Deep Poisson Factor Modeling

Ricardo Henao, Zhe Gan, James Lu and Lawrence Carin

Department of Electrical and Computer Engineering Duke University, Durham, NC 27708 {r.henao,zhe.gan,james.lu,lcarin}@duke.edu

1 Graphical Model

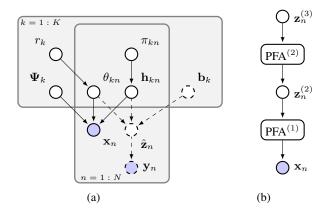


Figure 1: Graphical models. (a) Poisson Factor Analysis (PFA) module. Nodes $(\mathbf{b}_k, \hat{\mathbf{z}}_n \text{ and } \mathbf{y}_n)$ and edges drawn with dashed lines correspond to the discriminative PFA. (b) Deep Poisson factor model. Filled and empty nodes represent observed and latent variables, respectively.

2 Inference Details

2.1 MCMC

Conditional posteriors (layer index omitted for clarity):

$$\psi_{k} \sim \text{Dirichlet}(\eta + x_{1k}, \dots, \eta + x_{Mk}),$$

$$\theta_{kn} \sim \text{Gamma}(r_{k}h_{kn} + x_{kn}, b^{-1}),$$

$$h_{kn} \sim \delta(x_{kn} = 0) \text{Bernoulli}(\tilde{\pi}_{kn}(\tilde{\pi}_{kn} + 1 - \pi_{kn})^{-1}) + \delta(x_{kn} \ge 1),$$

$$r_{k} \sim \text{Gamma}\left(1 + \sum_{n} u_{kn}, 1 - \sum_{n} h_{kn} \log(1 - b)\right),$$

$$z_{kn} \sim \delta(h_{kn} = 1) \text{Poisson}_{+}(\tilde{\lambda}_{kn}),$$

where $Poisson_{+}(\cdot)$ is the zero-truncated Poisson distribution and

$$x_{mk} = \sum_{n=1}^{N} x_{mkn},$$

$$x_{\cdot kn} = \sum_{m=1}^{M} x_{mkn},$$

$$\tilde{\pi}_{kn} = \pi_{kn} (1-b)^{r_k},$$

$$u_{kn} = \sum_{j=1}^{x_{\cdot kn}} u_{knj}, \qquad u_{knj} \sim \text{Bernoulli}\left(\frac{r_k}{r_k + j - 1}\right).$$
(1)

Note that for multilayer models, $\pi_{kn}^{(\ell)} = 1 - \exp(\lambda_{kn}^{(\ell+1)})$. The data augmentation scheme for r_k via u_{kn} is described in [1].

For the discriminative DPFA, lets denote latent counts for \hat{y}_n as \hat{x}_{ckn} , with summaries analogous to (1), as \hat{x}_{ck} and $\hat{x}_{\cdot kn}$. Then,

$$\mathbf{b}_{k} \sim \text{Dirichlet}(\zeta + \widehat{x}_{1k}, \dots, \zeta + \widehat{x}_{Ck}),$$

$$\theta_{kn} \sim \text{Gamma}(r_{k}h_{kn} + x_{kn} + \widehat{x}_{kn}, b^{-1}),$$

$$h_{kn} \sim \delta(x_{kn} = 0 \wedge \widehat{x}_{kn} = 0) \text{Bernoulli}(\widetilde{\pi}_{kn}(\widetilde{\pi}_{kn} + 1 - \pi_{kn})^{-1}) + \delta(x_{kn} \geq 1 \vee \widehat{x}_{kn} \geq 1).$$

Provided that θ_n and \mathbf{h}_n are shared by two PFAs, one for the count data, \mathbf{x}_n , and the other for the labels, \widehat{y}_n , their conditional posteriors are functions of latent counts coming from both sources, $x_{\cdot kn}$ and $\widehat{x}_{\cdot kn}$, respectively.

2.2 SVI

Variational parameter updates using (layer index omitted for clarity):

$$\begin{split} \phi_{mkn} &\propto \exp(\mathbb{E}[\log \psi_{mk}] + \mathbb{E}[\log \theta_{kn}]) \,, \\ \theta_{kn} &\sim \operatorname{Gamma}(\mathbb{E}[r_k]\mathbb{E}[h_{kn}] + \sum_{m=1}^{M} \phi_{mkn}, b^{-1}) \,, \\ h_{kn} &\sim \mathbb{E}[p(x_{\cdot kn} = 0)] \operatorname{Bernoulli}(\mathbb{E}[\tilde{\pi}_{kn}](\mathbb{E}[\tilde{\pi}_{kn}] + 1 - \mathbb{E}[\pi_{kn}])^{-1}) + \mathbb{E}[p(x_{\cdot kn} \ge 1)] \,, \\ r_k &\sim \operatorname{Gamma}\left(1 + \sum_n \mathbb{E}[u_{kn}], 1 - \sum_n \mathbb{E}[p(h_{kn} = 1)] \log(1 - b)\right) \,, \\ z_{kn} &\sim \mathbb{E}[p(h_{kn} = 1)] \operatorname{Poisson}_+(\tilde{\lambda}_{kn}) \,, \end{split}$$

