

Stochastic Blockmodel with Cluster Overlap, Relevance Selection, and Similarity-Based Smoothing

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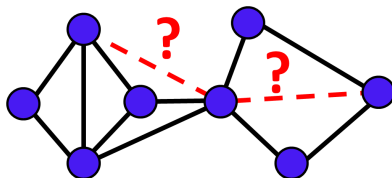
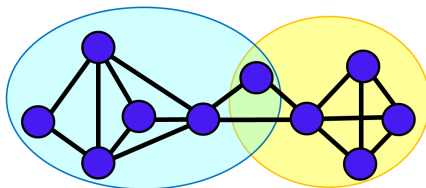
Introduction

- Stochastic Blockmodel

- Generative model
- Expresses objects as a low dimensional representation U_i, U_j
- Models the link probability of a pair of objects $P(A_{ij}) = f(U_i, U_j, \theta)$
- e.g., latent class model, mixed membership stochastic blockmodel

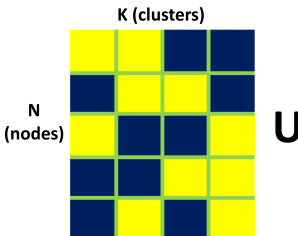
- Applications

- Revealing structures in networks
- (Overlapping) Clustering, Link prediction



Introduction

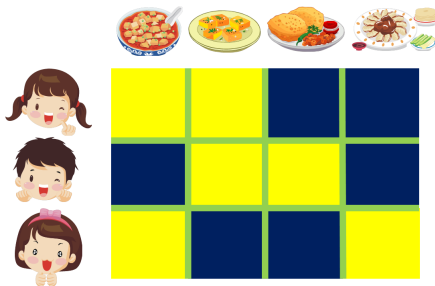
- Overlapping stochastic blockmodels
 - Objects have **hard** memberships in **multiple** clusters.



- Contributions of this paper
 - Extend the overlapping stochastic blockmodel to bipartite graphs
 - Relevance selection mechanism
 - Make use of additionally available object features
 - Nonparametric Bayesian approach

Background

- Indian Buffet Process (IBP) (Griffiths et al. 2011)
 - N objects, K clusters, overlapping clustering $\mathbf{U} \in \{0, 1\}^{N \times K}$.
 - Object: customer, cluster: dish
 - The first customer selects $Poisson(\alpha)$ dishes to begin with
 - Each subsequent customer n :
 - Selects an already selected dish k with probability $\frac{m_k}{n}$
 - Selects $Poisson(\alpha/n)$ new dishes



The Proposed Model

Basic Model

- Bipartite graph ($N \times M$ binary adjacency matrix, $|\mathcal{A}| = N$, $|\mathcal{B}| = M$)

$$\mathbf{U} \sim \text{IBP}(\alpha_u)$$

$$\mathbf{V} \sim \text{IBP}(\alpha_v)$$

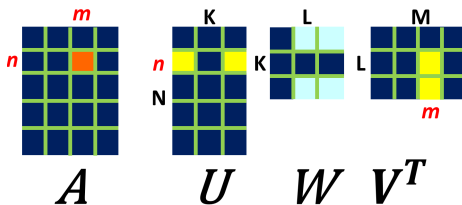
$$\mathbf{W} \sim \text{Nor}(0, \sigma_w^2)$$

$$\mathbf{A} \sim \text{Ber}(\sigma(\mathbf{U}\mathbf{W}\mathbf{V}^T))$$

- $\text{IBP}(\alpha)$: IBP prior distribution, $\text{Nor}(0, \sigma^2)$: Gaussian distribution,
- $\sigma(x) = \frac{1}{1 + \exp(-x)}$,
- $\text{Ber}(p)$: Bernoulli distribution,
- $\mathbf{U} \in \{0, 1\}^{N \times K}$, $\mathbf{V} \in \{0, 1\}^{M \times L}$: cluster assignment matrices

$$\begin{aligned} P(A_{nm} = 1) &= \sigma(\mathbf{u}_n \mathbf{W} \mathbf{v}_m^T) \\ &= \sigma\left(\sum_{k,l} u_{nk} W_{kl} v_{ml}\right) \end{aligned}$$

