Deep Learning PhD course HW#3

Muhammad Osama

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1 Ex1

Assuming $x^{(i)}$ is d dimensional and is either 0 or 1, we use the following data model conditioned on the latent variable z

$$p_{\theta}(\boldsymbol{x}^{(i)}|\boldsymbol{z}) = \prod_{k=1}^{d} Bernoulli(\theta_k)$$
 (1)

where $\theta = \{\theta_1, \dots, \theta_d\}$. For MNIST, d = 784. Hence given z, we take every pixel to be a Bernoulli random variable. Ideally, the pixel values lie in the interval [0, 1]. However, for simplicity we use the above approximation.

2 Ex2

The reason we cannot work with eq. (5) directly is because it is difficult to evaluate the expectation $\mathbb{E}_{q_{\boldsymbol{\theta}}(\boldsymbol{z}|\boldsymbol{x}^{(i)})} \left[\log p_{\boldsymbol{\theta}}(\boldsymbol{x}^{(i)}|\boldsymbol{z})\right]$ analytically.

3 Ex 3

$$q_{\phi}(\boldsymbol{z}|\boldsymbol{x}^{(i)}) = \mathcal{N}(\boldsymbol{\mu}^{(i)}, diag(\boldsymbol{\sigma}^{2(i)})) \qquad p(\boldsymbol{z}) = \mathcal{N}(0, \mathbf{I})$$

Let $\Sigma = diag(\sigma^{2(i)})$, hence the KL divergence is

$$\Delta_{KL}(q_{\phi}(\boldsymbol{z}|\boldsymbol{x}^{(i)})||p(\boldsymbol{z})) = \mathbb{E}_{q_{\phi}(\cdot)}\left[-\frac{(\boldsymbol{z}-\boldsymbol{\mu}^{(i)})^{\top} \Sigma^{-1}(\boldsymbol{z}-\boldsymbol{\mu}^{(i)})}{2}\right] + \mathbb{E}_{q_{\phi}(\cdot)}\left[\frac{\boldsymbol{z}^{\top} \boldsymbol{z}}{2}\right] - \frac{1}{2} \sum_{j=1}^{K} \log \sigma_{j}^{2(i)},$$

$$(2)$$

where

$$\mathbb{E}_{q_{\phi}(\cdot)}\left[-\frac{(\boldsymbol{z}-\boldsymbol{\mu}^{(i)})^{\top}\Sigma^{-1}(\boldsymbol{z}-\boldsymbol{\mu}^{(i)})}{2}\right] = \frac{1}{2}\sum_{j=1}^{K}(1),$$

$$\mathbb{E}_{q_{\phi}(\cdot)}\left[\frac{\boldsymbol{z}^{\top}\boldsymbol{z}}{2}\right] = \sum_{j=1}^{K}\sigma_{j}^{2(i)} + \mu_{j}^{2(i)}.$$

Hence,

$$\Delta_{KL}(q_{\phi}(\boldsymbol{z}|\boldsymbol{x}^{(i)})||p(\boldsymbol{z})) = -\frac{1}{2} \left[\sum_{i=1}^{K} 1 + \log \sigma_{j}^{2(i)} - \sigma_{j}^{2(i)} - \mu_{j}^{2(i)} \right]$$
(3)

Moreover,

$$g_{\phi}(\boldsymbol{\epsilon}^{(i,l)}, \boldsymbol{x}^{(i)}) = \boldsymbol{\mu}^{(i)} + \boldsymbol{\sigma}^{(i)} \odot \boldsymbol{\epsilon}^{(i,l)}$$

where

$$\boldsymbol{\epsilon}^{(i,l)} \sim \mathcal{N}(0, \mathbf{I})$$

4 Ex 4

The model summaries are given in figures 1 and 5 for latent dimensions of k=3 and k=20 respectively. For sampling from the generative network, a k dimensional vector \mathbf{z}_s was picked randomly in space \mathbb{R}^k . I looked at the mean and variance of the encoder distribution $q_{\boldsymbol{\phi}}(\mathbf{z}|\mathbf{x})$ for the test data to decide on what would be reasonable values for the elements of \mathbf{z}_s . The 784 probability parameters of the Bernoulli distribution $p_{\boldsymbol{\theta}}(\mathbf{z}|\mathbf{z})$ corresponding to this latent variable were obtained by using $\mathbf{p}_B = decoder.predict(\mathbf{z}_s)$. Then we simply independently sampled 784 random variables with these probabilities from the Bernoulli distribution i.e. the sample is given by $\mathbf{x}_s = Bernoulli(\mathbf{p}_B)$.

