## **AI701 HW2 Implementation Report**

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## **Code Implementation**

Before begin, we introduce our algorithm. For more details on the algorithm, please refer the homework submission.

```
Algorithm 1 P(\theta, Z|X) sampler using Gibbs + Metropolis-Hastings
Input: Dataset X = \{x_1, \dots, x_n\}
   Initialize \pi^{(1)}, \phi^{(1)} \sim p(\theta) and Z^{(1)} \sim P(Z|\pi^{(1)})
   for all t = 1, \cdots do
         Set Z^{(t+1)} = Z^{(t)}
         for all j=1,\cdots,n do
              \text{Sample } z_j^{(t+1)} \sim P\left(z_j | \left\{z_k^{(t+1)}\right\}_{k \neq j}\right) = P(z_j | Z^{(t+1)} \setminus \{z_j^{(t+1)}\}, X, \theta)
         for all j=1,\cdots,K do
              \alpha_i^{(t+1)} = \text{number of } i \text{ with } z_i^{(t+1)} = j
         Sample \pi^{(t+1)} \sim Dir(\mathbf{1} + \alpha^{(t+1)}) where \mathbf{1} = (1, 1, \dots, 1) \in \mathbb{Z}^K
         Propose \phi' \sim q(\phi'|\phi^{(t)})
         Compute A(\phi'|\phi) = min\left\{1, \frac{P(X|Z,\phi')P(\phi')q(\phi|\phi')}{P(X|Z,\phi)P(\phi)q(\phi'|\phi)}\right\}
         Sample u \sim \text{Unif } [0,1]
         if u \leq A(\phi'|\phi) then
               \phi^{(t+1)} = \phi'
               \phi^{(t+1)} = \phi^{(t)}
         end if
   end for
```

Now, we will briefly explain our code implementation with code blocks. Before we start, please note that this code implementation is same as the algorithm. (This code is just a direct implementation of the algorithm).

```
import numpy as np
from scipy.stats import multivariate_normal
import matplotlib
matplotlib.use('agg')
import matplotlib.pyplot as plt
# read data
x = []
with open('data.txt', 'r') as f:
    lines = f.readlines()
    for line in lines:
        x.append([float(val) for val in line.split()])
x = np.array(x)
alpha = [1 for _ in range(K)]
pi = np.random.dirichlet(alpha)
# sample z from categorical distribution conditioned on pi
z = np.random.choice([k+1 for k in range(K)], 1000, p = pi)
mu = np.random.multivariate_normal([0,0], [[5., 0.],[0., 5.]], size = K)
lmda = np.random.lognormal(.1, .1, size = K)
v = np.random.multivariate_normal([0,0], [[.25, 0.],[0., .25]], size = K)
```

This code block is for reading data and initializing the parameters. In other words, initialize  $\pi^{(1)}, \phi^{(1)} \sim p(\theta)$  and  $Z^{(1)} \sim P(Z|\pi^{(1)})$ 

```
# probability density of lognormal distribution
def lognormal_pdf(x, mu, sigma):
    return (1/(x * sigma * np.sqrt(2*np.pi))) * np.exp(-1 * ((np.log(x)-mu)**2)/(2*(sigma**2)))

# calculate p(z_i | {z_k}_(k != i))
def p_z(i, pi, x, mu, lmda, v):
    K = len(pi)
    p = []

for j in range(K):
    N = multivariate_normal(mean = mu[j], cov = lmda[j] * np.eye(2) + np.outer(v[j], v[j]))
    p.append(pi[j] * N.pdf(x[i]))

# normalize
sum = 0
for i in range(len(p)):
    sum += p[i]

return p/sum
```