- W_{kl} : the interaction strength between two nodes due to their memberships in cluster k and cluster l



$$P(A_{nm} = 1) = \sigma(W_{12} + W_{13} + W_{32} + W_{33})$$

Basic Model

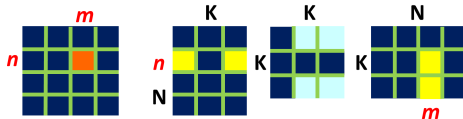
- Unipartite graph ($\mathbf{A} \in \{0, 1\}^{N \times N}$)

$$\mathbf{U} \sim \mathcal{IBP}(\alpha_u)$$

$$\mathbf{W} \sim \mathcal{Nor}(0, \sigma_w^2)$$

$$\mathbf{A} \sim \text{Ber}(\sigma(\mathbf{U}\mathbf{W}\mathbf{U}^\top))$$

$$\begin{aligned} P(A_{nm} = 1) &= \sigma(\mathbf{u}_n \mathbf{W} \mathbf{u}_m^\top) \\ &= \sigma\left(\sum_{k,l} u_{nk} W_{kl} u_{ml}\right) \end{aligned}$$



$$\mathbf{A} \quad \mathbf{U} \quad \mathbf{W} \quad \mathbf{U}^\top$$

- $\mathcal{IBP}(\alpha)$: IBP prior distribution,
 $\mathcal{Nor}(0, \sigma^2)$: Gaussian distribution,
- $\sigma(x) = \frac{1}{1 + \exp(-x)}$,
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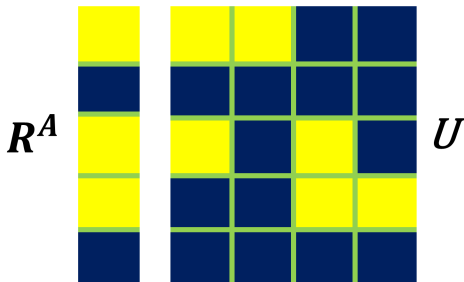
$$P(A_{nm} = 1) = \sigma(W_{12} + W_{13} + W_{32} + W_{33})$$

Relevance Selection Mechanism

- Motivation

- In real-world networks, there may be some noisy objects (e.g., spammer)
- May lead to bad parameter estimates

- Maintain two random binary vectors $\mathbf{R}^A \in \{0, 1\}^{N \times 1}$, $\mathbf{R}^B \in \{0, 1\}^{M \times 1}$



Relevance Selection Mechanism

- Background noise link probability $\phi \sim \text{Bet}(a, b)$
- If one or both objects $n \in \mathcal{A}$ and $m \in \mathcal{B}$ are irrelevant
 - A_{nm} is drawn from $\text{Ber}(\phi)$
- If both n and m are relevant,
 - A_{nm} is drawn from $\text{Ber}(p) = \text{Ber}(\sigma(\mathbf{u}_n \mathbf{W} \mathbf{v}_m^\top))$

$$\phi \sim \text{Bet}(a, b)$$

$$R_n^A \sim \text{Ber}(\rho_n^A), \quad R_m^B \sim \text{Ber}(\rho_m^B)$$

$$\mathbf{u}_n \sim \text{IBP}(\alpha_u) \quad \text{if } R_n^A = 1; \text{ zeros otherwise}$$

$$\mathbf{v}_m \sim \text{IBP}(\alpha_v) \quad \text{if } R_m^B = 1, \text{ zeros otherwise}$$

$$p = \sigma(\mathbf{u}_n \mathbf{W} \mathbf{v}_m^\top)$$

$$A_{nm} \sim \text{Ber}(p^{R_n^A R_m^B} \phi^{1 - R_n^A R_m^B})$$

Exploiting Pairwise Similarities

- We may have access to side information
 - e.g., a similarity matrix between objects
- The IBP does not consider the pairwise similarity information.
 - Customer n chooses an existing dish regardless of the similarity of this customer with other customers.
- Two objects n and m have a high pairwise similarity
 $\Rightarrow \mathbf{u}_n$ and $\mathbf{u}_m$ should also be similar.
 - Encourages a customer to select a dish if the customer has a high similarity with all other customers who chose that dish.
 - Let the customer select many new dishes if the customer has low similarity with previous customers.

Exploiting Pairwise Similarities

- Modify the sampling scheme in the IBP based generative model
 - The probability that object n gets membership in cluster k will be proportional to $\frac{\sum_{n' \neq n} S_{nn'}^A u_{n'k}}{\sum_{n'=1}^n S_{nn'}^A}$.