4.1 Number of latent variables K=3

- (a) Loss on test data: KL loss = 7.89, Reconstruction error = 131.80
- (b) Some samples x_s from the generative network for k=2 are shown in figure 2 (c) Some original test examples and their reconstructions are shown in figures 3 and 4.

4.2 Number of Latent variables K = 20

- (a) Loss on test data: KL loss = 25.23, Reconstruction error = 78.33
- (b) Some samples x_s from the generative network for k=20 are shown in figure 6. (c) Some original test examples and their reconstructions are shown in figures 7 and 8.

We observe that as the dimension of the latent variable increases from k=3

In [553]: vae.summary()

Layer (type)	Output Shape	Param #
encoder_input (InputLayer)	(None, 784)	0
encoder (Model)	[(None, 2), (None, 2), (N	403972
decoder (Model)	(None, 784)	403728

Total params: 807,700 Trainable params: 807,700 Non-trainable params: 0

Figure 1

(a) (b) (c)

Figure 2: Samples \boldsymbol{x}_s for latent variable (a) $\boldsymbol{z}_s=[-2,4]$ (b) $\boldsymbol{z}_s=[-0.2,0.4]$ (c) $\boldsymbol{z}_s=[0.1,0.1]$

to k=20, the overall loss decreases. It is interesting to see that the KL loss actually increases with dimension so that the posterior $q_{\phi}(\boldsymbol{z}|\boldsymbol{x})$ is less close to the prior $p(\boldsymbol{z})$. However, the reconstruction error decreases with dimension.

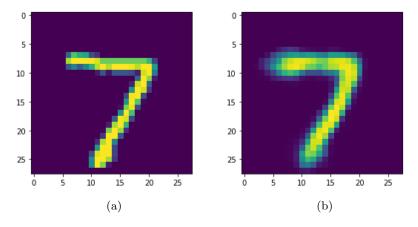


Figure 3: (a) Original (b) Reconstructed

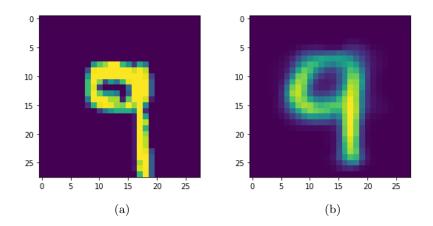


Figure 4: (a) Original (b) Reconstructed

Layer (type)	Output Shape	Param #
encoder_input (InputLayer)	(None, 784)	0
encoder (Model)	[(None, 20), (None, 20),	422440
decoder (Model)	(None, 784)	412944
Total params: 835,384 Trainable params: 835,384 Non-trainable params: 0		