where $\mathbb{E}[x_{mkn}] = \phi_{mkn}$, $\mathbb{E}[\tilde{\pi}_{kn}] = \mathbb{E}[\pi_{kn}](1-b)^{\mathbb{E}[r_k]}$ and $\mathbb{E}[u_{kn}] = \sum_{j=1}^{x_{\cdot kn}} \mathbb{E}[r_k](\mathbb{E}[r_k] + j - 1)^{-1}$.

List of Figures

1	Graphical models	1
2	Representative meta-topics obtained from 20 News	3
3	Graph representation obtained from 20 News	4
4	Graph representation obtained from RCV1	5
5	Graph representation obtained from Wiki	6
6	Representative meta-topics obtained from medical records data	7
7	Graph representation obtained from medical records data	8

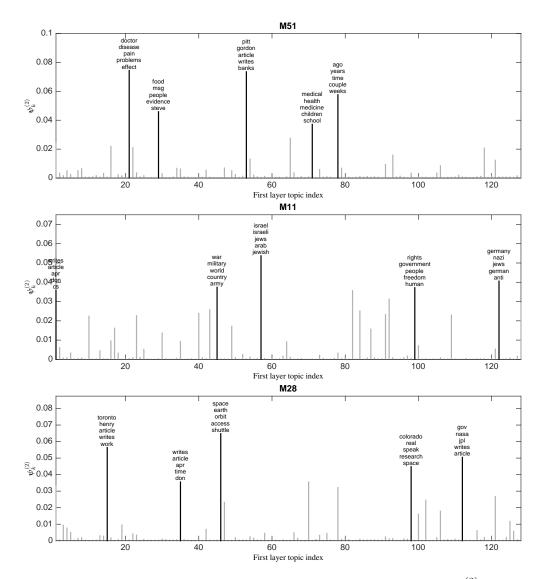


Figure 2: Representative meta-topics obtained from 20 News. Meta-topic weights $\psi_k^{(2)}$ vs. layer-1 topics indices, with word lists corresponding to the top four words in layer-1 topics, $\psi_k^{(1)}$.

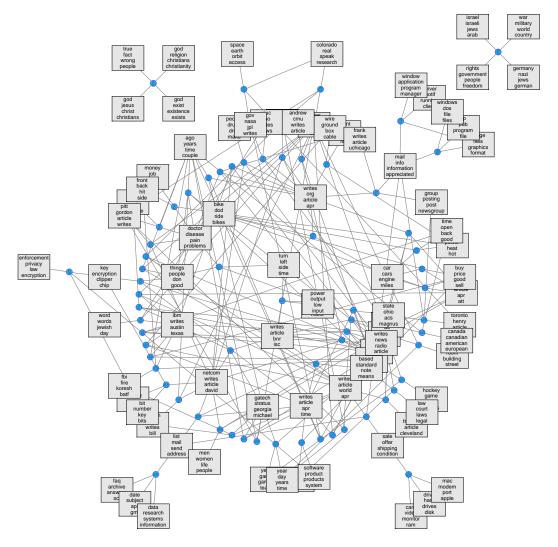


Figure 3: Graph representation obtained from 20 News. Meta-topics are denoted by circles and layer-1 topics as boxes, with word lists corresponding to the top four words in layer-1 topics, $\psi_k^{(1)}$. For clarity, we only show the top four connections between meta-topics and their associated topics

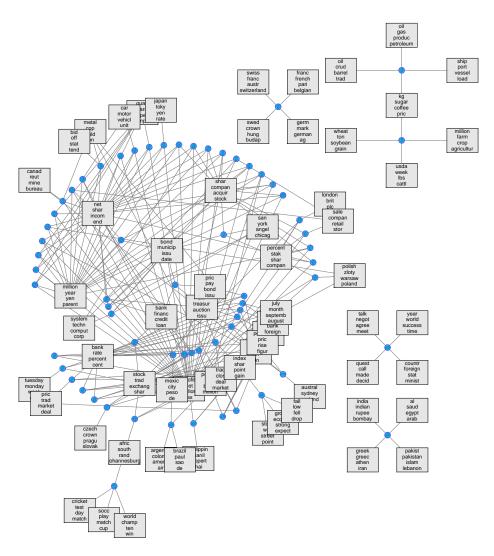


Figure 4: Graph representation obtained from RCV1. Meta-topics are denoted by circles and layer-1 topics as boxes, with word lists corresponding to the top four words in layer-1 topics, $\psi_k^{(1)}$. For clarity, we only show the top four connections between meta-topics and their associated topics.