$\sum_{n'=1}^n S_{nn'}^A$: effective total number of objects,
 $\sum_{n' \neq n} S_{nn'}^A u_{n'k}$: effective number of objects (other than n) that belong to cluster k

- IBP: $\frac{\sum_{n' \neq n} u_{n'k}}{n} = \frac{m_k}{n}$

- The number of new clusters for object n is given by $\text{Poisson}(\alpha / \sum_{n'=1}^n S_{nn'}^A)$.

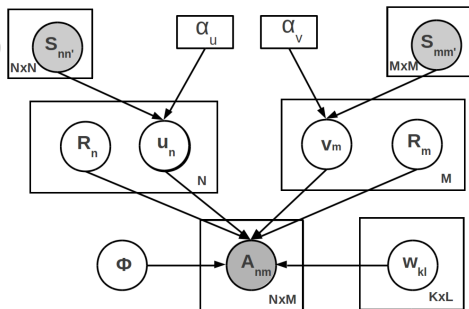
If the object n has low similarities with the previous objects, encourage it more to get memberships in its own new clusters

- IBP: $\text{Poisson}(\alpha/n)$

The Final Model

- **ROCS** (**R**elevance-based **O**verlapping **C**lustering with **S**imilarity-based-smoothing)

$$\begin{aligned}
 \phi &\sim \text{Bet}(a, b) \\
 \rho_n^A &\sim \text{Bet}(c, d), \quad \rho_m^B \sim \text{Bet}(e, f) \\
 R_n^A &\sim \text{Ber}(\rho_n^A), \quad R_m^B \sim \text{Ber}(\rho_m^B) \\
 \mathbf{u}_n &\sim \text{SimIBP}(\alpha_u, \mathbf{S}^A) \\
 \mathbf{v}_m &\sim \text{SimIBP}(\alpha_v, \mathbf{S}^B) \\
 p &= \sigma(\mathbf{u}_n \mathbf{W} \mathbf{v}_m^\top) \\
 A_{nm} &\sim \text{Ber}(p^{R_n^A R_m^B} \phi^{1-R_n^A R_m^B})
 \end{aligned}$$



- $\text{SimIBP}(\alpha_u, \mathbf{S}^A)$: similarity information augmented variant of the IBP

- For inference, we use MCMC (Gibbs sampling)

Experiments

Experiments

- Tasks

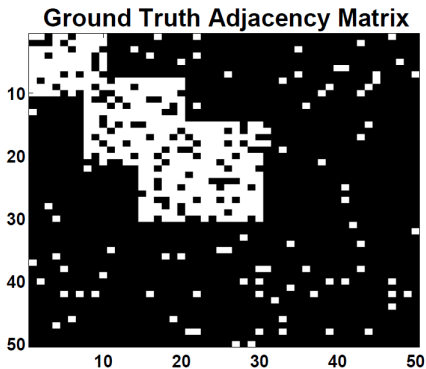
- The correct number of clusters
- Identify relevant objects
- Use pairwise similarity information
- Overlapping clustering
- Link prediction

- Baselines

- Overlapping Clustering using Nonnegative Matrix Factorization (OCNMF) (Psorakis et al. 2011)
- Kernelized Probabilistic Matrix Factorization (KPMF) (Zhou et al. 2012)
- Bayesian Community Detection (BCD) (Mørup et al. 2012)
- Latent Feature Relational Model (LFRM) (Miller et al. 2009)

Experiments

- Synthetic Data
 - 30 relevant objects, 20 irrelevant objects
 - Three overlapping clusters