This code block is for helper functions. The 'lognormal\_pdf' function calculates the probability density of the lognormal distribution. And the 'p\_z' function calculates

$$\begin{split} &P(z_i = j|Z \setminus \{z_i\}, X, \theta) \\ &= \frac{\pi_j \mathcal{N}(x_i|\mu_j, \lambda_j I_d + v_j v_j^T)}{\sum_{k=1}^K \pi_k \mathcal{N}(x_i|\mu_k, \lambda_k I_d + v_k v_k^T)} \\ &\quad \text{for all j = 1, ..., K} \end{split}$$

```
log_prob = 0.
    N_prime = multivariate_normal(mean = mu_prime[j], cov = lmda_prime[j] * np.eye(2) + np.outer(v_prime[j], v_prime[j]))
    log\_prob += np.log(N\_prime.pdf(x[i])) - np.log(N.pdf(x[i])) # log((X|Z,phi') / p(X|Z,phi))
N_v = multivariate_normal(mean = [0,0], cov = [[.25, 0.],[0., .25]])
p_phi, p_phi_prime = 0., 0.
    p_phi_prime += (np.log(N_mu.pdf(mu_prime[i])) + np.log(lognormal_pdf(lmda_prime[i], .1, .1))
                     + np.log(N_v.pdf(v_prime[i])))
log_prob += p_phi_prime - p_phi
q_phi, q_phi_prime = 0., 0.
    N_{mu} = multivariate_normal(mean = mu[i], cov = sigma_q2 * np.eye(2))
    N_v_{prime} = multivariate_{normal(mean} = v[i], cov = sigma_q2 * np.eye(2))
    N_mu = multivariate_normal(mean = mu_prime[i], cov = sigma_q2 * np.eye(2))
    N_v = multivariate_normal(mean = v_prime[i], cov = sigma_q2 * np.eye(2))
    q_phi += (np.log(N_mu.pdf(mu[i])) + np.log(lognormal_pdf(lmda[i], lmda_prime[i], sigma_q2)) + np.log(N_v.pdf(v[i])))
q_phi_prime += (np.log(N_mu_prime.pdf(mu_prime[i])) + np.log(lognormal_pdf(lmda_prime[i], lmda[i], sigma_q2))
                     + np.log(N_v_prime.pdf(v_prime[i])))
log_prob += q_phi - q_phi_prime
return min(np.exp(log_prob), 1.)
```

This code block is for calculating the acceptance probability

$$A(\phi'|\phi) = min\left\{1, \frac{P(X|Z,\phi')P(\phi')q(\phi|\phi')}{P(X|Z,\phi)P(\phi)q(\phi'|\phi)}\right\}$$

To prevent the likelihood overflow/underflow, use the log-exp trick. Calculate all the probabilities in log-scale, and exponent the log\_prob at the end.

```
def loglikelihood(x, z, pi, mu, lmda, v):
    log_prob = 0.

N_mu = multivariate_normal(mean = [0,0], cov = [[5., 0.],[0., 5.]])
N_v = multivariate_normal(mean = [0,0], cov = [[.25, 0.],[0., .25]])

for i in range(K):
    log_prob += np.log(N_mu.pdf(mu[i])) + np.log(N_v.pdf(v[i])) + lognormal_pdf(lmda[i], .1, .1)

for i in range(len(x)):
    j = z[i] - 1
    N = multivariate_normal(mean = mu[j], cov = lmda[j] * np.eye(2) + np.outer(v[j], v[j]))
    log_prob += np.log(pi[j] * N.pdf(x[i]))

return log_prob
```

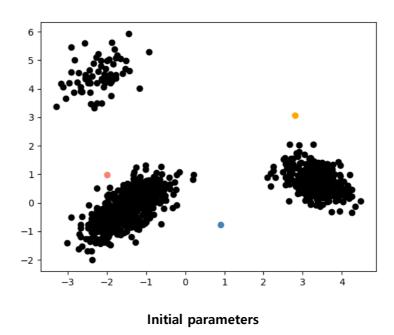
This code block is for calculating the marginal log-likelihood  $\log p(\overline{X,Z,\theta})$  to trace the convergence.

