Solution template for the question 1.6-1.7. This template consists of following steps. Except the step 2, you don't need to modify it to answer the questions.

- 1. Initialize libraries
- 2. Insert the answers for the questions 1.1~1.5 below (this is the part you need to fill)
- 3. Define data loaders
- 4. Define VAE network architecture
- 5. Initialize the model and optimizer
- 6. Train the model
- 7. Save the model
- 8. Load the model
- 9. Evaluate the model with importance sampling

Initialize libraries

In [22]:

```
import math
from torchvision.datasets import utils
import torch.utils.data as data_utils
import torch
import os
import numpy as np
from torch import nn
from torch.nn.modules import upsampling
from torch.functional import F
from torch.optim import Adam
```

Insert the answers for the questions 1.1~1.5 below

```
def log_likelihood_bernoulli(mu, target):
    COMPLETE ME. DONT MODIFY THE PARAMETERS OF THE FUNCTION. Otherwise, tests mi
ght fail.
    *** note. ***
    :param mu: (FloatTensor) - shape: (batch size x input size) - The mean of Be
rnoulli random variables p(x=1).
    :param target: (FloatTensor) - shape: (batch size x input size) - Target sam
ples (binary values).
    :return: (FloatTensor) - shape: (batch size,) - log-likelihood of target sam
ples on the Bernoulli random variables.
    # init
    batch size = mu.size(0)
    mu = mu.view(batch size, -1)
    target = target.view(batch size, -1)
    loss = (target*torch.log(mu) + (1-target)*torch.log(1-mu))
    out = torch.sum(loss, dim=-1)
    return out
def log likelihood normal(mu, logvar, z):
    COMPLETE ME. DONT MODIFY THE PARAMETERS OF THE FUNCTION. Otherwise, tests mi
ght fail.
    *** note. ***
    :param mu: (FloatTensor) - shape: (batch size x input size) - The mean of No
rmal distributions.
    :param logvar: (FloatTensor) - shape: (batch size x input size) - The log va
riance of Normal distributions.
    :param z: (FloatTensor) - shape: (batch size x input size) - Target samples.
    :return: (FloatTensor) - shape: (batch_size,) - log probability of the sames
on the given Normal distributions.
    0.00
    # init
    batch size = mu.size(0)
    mu = mu.view(batch_size, -1)
    logvar = logvar.view(batch size, -1)
    z = z.view(batch_size, -1)
    out = torch.zeros(batch size,)
    for i in range(batch size):
        m = torch.distributions.multivariate normal.MultivariateNormal(mu[i], to
rch.diag(logvar[i].exp()))
        out[i] = m.log prob(z[i])
    return out
def log_mean_exp(y):
    COMPLETE ME. DONT MODIFY THE PARAMETERS OF THE FUNCTION. Otherwise, tests mi
ght fail.
    *** note. ***
```

```
:param y: (FloatTensor) - shape: (batch_size x sample_size) - Values to be e
valuated for log_mean_exp. For example log proababilies
    :return: (FloatTensor) - shape: (batch size,) - Output for log mean exp.
    # init
    batch size = y.size(0)
    sample size = y.size(1)
    out = torch.zeros(batch size,)
    for i in range(batch size):
        yi max = torch.max(y[i])
        yi = y[i] - yi max
        out[i] = torch.log(torch.mean(torch.exp(yi))) + yi max
    return out
def kl gaussian gaussian analytic(mu q, logvar q, mu p, logvar p):
    COMPLETE ME. DONT MODIFY THE PARAMETERS OF THE FUNCTION. Otherwise, tests mi
ght fail.
