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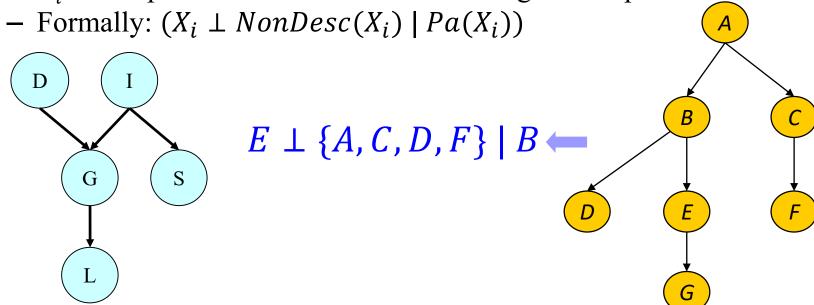
Chapter 3 Local Probabilistic Models

2021 Fall Jin Gu (古槿)

Bayesian Networks (Intuitive)

Can we find a simple graph model to equally or partially represent the probability with the same independences?

- Directed acyclic graph (DAG) G
 - Nodes X_1, \dots, X_n represent random variables
- G encodes local independence assumptions
 - $-X_i$ is independent of its non-descendants given its parents



Factorization Theorem ***

If we define the independences in G as $X_i \perp NonDesc(X_i) \mid Pa(X_i)$

• G is an I-Map of $P \to P(X_1,...,X_n) = \prod_{i=1}^n P(X_i | Pa(X_i))$

G is a given graph. If G is an I-Map of P, P can be factorized according to G.

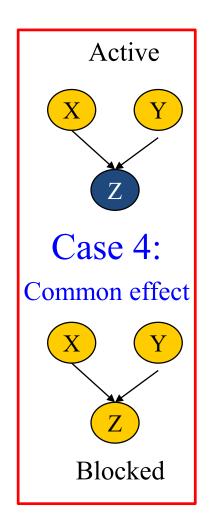
•
$$P(X_1,...,X_n) = \prod_{i=1}^n P(X_i | Pa(X_i)) \rightarrow G$$
 is an *I*-Map of *P*

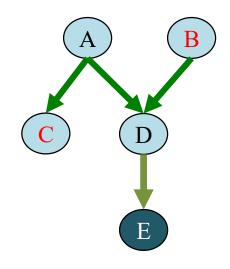
G is a given graph. If P can be factorized according to G, G is an I-Map of P.

Comments

- The Factorization Theorem Tells Us How to Deal With the Probability Based on Graph
 - Always put parent variables as the conditions
 - Represent you model according to logic order (generative model), but not always
- Any operation on graph is equal to the reasoning in original probabilistic problems
 - For example, d-Sep can find independences
 - Removal of edges means additional independence assumptions
- Why is the model called as "Bayesian" Network?
 - In real applications, we usually need to infer "reasons" based on the observed "results"
 - Another name is *belief* network

V-Structure Activation in BN





d-Sep(B,C|E) = no

Child or *its descendants* will activate v-structures in BNs (cause dependences)

Comments: trees have no v-structure!

"Independences" are NOT Enough

- As in the student example $I \rightarrow S$
- But what is P(I), and what is P(S|I)?

0.3

P(I,S)

I	5	P(I,S)
i ⁰	s^0	0.665
i ⁰	S^1	0.035
j ¹	s^0	0.06
j ¹	S ¹	0.24

Joint parameterization

P(I)

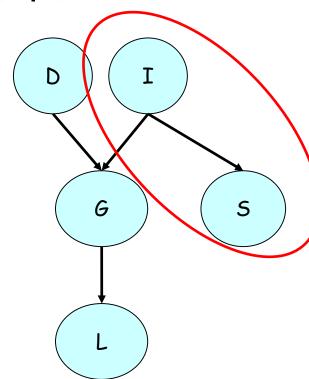
i⁰ 0.95 0.05 i¹ 0.2 0.8

P(S|I)

Conditional parameterization

Local Probabilistic Models
Conditional Probability Distributions (CPDs)

