# **Introduction to Probabilistic Graphical Models**

## **Practical Session 2**

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```
In [160]: %matplotlib inline
    import numpy as np
    from scipy.linalg import norm
    import matplotlib.pyplot as plt
    import networkx as nx
    import pyparsing
    from IPython.display import Math
    import copy
    import math
    from matplotlib.patches import Ellipse
    ## we first run the code with the warning enabled, if no error of relevant warning is to be found,
    ## we disable the warning then
    # import warnings
    # warnings.filterwarnings("ignore")
math_pi = math.pi
```

```
In [161]: # choose a large font size by default and use tex for math
    fontsize = 18
    params = {'axes.labelsize': fontsize + 2,
        'font.size': fontsize + 2,
        'legend.fontsize': fontsize + 2,
        'xtick.labelsize': fontsize,
        'ytick.labelsize': fontsize,
        'text.usetex': True}
    plt.rcParams.update(params)
```

#### **Question 1**

We need to compute

$$\gamma_i(x) = rac{\pi_i \mathcal{N}(x; \mu_i, \Sigma_i)}{\sum_{j=1}^K \pi_j \mathcal{N}(x; \mu_j, \Sigma_j)}$$

We define  $g_i(x)=\pi_i\mathcal{N}(x;\mu_i,\Sigma_i)$  and  $lg_i=\log g_i(x)$ , then the log of the nominator is equal to:

$$egin{aligned} \log g_i(x) &= \log(\pi_i \mathcal{N}(x; \mu_i, \Sigma_i)) \ &= \log \left( \pi_i rac{1}{(2\pi)^{K/2} |\Sigma_i|^{1/2}} \mathrm{exp} igg( -rac{1}{2} (x - \mu_i)^T \Sigma_i^{-1} (x - \mu_i) igg) 
ight) \ &= \log \pi_i - rac{1}{2} (K \log(2\pi) + \log |\Sigma_i|) - rac{1}{2} (x - \mu_i)^T \Sigma_i^{-1} (x - \mu_i) \end{aligned}$$

Then we could compute  $gamma_i(x)$  as follows:

$$egin{aligned} \gamma_i(x) &= rac{\pi_i \mathcal{N}(x; \mu_i, \Sigma_i)}{\sum_{j=1}^K \pi_j \mathcal{N}(x; \mu_j, \Sigma_j)} \ &= rac{g_i(x)}{\sum_{j=1}^K g_j(x)} = rac{\exp(\log(g_i(x)))}{\sum_{j=1}^K \exp(\log(g_j(x)))} \ &= rac{\exp(lg_i - maxlg) \exp(maxlg)}{\sum_{j=1}^K \exp(lg_j - maxlg) \exp(maxlg)} \ &= rac{\exp(lg_i - maxlg)}{\sum_{j=1}^K \exp(lg_j - maxlg)} \end{aligned}$$

where  $maxlg = \max_{j} lg_j = \max_{j} \log g_j(x), j = 1, \ldots, K$ 

## **Question 2**

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From lecture notes, we have:

$$\mathcal{L}_t( heta) = \sum_{n=1}^N \sum_{k=1}^K \gamma_k^{(t)}(x_n) \log \mathcal{N}(x_n; \mu_k, \Sigma_k) + \sum_{n=1}^N \sum_{k=1}^K \gamma_k^{(t)}(x_n) \log \pi_k$$

 ${\mathcal M}$ -step

$$heta^{(t+1)} = rg \max_{ heta} \mathcal{L}_t( heta)$$

where  $heta^{(t)} = (\mu_k^{(t)}, \Sigma_k^{(t)}, \pi_k^{(t)})$ 

For  $\mu_k^{(t+1)}$  :

$$egin{aligned} rac{\partial \mathcal{L}_t( heta)}{\partial \mu_k^*} & \propto \sum_{n=1}^N \gamma_k^{(t)}(\Sigma_k)^{-1} (x_n - \mu_k^*) = 0 \ & \sum_{n=1}^N \gamma_k^{(t)}(x_n) (\Sigma_k)^{-1} (x_n - \mu_k^*) = 0 \ & \sum_{n=1}^N \gamma_k^{(t)}(x_n) (\Sigma_k)^{-1} x_n = \sum_{n=1}^N \gamma_k^{(t)}(\Sigma_k)^{-1} \mu_k^* \ & \mu^* (:= \mu_k^{(t+1)}) = rac{\sum_{n=1}^N \gamma_k^{(t)}(x_n) . \, x_n}{\sum_{n=1}^N \gamma_k^{(t)}(x_n)} \end{aligned}$$

For 
$$\pi_k^{(t+1)}$$

We rewirte the formula for

$$egin{aligned} \mathcal{L}_t( heta) &= \sum_{n=1}^N \sum_{k=1}^K \gamma_k^{(t)}(x_n) \log(\pi_k \mathcal{N}(x_n; \mu_k, \Sigma_k)) \ &= \sum_{n=1}^N \sum_{k=1}^K \gamma_k^{(t)}(x_n) l_k(x_n) \ &= \sum_{n=1}^N \sum_{k=1}^K \gamma_k^{(t)}(x_n) \left( \log \pi_k - rac{1}{2} (K \log(2\pi) + \log |\Sigma_k|) - rac{1}{2} (x - \mu_k)^T \Sigma_k^{-1} (x - \mu_i)) 
ight) \ &= rac{\partial \mathcal{L}_t( heta)}{\partial \pi_k^*} = 0 \ : \ \sum_{n=1}^N \sum_{k=1}^K \gamma_k^{(t)}(x_n) rac{1}{\pi_k^*} = 0 \end{aligned}$$

Together With  $\sum_{k=1}^K \pi_k^* = 1$ , we get:

$$\pi_k^*(:\pi_k^{(t+1)}) = rac{1}{N} \sum_{n=1}^N \gamma_k^{(t)}(x_n)$$

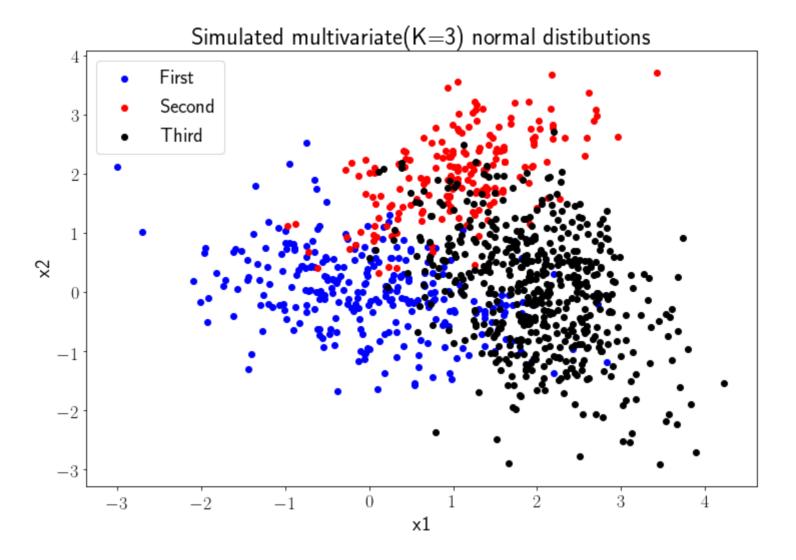
For 
$$\Sigma_k^{(t+1)}$$
 :