    *** note. ***
    :param mu q: (FloatTensor) - shape: (batch size x input size) - The mean of
 first distributions (Normal distributions).
    :param logvar q: (FloatTensor) - shape: (batch size x input size) - The log
 variance of first distributions (Normal distributions).
    :param mu p: (FloatTensor) - shape: (batch_size x input_size) - The mean of
 second distributions (Normal distributions).
    :param logvar p: (FloatTensor) - shape: (batch size x input size) - The log
 variance of second distributions (Normal distributions).
    :return: (FloatTensor) - shape: (batch size,) - kl-divergence of KL(q||p).
    0.00
    # init
    batch size = mu q.size(0)
    mu q = mu q.view(batch size, -1)
    logvar_q = logvar_q.view(batch size, -1)
    mu p = mu p.view(batch size, -1)
    logvar p = logvar p.view(batch size, -1)
    out = torch.ones(batch_size,).type(torch.FloatTensor)
    for i in range(batch_size):
        sigma0 = torch.diag(logvar_q[i].exp())
        sigma1 = torch.diag(logvar p[i].exp())
        mu0 = mu q[i]
        mu1 = mu p[i]
        ## find KL(N0||N1)
        ## https://en.wikipedia.org/wiki/Kullback%E2%80%93Leibler_divergence#Mul
tivariate normal distributions
        ## out[i] = (torch.trace(torch.matmul(torch.inverse(sigma1), sigma0)) +
 torch.matmul((mu1 - mu0), torch.matmul(torch.inverse(sigma1), (mu1 - mu0))) - l
en(mu1) + torch.log(torch.det(sigma1)/torch.det(sigma0)))/2.0
        dq = torch.distributions.multivariate_normal.MultivariateNormal(mu_q[i],
torch.diag(logvar q[i].exp()))
        dp = torch.distributions.multivariate normal.MultivariateNormal(mu p[i],
torch.diag(logvar p[i].exp()))
        out[i] = torch.distributions.kl.kl_divergence(dq, dp)
```

```
return out
def kl_gaussian_gaussian_mc(mu_q, logvar_q, mu_p, logvar_p, num_samples=1):
    COMPLETE ME. DONT MODIFY THE PARAMETERS OF THE FUNCTION. Otherwise, tests mi
ght fail.
    *** note. ***
    :param mu q: (FloatTensor) - shape: (batch size x input size) - The mean of
 first distributions (Normal distributions).
    :param logvar_q: (FloatTensor) - shape: (batch_size x input_size) - The log
 variance of first distributions (Normal distributions).
    :param mu p: (FloatTensor) - shape: (batch size x input size) - The mean of
 second distributions (Normal distributions).
    :param logvar p: (FloatTensor) - shape: (batch size x input size) - The log
 variance of second distributions (Normal distributions).
    :param num samples: (int) - shape: () - The number of sample for Monte Carlo
estimate for KL-divergence
    :return: (FloatTensor) - shape: (batch size,) - kl-divergence of KL(q \mid p).
    # init
    batch size = mu q.size(0)
    input size = np.prod(mu q.size()[1:])
    out = torch.ones(batch size,)
    out = out.type(torch.FloatTensor)
    for i in range(batch size):
        dq = torch.distributions.multivariate normal.MultivariateNormal(mu q[i],
torch.diag(logvar g[i].exp()))
        dp = torch.distributions.multivariate normal.MultivariateNormal(mu p[i],
torch.diag(logvar_p[i].exp()))
        kl div = 0
        for j in range(num_samples):
            sample = dq.sample()
            kl_div += dq.log_prob(sample) - dp.log prob(sample)
        out[i] = kl_div / float(num_samples)
    return out
```