0.7

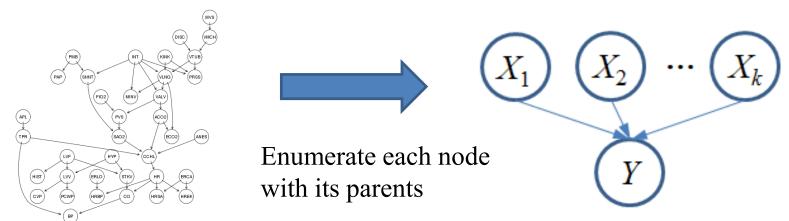


The Basic Structures in BNs

Recall the probability factorization over graph

$$P(X_1,...,X_n) = \prod_{i=1}^n P(X_i | Pa(X_i))$$

• All the required representations are the local conditional probability distributions (CPDs) encoded by the target variable given its parents



Outlines

- Local dependences in Bayesian networks
- Discrete CPDs
 - Tabular CPDs
 - Rule CPDs
 - Independence of causal influence
- Continuous CPDs
- A few CPDs in neural networks
- Bayesian networks representation examples

Textbook References

Textbook 1

- Chapter 5.1~5.4
- Chapter 5.6

Textbook 2

- Chapter 11 (Mixture Models)
- Chapter 12 (Latent Factor Models)

Tabular CPDs

- For a local structure: $(Y|X_1, X_2, \dots, X_k)$, if all variables are binary, there should be 2^k independent parameters.
 - -2^k different configurations for X_1, X_2, \dots, X_k
 - For each configuration for $X_1, X_2, \dots, X_k = \vec{x}$, we need $P(Y = y_0 | \vec{x})$
- The required parameters are exponentially increasing as the number of parents in BN local structures

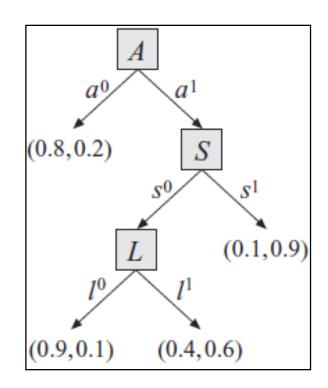
Rule CPDs

- We have a set of rules from the distribution
- For X = x and a configuration of a subset of its parents e, with a probability p, we can define a

RULE:

$$\rho:\langle e,x;p\rangle$$

 Tree CPDs can be directly transformed into a rule CPD



$$\rho_{1}:\langle a^{0}, j^{0}; 0.8 \rangle
\rho_{2}:\langle a^{0}, j^{1}; 0.2 \rangle
\rho_{3}:\langle a^{1}, s^{0}, l^{0}, j^{0}; 0.9 \rangle
\rho_{4}:\langle a^{1}, s^{0}, l^{0}, j^{1}; 0.1 \rangle
\rho_{5}:\langle a^{1}, s^{0}, l^{1}, j^{0}; 0.4 \rangle
\rho_{6}:\langle a^{1}, s^{0}, l^{1}, j^{1}; 0.6 \rangle
\rho_{7}:\langle a^{1}, s^{1}, j^{0}; 0.1 \rangle
\rho_{8}:\langle a^{1}, s^{1}, j^{1}; 0.9 \rangle$$

A rule-based CPD $P(X \mid Pa_X)$ is a set of rules \mathcal{R} such that:

- For each rule $\rho \in \mathcal{R}$, we have that $Scope[\rho] \subseteq \{X\} \cup Pa_X$.
- For each assignment (x, u) to $\{X\} \cup Pa_X$, we have precisely one rule $\langle c; p \rangle \in \mathcal{R}$ such that c is compatible with (x, u). In this case, we say that $P(X = x \mid Pa_X = u) = p$.
- The resulting CPD $P(X \mid U)$ is a legal CPD, in that

$$\sum_{x} P(x \mid u) = 1.$$

• Rule CPDs can model any complex CPDs, but not any rule CPD can be compactly transformed as a tree CPD. For example, no rule includes all the parent nodes.

COMMENTS: If you have big memory, use **hash table** to store rule-CPDs. You will get computational complexity O(1) for finding a target probability.

Independence of Causal Influence

- "Causal" / "Associative"
- The computer system will run down **if any one** of CPU, memory, hard disk, power or OS encounters some problems.
- The bank data center will run down if all the big computers run down at the same time.
- You will success if the addictive efforts of your body health, diligence, intelligence and persistence are enough.

Noisy-Or Model

- The failure rate of CPU is f_1
- The failure rate of MEM is f_2
- The failure rate of DISK is f_3
- The failure rate of POWER is f_4
- The failure rate of OS is f_5
- The failure rate of other events is f_0
- Question: the failure rate of your computer?