$$egin{aligned} rac{\partial \mathcal{L}_t( heta)}{\partial \Sigma_k^*} &= 0 \; : \; \propto \; \sum_{n=1}^N \gamma_k^{(t)}(x_n) \left( -(\Sigma_k^*) + (x-\mu_k)(x-\mu_k)^T 
ight) = 0 \ & \sum_{n=1}^N \gamma_k^{(t)}(x_n) \Sigma_k^* = \sum_{n=1}^N \gamma_k^{(t)}(x_n) \left( (x-\mu_k)(x-\mu_k)^T 
ight) \ & \Sigma_k^* (:= \Sigma_k^{t+1}) = rac{\sum_{n=1}^N \gamma_k^{(t)}(x_n)(x-\mu_k)(x-\mu_k)^T}{\sum_{n=1}^N \gamma_k^{(t)}(x_n)} \end{aligned}$$

### **Question 3**

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```
In [162]: # hyper-parameters
          K = 3
          colors = ['b', 'r', 'k']
          pi = np.array([0.3, 0.2, 0.5])
          us = np.array([[0, 0], [1, 2], [2,0]])
          \# sq1 = np.array([[1, -0.25], [-0.25, 0.5]])
          \# sg2 = np.array([[0.5, 0.25], [0.25, 0.5]])
          \# sg3 = np.array([[0.5, -0.25], [-0.25, 1]])
          \# sigmas = [sq1, sq2, sq3]
          sgs = np.array([[[1, -0.25], [-0.25, 0.5]], [[0.5, 0.25], [0.25, 0.5]], [[0.5, -0.25], [-0.25, 1]]])
          def sample loc (prob, pi distrbution):
              # return the idx loction of prob in the prob distrbution
              nr loc = len(pi distrbution)
              cum sum = np.cumsum(pi distrbution)
              idx = np.where(prob < cum sum)[0]
              return idx[0]
          N = 1000 # number of samples
          xs = np.zeros((N, 3))
          # simulator the trajectory
          for i in np.arange(0,N):
              si pi = np.random.uniform(0,1)
              gs_id = sample_loc(si_pi, pi)
              ui = us[gs id]
              sqi = sqs[qs id]
              xs[i,[0,1]] = np.random.multivariate normal(ui, sgi)
              xs[i,2] = gs id
          x a1 = xs[xs[:.2] == 0][:.[0.1]]
```



```
In [163]: def cal loggi(pi, ui, sigi, x, nr x):
              sigi inv = np.linalq.inv(sigi)
              log2pi = np.log(2* math pi)
              log sigi det = np.log(np.linalg.det(sigi))
              diff = x - ui
              exp part = [- diff[j,:].dot(sigi inv).dot(diff[j,:]) / 2 for j in np.arange(nr x)]
              logqi = np.log(pi) - (K * log2pi + log siqi det) / 2 + exp part
              return loggi
          def cal gammas(K, ps, us, sigs, xs):
              gamma KNs = np.zeros((K,N))
              nr x = N
              logg KN = np.zeros((K, N))
              for j in np.arange(K):
                  logg KN[j,:] = cal loggi(ps[j], us[j], sigs[j], xs, nr x)
              max lgs = np.max(logg KN, axis=0)
              lg diff = logg KN - max lgs
              sum exp = np.sum(np.exp(lg diff), axis=0)
              for j in np.arange(K):
                  gamma KNs[j,:] = np.divide(np.exp(logg KN[j,:] - max lgs), sum exp)
              return gamma KNs
```

```
In [164]: def update_para(gammas, K, ps, us, sigs, xs):
    ps_new = ps.copy()
    us_new = us.copy()
    sigs_new = sigs.copy()

    for j in np.arange(K):

        sum_gamma = np.sum(gammas[j,:])
        us_new[j,:] = np.sum(xs.T * gammas[j,:], axis=1) / sum_gamma
        ps_new[j] = sum_gamma / N
        diff = xs - us_new[j,:]
        sig_nom = [gammas[j,i] * np.outer(diff[i,:], diff[i,:]) for i in np.arange(N)]
        sigs_new[j,:] = np.sum(sig_nom, axis=0) / sum_gamma

    return us_new, ps_new, sigs_new
```

```
In [165]: def cal_loglik(K, ps, us, sigs, xs, gammas):
    loglik = 0
    for k in range(K):
        sigi = sigs[k]
        sigi_inv = np.linalg.inv(sigi)
        log2pi = np.log(2* math_pi)
        log_sigi_det = np.log(np.linalg.det(sigi))
        ui = us[k]
        pi = ps[k]
        diff = xs - ui

    exp_part = [- diff[j,:].dot(sigi_inv).dot(diff[j,:]) / 2 for j in np.arange(N)]
    log_part = - (K * log2pi + log_sigi_det) / 2 + exp_part + np.log(pi)
    loglik += np.sum(gammas[k,:] * (log_part + exp_part))
    return loglik
```