Define data loaders

In [24]:

```
def get data loader(dataset location, batch size):
   URL = "http://www.cs.toronto.edu/~larocheh/public/datasets/binarized_mnist/"
   # start processing
   def lines to np array(lines):
        return np.array([[int(i) for i in line.split()] for line in lines])
   splitdata = []
    for splitname in ["train", "valid", "test"]:
        filename = "binarized mnist %s.amat" % splitname
        filepath = os.path.join(dataset_location, filename)
        utils.download url(URL + filename, dataset location)
        with open(filepath) as f:
            lines = f.readlines()
        x = lines to np array(lines).astype('float32')
        x = x.reshape(x.shape[0], 1, 28, 28)
        # pytorch data loader
        dataset = data utils.TensorDataset(torch.from numpv(x))
        dataset loader = data utils.DataLoader(x, batch size=batch size, shuffle
=splitname == "train")
        splitdata.append(dataset loader)
    return splitdata
```

In [25]:

```
train, valid, test = get_data_loader("binarized_mnist", 64)

Using downloaded and verified file: binarized_mnist/binarized_mnist_
train.amat
Using downloaded and verified file: binarized_mnist/binarized_mnist_
valid.amat
Using downloaded and verified file: binarized_mnist/binarized_mnist_
test.amat
```

Define VAE network architecture

```
class Encoder(nn.Module):
    def __init__(self, latent_size):
        super(Encoder, self). init ()
        self.mlp = nn.Sequential(
            nn.Linear(784, 300),
            nn.ELU(),
            nn.Linear(300, 300),
            nn.ELU(),
            nn.Linear(300, 2 * latent_size),
        )
    def forward(self, x):
        batch size = x.size(0)
        z mean, z logvar = self.mlp(x.view(batch size, 784)).chunk(2, dim=-1)
        return z mean, z logvar
class Decoder(nn.Module):
    def init (self, latent size):
        super(Decoder, self).__init__()
        self.mlp = nn.Sequential(
            nn.Linear(latent size, 300),
            nn.ELU(),
            nn.Linear(300, 300),
            nn.ELU(),
            nn.Linear(300, 784),
        )
    def forward(self, z):
        return self.mlp(z) - 5.
class VAE(nn.Module):
    def init (self, latent size):
        super(VAE, self). init ()
        self.encode = Encoder(latent size)
        self.decode = Decoder(latent size)
    def forward(self, x):
        z_mean, z_logvar = self.encode(x)
        z_sample = z_mean + torch.exp(z_logvar / 2.) * torch.randn_like(z_logvar
)
        x mean = self.decode(z sample)
        return z_mean, z_logvar, x_mean
    def loss(self, x, z_mean, z_logvar, x_mean):
        ZER0 = torch.zeros(z_mean.size())
        #kl = kl gaussian gaussian mc(z mean, z logvar, ZERO, zero, num samples=
1000).mean()
        kl = kl gaussian_gaussian_analytic(z_mean, z_logvar, ZERO, ZERO).mean()
        recon loss = -log likelihood bernoulli(
            torch.sigmoid(x mean.view(x.size(0), -1)),
            x.view(x.size(0), -1),
        ).mean()
        return recon_loss + kl
```

In [27]:

```
vae = VAE(100)
params = vae.parameters()
optimizer = Adam(params, lr=3e-4)
print(vae)
VAE(
  (encode): Encoder(
    (mlp): Sequential(
      (0): Linear(in_features=784, out_features=300, bias=True)
      (1): ELU(alpha=1.0)
      (2): Linear(in features=300, out features=300, bias=True)
      (3): ELU(alpha=1.0)
      (4): Linear(in features=300, out features=200, bias=True)
    )
  (decode): Decoder(
    (mlp): Sequential(
      (0): Linear(in features=100, out features=300, bias=True)
      (1): ELU(alpha=1.0)
      (2): Linear(in_features=300, out_features=300, bias=True)
      (3): ELU(alpha=1.0)
      (4): Linear(in_features=300, out_features=784, bias=True)
    )
  )
)
```

Train the model

In [28]:

```
for i in range(20):
    # train
    for x in train:
        optimizer.zero grad()
        z_{mean}, z_{logvar}, x_{mean} = vae(x)
        loss = vae.loss(x, z mean, z logvar, x mean)
        loss.backward()
        optimizer.step()
    # evaluate ELBO on the valid dataset
    with torch.no grad():
        total_loss = 0.
        total\_count = 0
        for x in valid:
            total loss += vae.loss(x, *vae(x)) * x.size(0)
            total count += x.size(0)
        print('-elbo: ', (total loss / total count).item())
-elbo:
        167.1219940185547
```

-elbo: 142.9504852294922 -elbo: 129.03280639648438 -elbo: 122.30564880371094 -elbo: 116.72787475585938 -elbo: 113.34302520751953 -elbo: 110.86123657226562 -elbo: 108.96890258789062 -elbo: 107.67586517333984 -elbo: 106.59246063232422 -elbo: 105.75444793701172 -elbo: 104.75553131103516 -elbo: 104.35298919677734 -elbo: 104.01703643798828 -elbo: 103.03252410888672 -elbo: 102.75845336914062 -elbo: 102.19798278808594 -elbo: 102.01860046386719 -elbo: 101.64897155761719 -elbo: 101.37193298339844

Save the model

```
torch.save(vae, 'model.pt')
```

/home/dhaivat1729/anaconda3/envs/detectron_fair/lib/python3.6/site-p ackages/torch/serialization.py:292: UserWarning: Couldn't retrieve s ource code for container of type VAE. It won't be checked for correc tness upon loading.

