CS5340 - Lab3 Report

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1. E-step

Emission probability we first compute emission probability with scipy.stats.norm.pdf function.

$$emission[i, n, k] = p(x_n^{(i)}|z_{nk} = 1, \mu_k, \sigma_k)$$

Here $i=1,2,\ldots,T$ is the index of observed sequence, $n=1,2,\ldots,N$ is the index of node inside each sequence, and $k=1,2,\ldots,K$ refers to the hidden state.

 $\hat{\alpha}^{(i)}(z_1)$ and $\hat{\beta}^{(i)}(z_N)$ initialize $\hat{\alpha}^{(i)}(z_1) = \frac{\alpha^{(i)}(z_1)}{c_1}$, with $\alpha^{(i)}(z_1) = p(x_1^{(i)}, z_1) = p(x_1^{(i)}|z_1)p(z_1)$ and normalizer $c_1 = \sum_{z_1} \alpha^{(i)}(z_1)$, and initialize $\hat{\beta}^{(i)}(z_N) = \mathbf{1}$

forward process for n = 2 to N, compute $\hat{\alpha}^{(i)}(z_n), c_n$ with

$$c_n \hat{\alpha}^{(i)}(z_n) = p(x_n^{(i)}|z_n) \sum_{z_{n-1}} \hat{\alpha}^{(i)}(z_{n-1}) p(z_n|z_{n-1}) := \tilde{\alpha}^{(i)}(z_n)$$
$$c_n = \sum_{z} \tilde{\alpha}^{(i)}(z_n)$$

backward process for n = N-1 to 1, compute $\hat{\beta}^{(i)}(z_n)$ with

$$c_{n+1}\hat{\beta}^{(i)}(z_n) = \sum_{z_{n+1}} p(x_{n+1}^{(i)}|z_{n+1}) p(z_{n+1}|z_n) \hat{\beta}^{(i)}(z_{n+1}) := \tilde{\beta}^{(i)}(z_n)$$

compute γ and ξ with

$$\gamma^{(i)}(z_n) = \hat{\alpha}^{(i)}(z_n)\hat{\beta}^{(i)}(z_n)$$

$$\xi^{(i)}(z_{n-1}, z_n) = \frac{\hat{\alpha}^{(i)}(z_{n-1})p(x_n^{(i)}|z_n)p(z_n|z_{n-1})\hat{\beta}^{(i)}(z_n)}{c_n}$$

2. M-step

update π with

$$\pi_k = \frac{\sum_{i=1}^T \gamma^{(i)}(z_{1k})}{T}$$

update A with

$$A_{jk} = \frac{\sum_{i=1}^{T} \sum_{n=2}^{N} \xi^{(i)}(z_{n-1,j}, z_{nk})}{\sum_{i=1}^{T} \sum_{n=2}^{N} \sum_{l=1}^{K} \xi^{(i)}(z_{n-1,j}, z_{nl})}$$

update μ_k and σ_k here $x_n^{(i)} \in \mathbb{R}$ is sampled from Gaussian distribution $\mathcal{N}(\mu_k, \sigma_k)$

$$\mu_k = \frac{\sum_{i=1}^{T} \sum_{n=1}^{N} \gamma^{(i)}(z_{nk}) x_n^{(i)}}{\sum_{i=1}^{T} \sum_{n=1}^{N} \gamma^{(i)}(z_{nk})}$$

$$\sigma_k = \frac{\sum_{i=1}^{T} \sum_{n=1}^{N} \gamma^{(i)}(z_{nk}) (x_n^{(i)} - \mu_k)^2}{\sum_{i=1}^{T} \sum_{n=1}^{N} \gamma^{(i)}(z_{nk})}$$

3. EM algorithm

initialize π, A, μ, σ with K-means clustering update params run E-step and M-step iteratively until convergence.