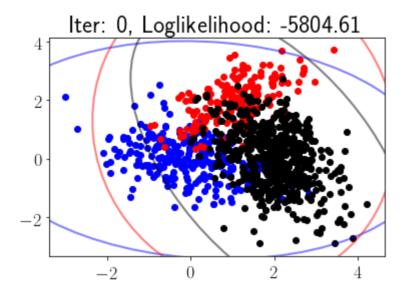
```
In [167]: xs_nl = xs[:,:2] # not labeled data
xs_mean = np.mean(xs_nl, axis=0)
xs_cov = np.cov(xs_nl.T)

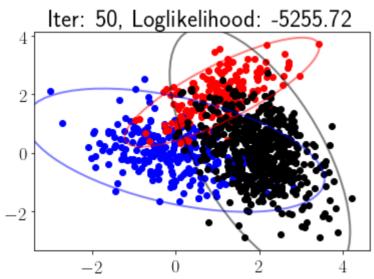
# initilize the parameters: set them to be equal
xs = xs_nl.copy()
ps = np.ones((3,1)) / K

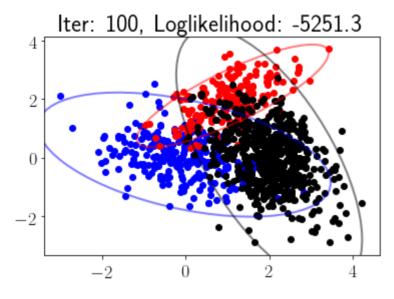
us = np.array([[0, 0.5], [1, 1.2], [2,0.1]])
#us = np.array([xs_mean, xs_mean, xs_mean])
sigs = np.array([xs_cov, xs_cov, xs_cov])
gammas = cal_gammas(K, ps, us, sigs, xs)
log_lik = cal_loglik(K, ps, us, sigs, xs, gammas)
print (log_lik)
```