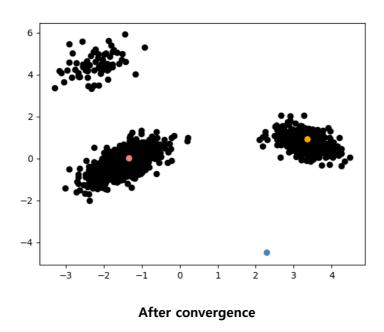
```
plt.scatter(x[:,0], x[:,1])
plt.savefig('data.png')
# main procedure
loglikelihoods = []
max_iter = 1000
for iter in range(max_iter):
    for j in range(len(x)):
        z[j] = np.random.choice([k+1 for k in range(K)], p = p_z(j, pi, x, mu, lmda, v))
    new_alpha = []
    for j in range(1, K+1):
       new_alpha.append(list(z).count(j))
    alpha = [x+1 for x in new_alpha]
    pi = np.random.dirichlet(alpha)
    mu_prime, lmda_prime, v_prime = [], [], []
    sigma_q2 = 1.
       mu_prime.append(np.random.multivariate_normal(mu[i], sigma_q2 * np.eye(2)))
        lmda_prime.append(np.random.lognormal(np.log(lmda[i]), sigma_q2))
       v_prime.append(np.random.multivariate_normal(v[i], sigma_q2 * np.eye(2)))
    mu_prime = np.vstack(mu_prime)
    lmda_prime = np.hstack(lmda_prime)
    v_prime = np.vstack(v_prime)
   u = np.random.random()
    a = accept_prob(x, z, mu, lmda, v, mu_prime, lmda_prime, v_prime)
    print('Acceptance probability : %.4f'%(a))
    if u < a:
       print('Accepted')
        mu = mu_prime
       lmda = lmda_prime
       v = v_prime
       plt.scatter(mu[0][0], mu[0][1], color='salmon')
        \verb"plt.scatter(mu[1][0], mu[1][1], color='orange')
        plt.savefig('results/estimation_iter%d.png'%(iter+1))
    ll = loglikelihood(x, z, pi, mu, lmda, v)
    loglikelihoods.append(ll)
    plt.plot(range(1, iter+2), loglikelihoods)
    plt.xlabel('iteration')
    plt.ylabel('log-likelihood')
    plt.title('Trace of log-likelihood')
    plt.savefig('trace.png')
    plt.close()
    print('[%d/%d] Marginal LogLikelihood : %.4f'%(iter+1, max_iter, ll))
```

This code block is the implementation of our main algorithm. First, sample Z from the categorical distribution conditioned on pi using the Gibbs sampling. The, calculate the new alpha (will be the parameter of the Dirichlet distribution). And sample the pi from the resulting Dirichlet distribution. And then, sample mu, lambda, v using the Metropolis-Hastings algorithm. First propose mu', lambda', v' from the proposal distribution. Second, using 'accept\_prob' function, calculate the acceptance probability. Next sample u from Uniform[0,1] and if u < acceptance probability, then accept the proposed parameters. Otherwise, do not update the parameters mu, lambda and v. When the new parameters are accepted, draw a picture to see the new parameter. At the end of each loop, calculate the marginal log-likelihood and print/plot it to trace the convergence.

## **Experimental Results**

In the following figures, the black points are the data points and red, orange, blue points are the mean of each Gaussian component.

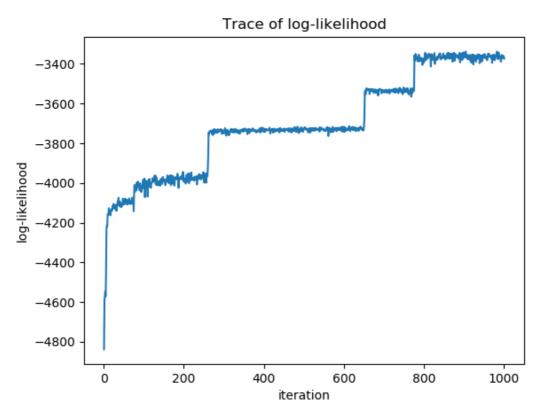




As we can see in the figure, means of two Gaussian components (red, orange points) are

seems quite reasonable. However, the other mean (blue point) seems couldn't find the right position.

The following figure is the trace of the marginal log-likelihood.



Trace of the marginal log-likelihood

This figure shows that even if our sampler is a little slow, it's going in the right direction.