4. Results

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yichao@macbook:~/Documents/CS5340/lab3$ python test_lab3.py
e step: PASSED
m_step: PASSED
Running on seq_short
Loaded 200 sequences, with average length = 8.0
Groundtruth pi:
    [0.9 0.1 0.]
Groundtruth A:
   [[0.3 0.7 0.]
    [0. 0.6 0.4]
    [0. 0. 1.]]
Groundtruth phi:
   {'mu': array([-1.5, 0.5, -0.2]), 'sigma': array([0.5, 0.2, 0.3])}
Your pi:
    [0. 0.91 0.09]
Your A:
   [[1. 0. 0.]
    [0. 0.28 0.72]
   [0.4 0. 0.6]]
Your phi:
   {'mu': array([-0.21, -1.5, 0.52]), 'sigma': array([0.29, 0.51, 0.21])}
Time running fit_hmm() on dataset seq_short : 1.21s
Running on seq_long
Loaded 5 sequences, with average length = 1000.0
Groundtruth pi:
    [0.6 0. 0. 0.4]
Groundtruth A:
    [[0.3 0. 0. 0.7]
    [0.7 0.3 0. 0.]
    [0. 0.6 0.4 0.]
    [0. 0. 0.4 0.6]]
Groundtruth phi:
   {'mu': array([0., 1.1, 2., 1.]), 'sigma': array([0.5, 0.3, 0.4, 0.2])}
Your pi:
    [0.64 0. 0.36 0. ]
Your A:
   [[0.3 0. 0.7 0. ]
    [0.7 0.3 0. 0. ]
    [0. 0. 0.57 0.43]
   [0.
         0.58 0. 0.42]]
Your phi:
   {'mu': array([0. , 1.07, 1. , 2. ]), 'sigma': array([0.52, 0.31, 0.19, 0.41])}
Time running fit_hmm() on dataset seq_long : 3.56s
Running on seq_short2
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```
Loaded 100 sequences, with average length = 9.0
Groundtruth pi:
[0.4 0.3 0. 0.3]
Groundtruth A:
[[0.4 0.6 0. 0.]
[0.2 0.2 0.6 0. ]
[0. 0. 0.1 0.9]
[0. 0. 0. 1.]]
Groundtruth phi:
{\text{'mu': array([ 0. , -1. , 0.5, -0.5]), 'sigma': array([0.2, 0.2, 0.1, 0.1])}}
Your pi:
[0.33 0.25 0.42 0. ]
Your A:
[[0.18 0. 0.2 0.62]
[0. 1. 0. 0. ]
[0.58 0. 0.42 0. ]
[0. 0.96 0. 0.04]]
Your phi:
{'mu': array([-0.99, -0.5, 0.01, 0.5]), 'sigma': array([0.22, 0.1, 0.17, 0.09])}
Time running fit_hmm() on dataset seq_short2 : 0.45s
Running on seq_long2
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Loaded 5 sequences, with average length = 800.0
Groundtruth pi:
    [1. 0. 0. 0. 0. 0.]
Groundtruth A:
    [[0.3 0.7 0. 0. 0. 0.]
    [0. 0.6 0.4 0. 0. 0.]
    [0.2 0. 0.3 0.5 0. 0.]
    [0. 0. 0. 0.4 0.5 0.1]
    [0. 0. 0. 0. 0.8 0.2]
    [0.5 0. 0. 0.1 0. 0.4]]
Groundtruth phi:
   {'mu': array([ 0.5,  0. ,  2. , -1. , -0.2,  1.1]), 'sigma': array([0.5 , 0.25,
   0.2 , 0.2 , 0.4 , 0.1 ])}
Your pi:
   [0. 0. 0. 0. 1. 0.]
Your A:
    [[0.6 0.4 0. 0. 0. 0.]
    [0. 0.26 0.53 0. 0.2 0. ]
    [0. 0. 0.43 0.09 0. 0.47]
    [0. 0. 0.08 0.42 0.49 0. ]
    [0.69 0. 0. 0. 0.31 0. ]
    [0. 0. 0. 0.21 0. 0.79]]
Your phi:
   \{\text{'mu': array}([-0.01, 1.99, -0.98, 1.1, 0.49, -0.21]), 'sigma': array([0.25, 0.19,
   0.2 , 0.1 , 0.49, 0.39])}
Time running fit_hmm() on dataset seq_long2 : 3.33s
```