Figure 5

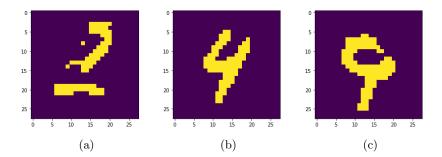


Figure 6: Samples \boldsymbol{x}_s

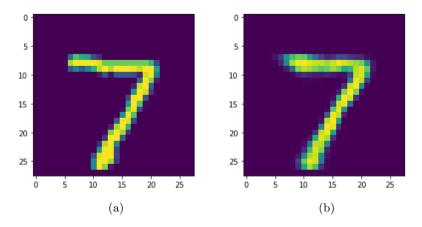


Figure 7: (a) Original (b) Reconstructed

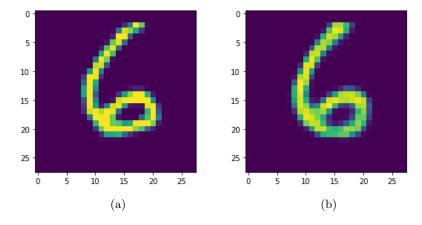


Figure 8: (a) Original (b) Reconstructed

Appendix

```
from keras import backend as K
from keras.losses import
   binary_crossentropy
from keras.datasets import mnist
from keras.layers import Lambda, Input,
from keras.models import Model
import tensorflow as tf
import numpy as np
import matplotlib.pyplot as plt
#sampling function-reparametrization
   trick
def sampling (args):
z_{mean}, z_{log}var = args
z_std = K. sqrt(K. exp(z_log_var))
batch = K.shape(z_mean)[0]
\dim = K. \operatorname{int\_shape} (z_{-mean}) [1]
epsl = K.random_normal(shape=(batch,dim))
return z_mean + z_std*epsl
#define losses
def reg_loss(args):
z_{mean}, z_{log_var} = args
#loss per datapoint
kl_loss = 1 + z_log_var - K.exp(z_log_var)
   ) - K. square (z_mean)
kl_loss = K.sum(kl_loss, axis = -1)
kl_{-}loss *= -0.5;
#average over all datapoints
kl_loss = K.mean(kl_loss)
return kl_loss
def recn_loss (varargs):
inputs, outputs = varargs
#loss per data point assuming nos. of MC
   samples r=1
```

```
\#recons_{loss} = K.sum(inputs * K.log(
   outputs) + (1-inputs)*K.log(1-outputs)
    , axis=-1)
recons_loss = binary_crossentropy(inputs,
    outputs)
recons_loss *= original_dim
#average over all data points
recons_{loss} = K.mean(recons_{loss})
return recons_loss
# MNIST dataset
(x_{train}, y_{train}), (x_{test}, y_{test}) =
   mnist.load_data()
image\_size = x\_train.shape[1]
original_dim = image_size * image_size
x_{train} = np.reshape(x_{train}, [-1,
    original_dim])
x_{test} = np.reshape(x_{test}, [-1,
   original_dim])
x_train = x_train.astype('float32') / 255
x_{test} = x_{test} \cdot astype('float32') / 255
# network parameters
input_shape = (original_dim,)
intermediate_dim = 512
batch\_size = 128
latent_dim = 3
epochs = 20
# build encoder model
inputs = Input(shape=input_shape, name='
   encoder_input')
x = Dense(intermediate_dim, activation='
   relu') (inputs)
z_mean = Dense(latent_dim , name='z_mean')
z_log_var = Dense(latent_dim, name='
   z_{\log_{v}}(x)
z = Lambda(sampling, output_shape=(
   latent_dim ,) , name='z') ([z_mean ,
   z \log_v var])
encoder = Model(inputs, [z_mean,
   z_log_var, z], name='encoder')
# build decoder model
```

```
latent_inputs = Input(shape=(latent_dim,)
    , name='z_sampling')
x = Dense(intermediate_dim, activation='
   relu') (latent_inputs)
outputs = Dense(original_dim, activation
   ='sigmoid')(x)
decoder = Model(latent_inputs, outputs,
   name='decoder')
outputs = decoder (encoder (inputs) [2])
vae = Model(inputs, outputs, name='vae')
vae\_loss = reg\_loss([z\_mean, z\_log\_var])
   + recn_loss([inputs, outputs])
vae.add_loss(vae_loss)
vae.compile(optimizer='adam', metrics =
   ['reg_loss', 'recn_loss'])
#fit
history = vae. fit (x_train, epochs=epochs,
   batch_size=batch_size, validation_data
   =(x_{test}, None)
#predict
ztest_mean, ztest_log_var, ztest =
   encoder.predict(x_test)
x_pred = decoder.predict(ztest)
#Ex4.1
x_{pred} = tf.convert_{to_{tensor}}(x_{pred}, np.
   float32)
kl\_loss = reg\_loss([ztest\_mean],
    ztest_log_var])
recons_{loss} = recn_{loss} ([x_{test}, x_{pred}])
with tf. Session() as sess:
kl_loss = kl_loss.eval()
with tf. Session() as sess:
recons_loss = recons_loss.eval()
print(kl_loss)
print(recons_loss)
\#\text{Ex}4.2
def sample_generative_network(laten_var):
```

```
#probabilities of bernoulli distribution
prb = decoder.predict(laten_var)
#sample from bernoulli
xhat = np.random.binomial(1, prb)
xhat = np.reshape(xhat,(image_size,
   image_size))
plt.imshow(xhat)
\#ex4.3
x_pred = decoder.predict(ztest);
idx = 700
#original image
plt.imshow(np.reshape(x_test[idx,:],(
   image_size , image_size)))
plt.show()
#reconstructed image
plt.show()
plt.imshow(np.reshape(x_pred[idx,:],(
   image_size , image_size ) ) )
```