Noisy-Or Model

- Define $f_1 \sim f_k$ as noisy parameters (in normal condition, there should be no hardware / software failure, so we treat the failure as a noise)
- Define f_0 as leak probability

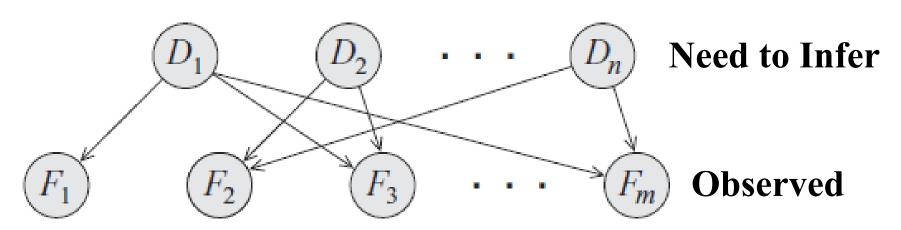
$$P(F = False) = (1 - f_0) \prod_{i} (1 - f_i)$$

$$P(F = True) = 1 - (1 - f_0) \prod_{i} (1 - f_i)$$

• Noisy-or: any parent can cause the same effect of its child with some probability

BN2O Networks

- Extension from Noisy-or model & Naive Bayes
- A disease can cause multiple phenotypes or symptoms, and multiple diseases may have share symptom(s).
- We can only observe symptoms, but we want to know diseases (most drugs are used for treating diseases).



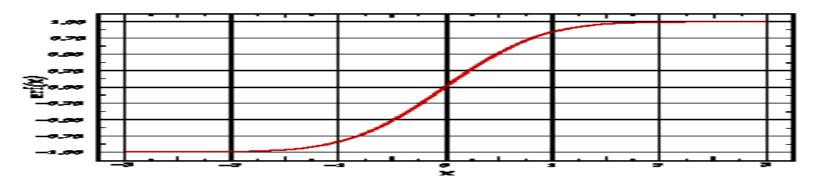
You need to specify the symptoms for each disease from big clinical data or medical instructions.

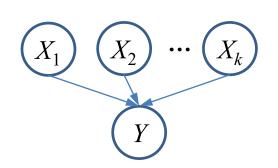
The Generalized Linear Models

• You will be more likely to be successful **if the summarized** score of the body health, diligence, intelligence, and persistence is higher.

$$S = w_h S_h + w_d S_d + w_i S_i + w_p S_p$$

$$P(success) = P(s) = ?$$
 Sigmoid Function!!

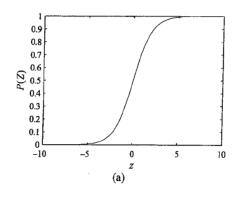


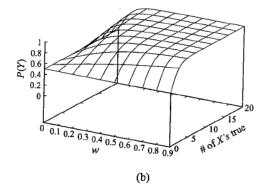


Logistic CPDs

• Sigmoid function

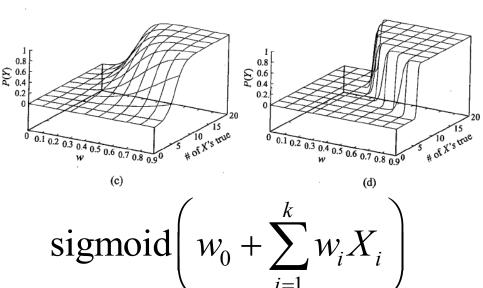
$$\operatorname{sigmoid}(s) = \frac{e^{s}}{1 + e^{s}}$$

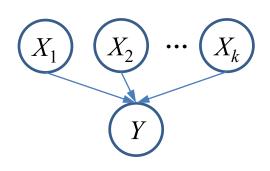




Logistic CPD

$$P(Y=y^1 \mid X_1, \cdots$$





Logistic CPDs

Odds

$$O = \frac{P(Y = y^1 \mid X_1, \dots)}{P(Y = y^0 \mid X_1, \dots)} e^{Z}$$

 Larger odds (>1) mean the configuration is more likely to cause the positive result

- The change of log odds of binary variables
 - -X are binary

$$\Delta O = \frac{O(X_{-j}, X_{j} = x_{j}^{1})}{O(X_{-j}, X_{j} = x_{j}^{0})} = e^{w_{j}}$$

• $w_j > 0$ means positive contribution

Generalized Linear Models

Linear Gaussian

$$-y = w_0 + \sum_{i=1}^k w_i x_i + \varepsilon, \varepsilon \in \mathcal{N}(0, \sigma^2)$$