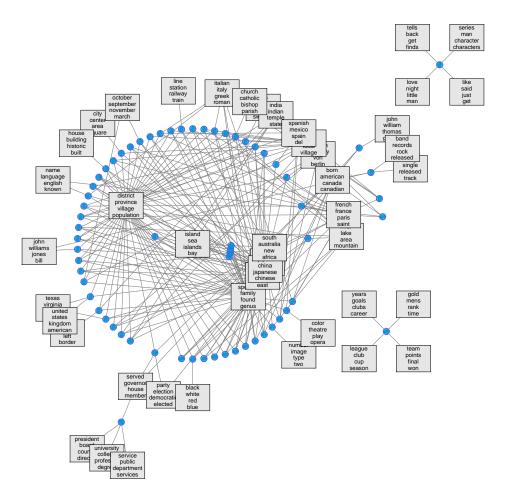


Figure 5: Graph representation obtained from Wiki. Meta-topics are denoted by circles and layer-1 topics as boxes, with word lists corresponding to the top four words in layer-1 topics, $\psi_k^{(1)}$. For clarity, we only show the top four connections between meta-topics and their associated topics.

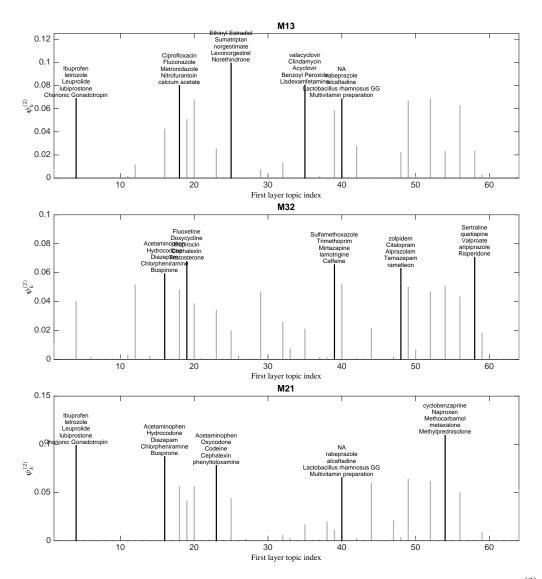


Figure 6: Representative meta-topics obtained from medical records data. Meta-topic weights $\psi_k^{(2)}$ vs. layer-1 topics indices, with word lists corresponding to the top four words in layer-1 topics, $\psi_k^{(1)}$.

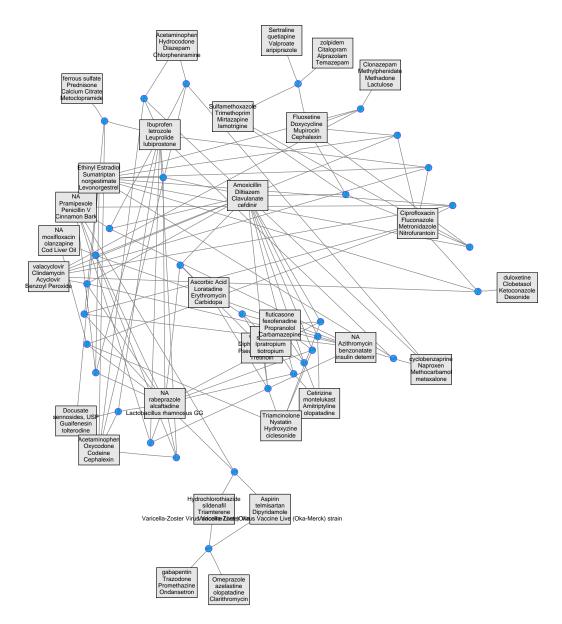


Figure 7: Graph representation obtained from medical records data. Meta-topics are denoted by circles and layer-1 topics as boxes, with word lists corresponding to the top four words in layer-1 topics, $\psi_k^{(1)}$. For clarity, we only show the top four connections between meta-topics and their associated topics.

References

[1] M. Zhou and L. Carin. Negative binomial process count and mixture modeling. *PAMI*, 2015.