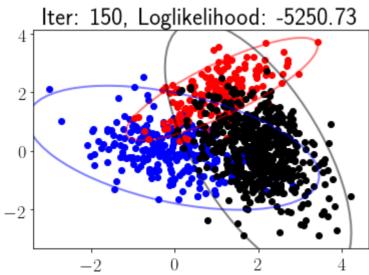
-5804.6052496393695

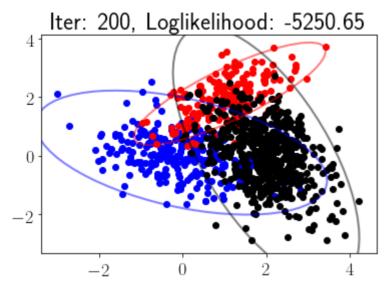
```
In [168]: #### 2. Implement the EM algorithm for GMMs
          xs nl = xs[:,:2] # not labeled data
          xs mean = np.mean(xs nl, axis=0)
          xs cov = np.cov(xs nl.T)
          # initilize the parameters: set them to be equal
          xs = xs \ nl.copy()
          ps = np.ones((3,1)) / K
          us = np.array([[0, 0.5], [1, 1.2], [2,0.1]]) # random choose some centers
          sigs = np.array([xs cov, xs cov, xs cov])
          Nr iter = 500
          log liks = np.zeros((Nr iter, 1))
          for it in np.arange(Nr iter):
              gammas = cal gammas(K, ps, us, sigs, xs)
              log lik = cal loglik(K, ps, us, sigs, xs, gammas)
              us new, ps new, sigs new = update para(gammas, K, ps, us, sigs, xs)
              log liks[it] = log lik
              title str = "Nr Iter: " + str(it) + ", Loglikelihood: " + str(round(log lik, 2))
              if it % 50 == 0:
                  #fig, ax = plt.subplots(figsize=(8,8))
                  fig, ax = plt.subplots()
                  plt.scatter(x g1[:,0], x g1[:,1], c='b', label='First')
                  plt.scatter(x g2[:,0], x g2[:,1], c='r', label='Second')
                  plt.scatter(x g3[:,0], x g3[:,1], c='k', label='Third')
```

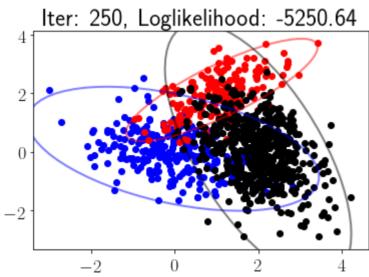


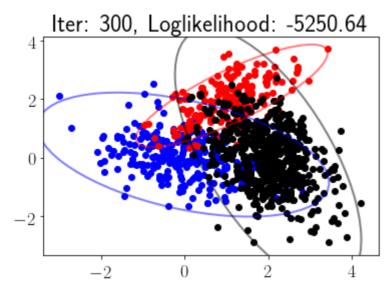


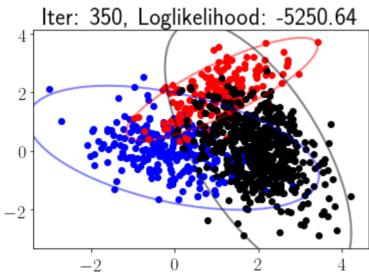


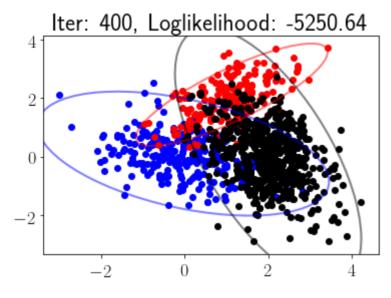


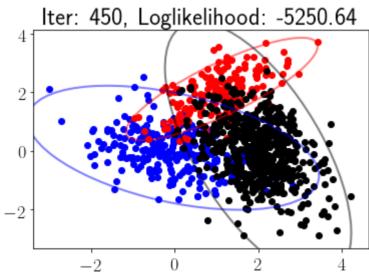






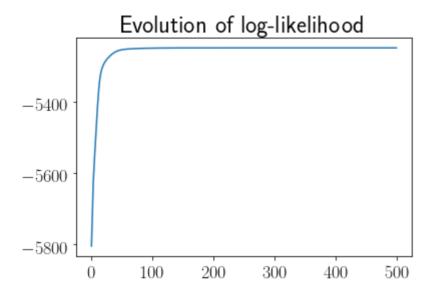




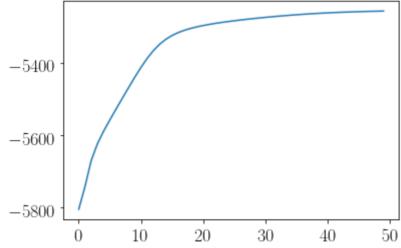


```
In [169]: # Evolution of log-likelihood
    plt.plot(np.arange(Nr_iter), log_liks)
    plt.title("Evolution of log-likelihood")
    plt.show()

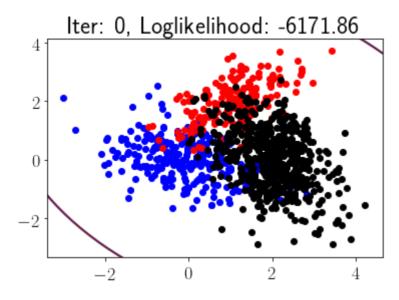
plt.plot(np.arange(50), log_liks[:50])
    plt.title("Zoom In: show the first 50 iterations: Evolution of log-likelihood")
    plt.show()
```

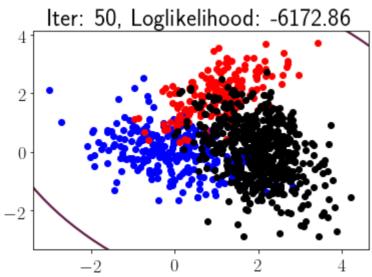


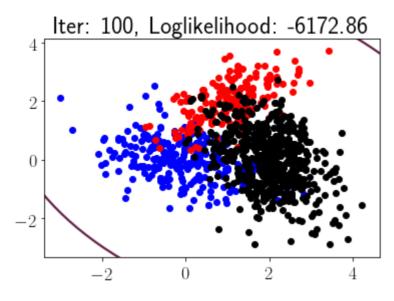
Zoom In: show the first 50 iterations: Evolution of log-likelihood