"type " + obj.__name__ + ". It won't be checked "
/home/dhaivat1729/anaconda3/envs/detectron_fair/lib/python3.6/site-p
ackages/torch/serialization.py:292: UserWarning: Couldn't retrieve s
ource code for container of type Encoder. It won't be checked for co
rrectness upon loading.

"type " + obj.__name__ + ". It won't be checked "
/home/dhaivat1729/anaconda3/envs/detectron_fair/lib/python3.6/site-p
ackages/torch/serialization.py:292: UserWarning: Couldn't retrieve s
ource code for container of type Sequential. It won't be checked for
correctness upon loading.

"type " + obj.__name__ + ". It won't be checked "
/home/dhaivat1729/anaconda3/envs/detectron_fair/lib/python3.6/site-p
ackages/torch/serialization.py:292: UserWarning: Couldn't retrieve s
ource code for container of type Linear. It won't be checked for cor
rectness upon loading.

"type " + obj.__name__ + ". It won't be checked "
/home/dhaivat1729/anaconda3/envs/detectron_fair/lib/python3.6/site-p
ackages/torch/serialization.py:292: UserWarning: Couldn't retrieve s
ource code for container of type ELU. It won't be checked for correc
tness upon loading.

"type " + obj.__name__ + ". It won't be checked "
/home/dhaivat1729/anaconda3/envs/detectron_fair/lib/python3.6/site-p
ackages/torch/serialization.py:292: UserWarning: Couldn't retrieve s
ource code for container of type Decoder. It won't be checked for co
rrectness upon loading.

"type " + obj.__name__ + ". It won't be checked "

Load the model

In [30]:

```
vae = torch.load('model.pt')
```

Evaluate the $\log p_{\theta}(x)$ of the model on test by using importance sampling

In [31]:

```
total loss = 0.
total\_count = 0
with torch.no grad():
    #x = next(iter(test))
    for x in test:
        # init
        K = 200
        M = x.size(0)
        # Sample from the posterior
        z mean, z logvar = vae.encode(x)
        eps = torch.randn(z mean.size(0), K, z mean.size(1))
        z_{samples} = z_{mean}[:, None, :] + torch_exp(z_logvar / 2.)[:, None, :] *
eps # Broadcast the noise over the mean and variance
        # Decode samples
        z samples flat = z samples.view(-1, z samples.size(-1)) # Flatten out th
e z samples
        x mean flat = vae.decode(z samples flat) # Push it through
        # Reshape images and posterior to evaluate probabilities
        x \text{ flat} = x[:, None].repeat(1, K, 1, 1, 1).reshape(M*K, -1)
        z mean flat = z mean[:, None, :].expand as(z samples).reshape(M*K, -1)
        z logvar flat = z logvar[:, None, :].expand as(z samples).reshape(M*K,
-1)
        ZEROS = torch.zeros(z mean flat.size())
        # Calculate all the probabilities!
        log p \times z = log likelihood bernoulli(torch.sigmoid(x mean flat), x flat)
.view(M, K)
        log_q_z_x = log_likelihood_normal(z_mean_flat, z_logvar_flat, z_samples_
flat).view(M, K)
        log p z = log likelihood normal(ZEROS, ZEROS, z samples flat).view(M, K)
        # Recombine them.
        W = log p x z + log p z - log q z x
        log p = log mean exp(w)
        # Accumulate
        total loss += log p.sum()
        total count += M
print('log p(x):', (total_loss / total_count).item())
```

log p(x): -95.58343505859375

In []: