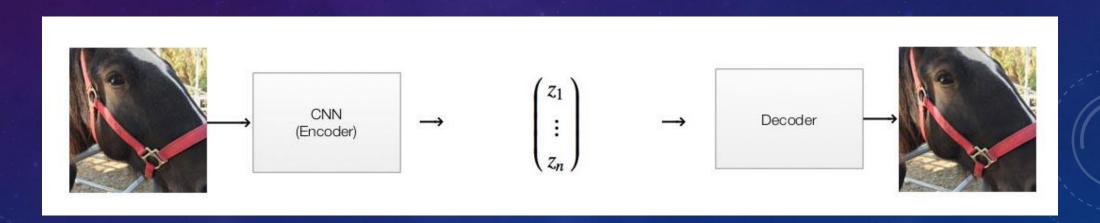
$$x \sim N\left(\binom{190}{70}, \binom{100}{50} \right)$$

Autoencoders

- In an autoencoders, we use a deep network to map the input image (for example 256x256 pixels = 256x256 = 65536 dimension) to a lower dimension latent variables (latent vector say 100-D vector: $(x_1, x_2, \dots x_{100})$).
- We use another deep network to decode the latent variables to restore the image.
 We train both encoder and decoder network to minimize the difference between the original image and the decoded image.
- By forcing the image to a lower dimension, we hope the network learns to encode the image by extracting core features.

Autoencoders

- For example, we enter a 256x256 image to the encoder, we use a CNN to encode the image to 20-D latent variables $(x_1, x_2, ..., x_{20}) = (0.1, 0, ..., -0.05)$.
- We use another network to decode the latent variables into a 256x256 image.
- We use backpropagation with cost function comparing the decoded and input image to train both encoding and decoding network



 For VAEs, we replace the middle part with a stochastic model using a gaussian distribution. Let's get into an example to demonstrate the flow:

$$\xrightarrow{\text{CNN}} \rightarrow \mathcal{N}\left(\begin{pmatrix} \mu_1 \\ \vdots \\ \mu_n \end{pmatrix}, \begin{pmatrix} \sigma_1 \\ \vdots \\ \sigma_n \end{pmatrix}\right) \rightarrow \begin{pmatrix} z_1 \\ \vdots \\ z_n \end{pmatrix} \sim \mathcal{N}\left(\begin{pmatrix} \mu_1 \\ \vdots \\ \mu_n \end{pmatrix}, \begin{pmatrix} \sigma_1 \\ \vdots \\ \sigma_n \end{pmatrix}\right) \rightarrow \text{Decoder}$$

For a variation autoencoder, we replace the middle part with 2 separate steps. VAE
does not generate the latent vector directly.

- It generates 100 Gaussian distributions each represented by a mean (μ_i) and a standard deviation (σ_i) .
- Then it samples a latent vector, say (0.1, 0.03, ..., -0.01), from these distributions.
- For example, if element x_i of the latent vector has $\mu_i = 0.1$ and $\sigma_i = 0.5$. We randomly select x_i with probability based on this Gaussian distribution:

$$egin{align} p(X=x_i) &= rac{e^{-(x_i-\mu_i)^2/(2\sigma_i^2)}}{\sigma_i\sqrt{2\pi}} \ z &= egin{pmatrix} z_1 \ dots \ z_{20} \end{pmatrix} \sim \mathcal{N} \left(egin{pmatrix} \mu_1 \ dots \ \mu_{20} \end{pmatrix}, egin{pmatrix} \sigma_1 \ dots \ \sigma_{20} \end{pmatrix}
ight) \end{aligned}$$

• Say, the encoder generates $\mu = (0, -0.01, ..., 0.2)$ and $\sigma = (0.05, 0.01, ..., 0.02)$. We can sample a value from this distribution:

$$N\left(\begin{pmatrix}0\\-0.01\\\vdots\\0.2\end{pmatrix},\begin{pmatrix}0.05\\0.01\\\vdots\\0.02\end{pmatrix}\right)$$

with the latent variables as (say) :

$$z = \begin{pmatrix} z_1 \\ \vdots \\ z_{20} \end{pmatrix} = \begin{pmatrix} 0.03 \\ -0.015 \\ \vdots \\ 0.197 \end{pmatrix}$$

- The autoencoder in the previous section is very hard to train with not much guarantee that the network is generalize enough to make good predictions.(We say the network simply memorize the training data.)
- In VAEs, we add a constrain to make sure:
 - 1. The latent variable are relative independent of each other, i.e. the 20 variables are relatively independent of each other (not co-related). This maximizes what a 20-D latent vectors can represent.
 - 2. Latent variables z which are similar in values should generate similar looking images. This is a good indication that the network is not trying to memorize individual image.

- To achieve this, we want the gaussian distribution model generated by the encoder to be as close to a normal gaussian distribution function.
- We penalize the cost function if the gaussian function is deviate from a normal distribution. This is very similar to the L_2 regularization in a fully connected network in avoiding overfitting.

$$z \sim N(0,1) = normal distribution$$

- In a normal gaussian distribution, the covariance E_{ij} is 0 for $i \neq j$. That is the latent variables are independent of each other.
- If the distribution is normalize, the distance between different z will be a good measure of its similarity.
- With sampling and the gaussian distribution, we encourage the network to have similar value of z for similar images.

• In VAE, we want to model the data distribution p(x) with an encoder $q_{\emptyset}(z|x)$, a decoder $p_{\theta}(z|x)$ and a latent variable model p(z) through the VAE objective function:

$$\log p(x) \approx E_q[log p_{\theta}(x|z)] - D_{KL}[q_{\emptyset}(z|x)||p(z)]$$

 To draw this conclusion, we start with the KL divergence which measures the difference of 2 distributions. By definition, KL divergence is defined as:

$$D_{KL}(q||p) = \sum_{x} q(x) \log \left(\frac{q(x)}{p(x)}\right) = E_q[\log(q(x)) - \log(p(x))]$$

Apply it with:

$$D_{KL}[(q||p)||p(z|x)] = E[logq(z|x) - logp(z|x)]$$

• Let $q_{\lambda}(z|x)$ be the distribution of z predicted by our encoder deep network. We want it to match the true distribution p(z|x). We want the distribution approximated by the deep network has little divergence from the true distribution. i.e. we want to optimize λ with the smallest KL divergence.

$$D_{KL}[(q_{\lambda}(z|x)||p(z|x)] = E[logq_{\lambda}(z|x) - logp(z|x)]$$

Apply:

$$\begin{split} p(z|x) &= \frac{p(x|z)p(z)}{p(x)} \\ D_{KL}[q_{\lambda}(z|x) \| p(z|x)] &= \mathbb{E}_q[\log q_{\lambda}(z|x) - \log \frac{p(x|z)p(z)}{p(x)}] \\ &= \mathbb{E}_q[\log q_{\lambda}(z|x) - \log p(x|z) - \log p(z) + \log p(x)] \\ &= \mathbb{E}_q[\log q_{\lambda}(z|x) - \log p(x|z) - \log p(z)] + \log p(x) \\ D_{KL}[q_{\lambda}(z|x) \| p(z|x)] - \log p(x) &= \mathbb{E}_q[\log q_{\lambda}(z|x) - \log p(x|z) - \log p(z)] \\ \log p(x) - D_{KL}[q_{\lambda}(z|x) \| p(z|x)] &= \mathbb{E}_q[\log p(x|z) - (\log q_{\lambda}(z|x) - \log p(z))] \\ &= \mathbb{E}_q[\log p(x|z)] - \mathbb{E}_q[\log q_{\lambda}(z|x) - \log p(z)] \\ &= \mathbb{E}_q[\log p(x|z)] - D_{KL}[q_{\lambda}(z|x) \| p(z)] \end{split}$$

• Define the term ELBO (Evidence lower bound) as:

$$ELBO(\lambda) = E_q[logp(x|z)] - D_{KL}[q_{\lambda}(z|x)||p(z)]$$
$$logp(x) - D_{KL}[q_{\lambda}(z|x)||p(z|x)] = ELBO(\lambda)$$

We call ELBO the evidence lower bound because:

$$log p(x) - D_{KL}[q_{\lambda}(z|x)||p(z|x)] = ELBO(\lambda)$$

 $logp(x) \ge ELBO(\lambda)$ since KL is always positive

Here, we define our VAE objective function

$$log p(x) - D_{KL}[q_{\lambda}(z|x)||p(z|x)] = E_q[log p(x|z)] - D_{KL}[q_{\lambda}(z|x)||p(z)]$$

- Instead of the distribution p(x), we can model the data x with $\log p(x)$. With the error term, $D_{KL}[q_{\lambda}(z|x)||p(z|x)]$, we can establish a lower bound ELBO for $\log p(x)$ which in practice is good enough in modeling the data distribution.
- In the VAE objective function, maximize our model probability $\log p(x)$. is the same as maximize $\log p(x|z)$. while minimize the divergence of $D_{KL}[q_{\lambda}(z|x)||p(z)]$.

• Maximizing $\log p(x|z)$ can be done by building a decoder network and maximize its likelihood. So with an encoder $q_{\emptyset}(z|x)$, a decoder $p_{\theta}(z|x)$, our objective become optimizing:

$$ELBO(\theta, \emptyset) = E_{q\theta(z|x)} \left[\log \left(p_{\theta}(x_i|z) \right) \right] - D_{KL} \left[q_{\emptyset}(z|x) || p(z) \right]$$

• We can apply a constrain to p(z) such that we can evaluate $D_{KL}[q_{\emptyset}(z|x)||p(z)]$ easily. In VAE, we use p(z) = N(0,1). For optimal solution, we want $q_{\emptyset}(z|x)$ to be as close as N(0,1).

• In VAE, we model $q_{\emptyset}(z|x)$ as $N(\mu, \Sigma)$

$$\begin{split} D_{KL}[q_{\phi}(z|x) \| p(z)] &= D_{KL}[N(\mu, \Sigma) \| N(0, 1)] \\ &= \frac{1}{2} \left(\operatorname{tr}(\Sigma) + \mu^T \mu - k - \log \det(\Sigma) \right) \\ &= \frac{1}{2} \left(\sum_k \Sigma + \sum_k \mu^2 - \sum_k 1 - \log \prod_k \Sigma \right) \\ &= \frac{1}{2} \left(\sum_k \Sigma(X) + \sum_k \mu^2(X) - \sum_k 1 - \sum_k \log \Sigma(X) \right) \\ &= \frac{1}{2} \sum_k (\Sigma + \mu^2 - 1 - \log \Sigma) \end{split}$$

Example 3.3 : VAE

Define the model architecture

Define the latent vector dimension

```
class VAE(nn.Module):
  def __init__(self, image_channels=1, h_dim=1600, z_dim=20):
    super(VAE, self).__init__()
    self.conv1 = nn.Conv2d(image_channels, 32, kernel_size=4, stride=2)
    self.conv2 = nn.Conv2d(32, 64, kernel size=4, stride=2)
    self.fc1 = nn.Linear(h_dim, z_dim)
    self.fc2 = nn.Linear(h_dim, z_dim)
    self.fc3 = nn.Linear(z_dim, h_dim)
    self.deconv2 = nn.ConvTranspose2d(64, 32, kernel_size=5, stride=2)
    self.deconv1 = nn.ConvTranspose2d(32, image_channels, kernel_size=4, stride=2)
```

Example 3.3 : VAE

```
if epoch % 1 == 0:
    save = img.cpu().data
    save_image(img, './vae_img/input_{}.png'.format(epoch))
    save = recon_batch.cpu().data
    save_image(save, './vae_img/output_{}.png'.format(epoch))
    Save the input images
    Save the reconstruction images
```