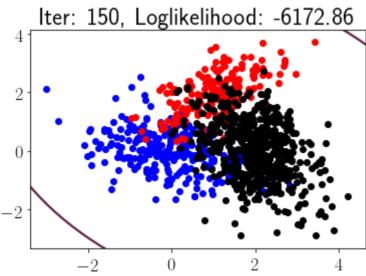


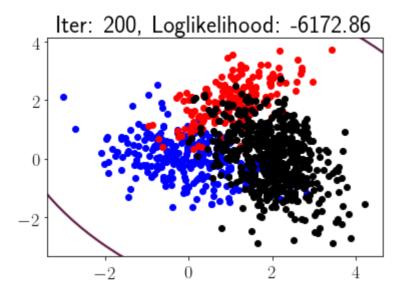
```
In [170]: ### Test with different initial values:
          xs nl = xs[:,:2] # not labeled data
          xs mean = np.mean(xs nl, axis=0)
          xs cov = np.cov(xs nl.T)
          # initilize the parameters with the same values
          ps = np.ones((3,1)) / K
          us = np.array([xs mean, xs mean, xs mean])
          sigs = np.array([xs_cov, xs cov, xs cov])
          Nr iter = 500
          log liks = np.zeros((Nr iter, 1))
          for it in np.arange(Nr iter):
              gammas = cal gammas(K, ps, us, sigs, xs)
              log lik = cal loglik(K, ps, us, sigs, xs, gammas)
              us new, ps new, sigs new = update para(gammas, K, ps, us, sigs, xs)
              log liks[it] = log lik
              title str = "Nr Iter: " + str(it) + ", Loglikelihood: " + str(round(log lik, 2))
              if it % 50 == 0:
                  #fig, ax = plt.subplots(figsize=(8,8))
                  fig, ax = plt.subplots()
                  plt.scatter(x g1[:,0], x g1[:,1], c='b', label='First')
                  plt.scatter(x g2[:,0], x g2[:,1], c='r', label='Second')
                  plt.scatter(x g3[:,0], x g3[:,1], c='k', label='Third')
                  plot contour(ax, us new, sigs new, colors)
                  ax.set title("Iter: " + str(it) + ", Loglikelihood: " + str(np.round(log lik, 2)))
                  nlt.show()
```

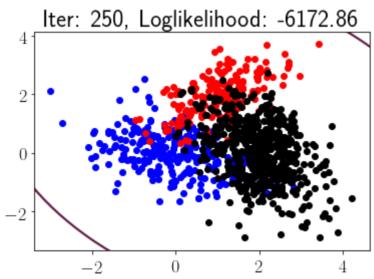


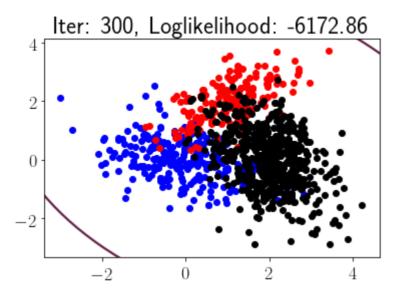


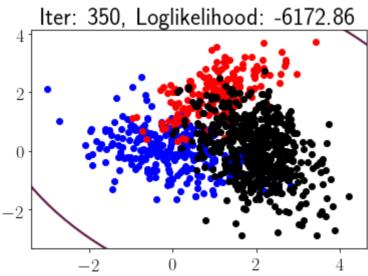


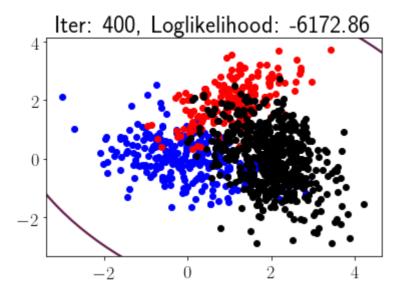


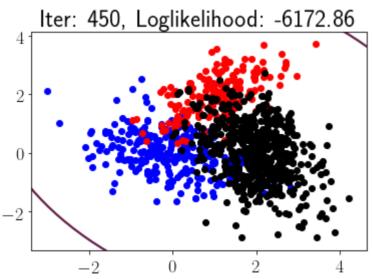






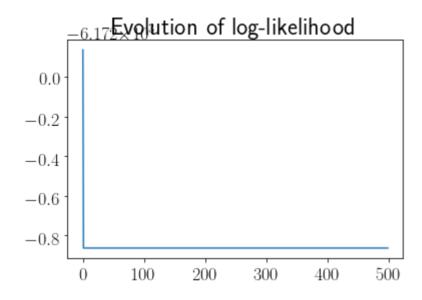




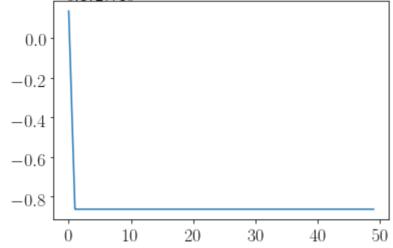


```
In [171]: # Evolution of log-likelihood
    plt.plot(np.arange(Nr_iter), log_liks)
    plt.title("Evolution of log-likelihood")
    plt.show()

plt.plot(np.arange(50), log_liks[:50])
    plt.title("Zoom In: show the first 50 iterations: Evolution of log-likelihood")
    plt.show()
```



Zoom In: show\_the2first 50 iterations: Evolution of log-likelihood



Conclunsion: the model is very sensitive to the inital data