Experiments

- Overlapping clustering

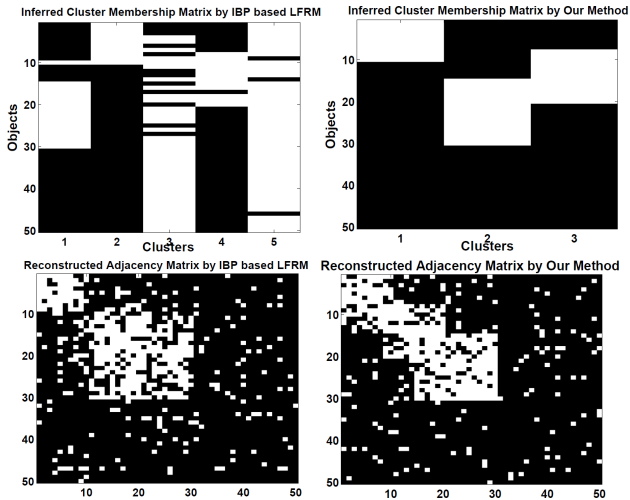


Table 1: Link Prediction on Synthetic Data

Method	0-1 Test Error (%)	AUC
OCNMF	44.82 (± 12.59)	0.7164 (± 0.1987)
KPMF	39.70 (± 1.78)	0.6042 (± 0.0517)
BCD	20.05 (± 1.49)	0.8504 (± 0.0197)
LFRM	9.59 (± 0.36)	0.8619 (± 0.0374)
ROCS	9.05 (± 0.42)	0.8787 (± 0.0303)

• Results Summary

- ROCS perfectly identifies relevant/irrelevant objects
- ROCS identifies the correct number of clusters
- For link prediction task, ROCS is better than other methods in terms of both 0-1 test error and AUC score.

Experiments

- Facebook Data

- An ego-network in Facebook (228 nodes)
- User profile (e.g., age, gender, etc.) – select 92 features.
- Known number of clusters: 14

Table 2: Link Prediction on Facebook Data

Method	0-1 Test Error (%)	AUC
OCNMF	36.58 (± 19.74)	0.7215 (± 0.1666)
KPMF	35.76 (± 2.76)	0.7013 (± 0.0174)
BCD	13.59 (± 0.31)	0.9187 (± 0.0242)
LFRM	12.38 (± 2.82)	0.9156 (± 0.0134)
ROCS	11.96 (± 1.44)	0.9388 (± 0.0156)

- BCD overestimated the number of clusters (20-22 across multiple runs).
- LFRM and ROCS almost correctly inferred the ground truth number of clusters (13-15 across multiple runs).

Experiments

- Drug-Protein Interaction Data
 - Bipartite graph (200 drug molecules, 150 target proteins)
 - Drug-drug similarity matrix, Protein-protein similarity matrix

Table 3: Link Prediction on Drug-Protein Interaction Data

Method	0-1 Test Error (%)	AUC
KPMF	16.65 ($\pm$ 0.36)	0.8734 ($\pm$ 0.0133)
LFRM	2.75 ($\pm$ 0.04)	0.9032 ($\pm$ 0.0156)
ROCS	2.31 ($\pm$ 0.06)	0.9276 ($\pm$ 0.0142)

- OCNMF and BCD are not applicable for bipartite graphs.
- LFRM here denotes ROCS without similarity information.
- KPMF takes into account the similarity information but does not assume overlapping clustering.

Experiments

- Lazega Lawyers Data
 - Directed graph, social networks (71 partners)
 - Each entry has features (gender, office-location, age, etc.)

Table 4: Link Prediction on Lazega-Lawyers Data

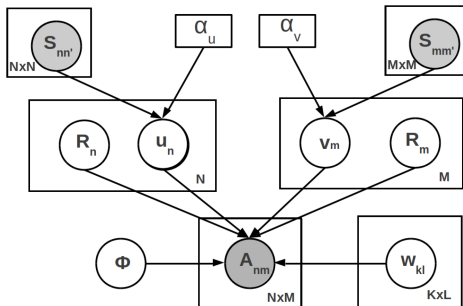
Method	0-1 Test Error (%)	AUC
OCNMF	35.36 (± 20.71)	0.6388 (± 0.1527)
KPMF	34.69 (± 1.13)	0.7203 (± 0.0229)
BCD	16.58 (± 0.56)	0.7876 (± 0.0168)
LFRM	14.05 (± 2.04)	0.8025 (± 0.0205)
ROCS	12.98 (± 0.32)	0.8248 (± 0.01642)

- Even weak similarity information can yield reasonable improvements in the prediction accuracy

Conclusions

Conclusions

- ROCS: a flexible model for modelling unipartite/bipartite graphs.
 - Each object can belong to multiple clusters (hard membership).
 - Nonparametric Bayesian approach.
 - Irrelevant objects can be dealt with in a principled manner.
 - Pairwise similarity between objects can be exploited to regularize the cluster memberships of objects.
 - Future work: make the model scalable.



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