# Variational Inference for Bayesian Mixture of Factor Analysers

### Angermueller Christof

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### 1 Notations

| Symbol         | Description                  |
|----------------|------------------------------|
| $\overline{P}$ | Dimensionality data          |
| Q              | Number of factors/components |
| S              | Number of factor analysers   |
| N              | Number of samples            |
| G              | Number of genes              |

## 2 Factor analysis

| Symbol           | Dimension    | Description                               |
|------------------|--------------|---|
| $\overline{y}$   | P            | Observed data vector                      |
| $oldsymbol{x}$   | Q            | Latent factors                            |
| $\eta$           | P            | Random Gaussian noise                     |
| $\Lambda$        | $P \times Q$ | Factor loading matrix                     |
| $\Lambda_{:q}$   | P            | Column loading matrix/principal component |
| $\boldsymbol{z}$ | Q            | Indicators factor analysers               |
| $\pi$            | S            | Mixing proportions                        |

$$P(\boldsymbol{x}) = \mathcal{N}(\boldsymbol{x}|0, I_Q) \tag{1}$$

$$P(\eta) = \mathcal{N}(\eta|0, \Psi) \tag{2}$$

### 2.1 Single factor analyser

$$y = \Lambda x + \eta \tag{3}$$

$$P(\boldsymbol{y}|\boldsymbol{x}, \Lambda, \Psi) = \mathcal{N}(\boldsymbol{y}|\Lambda \boldsymbol{x}, \Lambda \Lambda^T + \Psi)$$
(4)

## 2.2 Mixture factor analysers

$$\boldsymbol{y} = \boldsymbol{\eta} + \sum_{s=1}^{S} \Lambda^{s} \boldsymbol{x} P(z_{s} | \boldsymbol{\pi})$$
 (5)

$$P(\boldsymbol{y}|\boldsymbol{x}, \Lambda, \Psi) = \sum_{s=1}^{S} \mathcal{N}(\boldsymbol{y}|\Lambda \boldsymbol{x}, \Lambda^{s} \Lambda^{sT} + \Psi) P(z_{s}|\boldsymbol{\pi})$$
(6)

### 3 Gene expression model

| Symbol               | Dimension    | Description                                     |
|----------------------|--------------|---|
| $oldsymbol{y}^{n,g}$ | P            | Counts sample $n$ gene $g$                      |
| $oldsymbol{x}^{n,g}$ | Q            | Factors sample $n$ gene $g$                     |
| $oldsymbol{z}^{n,g}$ | Q            | Indicators factor analysers sample $n$ gene $g$ |
| $\pi$                | S            | Mixing proportions                              |
| $\Lambda^s$          | $P \times Q$ | Loading matrix factor analyser $s$              |
| $oldsymbol{ u}^s$    | Q            | Precisions components factor analyser $s$       |
| $\overline{a}$       | 1            | Shape gamma distribution                        |
| b                    | 1            | Rate gamma distribution                         |
| $\alpha$             | 1            | Dirichlet parameter                             |
| $\Psi$               | $Q \times Q$ | Noise matrix                                    |

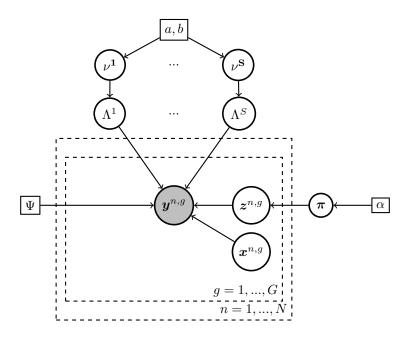


Figure 1: Graphical model

### 3.1 Conditional probability distributions

$$P(\boldsymbol{y}^{n,g}|\boldsymbol{x}^{n,g},\boldsymbol{z}^{n,g},\boldsymbol{\pi},\Lambda) = \sum_{s=1}^{S} \mathcal{N}(\boldsymbol{y}^{n,g}|\Lambda^{s}\boldsymbol{x}^{n,g},\Lambda^{s}\Lambda^{sT} + \Psi)P(z_{s}^{n,g}|\boldsymbol{\pi})$$
(7)

$$P(\boldsymbol{x}^{n,g}) = N(\boldsymbol{x}^{n,g}|0, I_Q)$$
(8)

$$P(\boldsymbol{z}^{n,g}|\boldsymbol{\pi}) = Mult(\boldsymbol{z}^{n,g}|\boldsymbol{\pi}) \tag{9}$$

$$P(\boldsymbol{\pi}|\alpha) = Dir(\boldsymbol{\pi}|\alpha) \tag{10}$$

$$P(\Lambda^{s}) = \prod_{q=1}^{Q} N(\Lambda^{s}_{:q}|0, I_{P} \frac{1}{\nu_{q}^{s}})$$
(11)

$$P(\boldsymbol{\nu}^s) = \prod_{q=1}^{Q} \Gamma(\nu_q^s | a, b)$$
 (12)