• $p(y) \sim \mathcal{N}\left(w_0 + \sum_{i=1}^k w_i x_i, \sigma^2\right)$

• Binomial distribution (Logistic CPD)

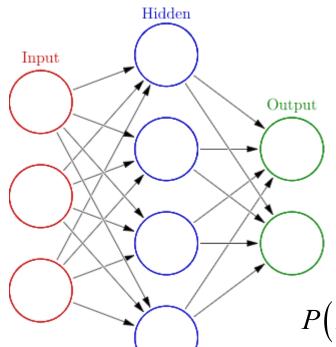
$$-P(y=1|x) = x, x = sigmoid(w_0 + \sum_{i=1}^k w_i x_i)$$

Poisson distribution

$$-P(y = k|X) = \lambda^k e^{-\lambda}/k!, \lambda = w_0 + \sum_{i=1}^k w_i x_i$$

Activation Function in NNs

Three-layer NNs



- Output of previous neurons
 - Activated $(x_i=1)$
 - Non-activated (x_i =0)
- Input of a neuron

$$-s_j = w_0 + \sum_i w_{i,j} x_i$$

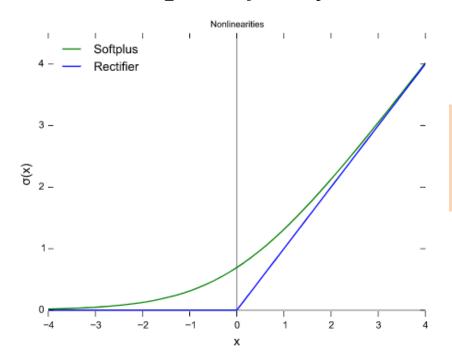
- Activation function
 - Sigmoid function

$$P(Y = y^1 | X_1, \dots$$
 sigmoid $\left(w_0 + \sum_{i=1}^k w_i X_i\right)$

Drawback: the gradient is too small during the two extremes of the sigmoid function!

ReLU: Rectified Linear Units

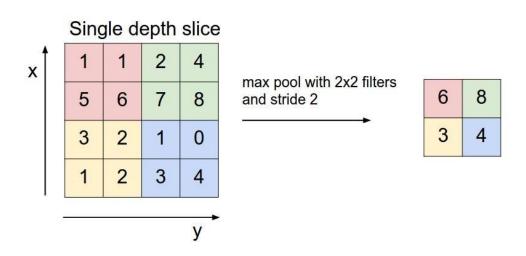
- Rectifier & softplus
 - Rectifier: $y = f(s) = \max(0, s)$
 - Softplus: $y = f(s) = \ln(1 + \exp(s))$



Comments: ReLU is the most used function for avoiding gradient dropping during backward propagation

Pooling Function

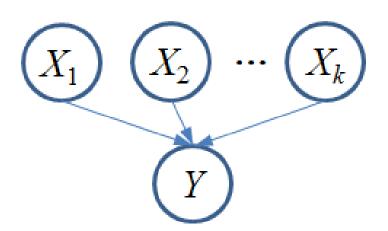
- Max pooling / median pooling (filtering)
- Advantages
 - Reduce noises for median pooling
 - Shift invariant for max pooling
 - − *Q*: How about rotation & inflation invariant?



The General Formulation

- Noisy-or model
- Generalized linear model
- We can significantly reduce the number of parameters for proper representation of CPDs

- Noisy-max model
- Noisy-and model
- •
- •



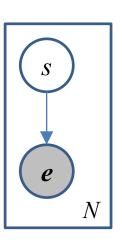
Summary: From I-Map to CPDs

- "Independences" are not enough to represent a probability.
- Bayesian Networks separate a large joint distribution into a set of local probabilistic models or conditional probability distributions.
 We further need above models to represent the local distributions efficiently.
- Local models are designed for $P(X|Pa_X)$

- Cancer is a general disease category consisting of many different types of diseases.
 For example, breast cancer has four major subtypes: normal-like, basal, luminal A and luminal B, with distinct clinical outcomes.

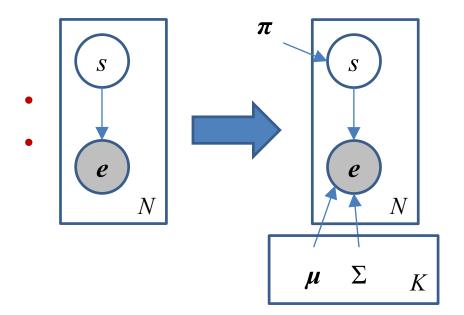
 Each subtype has different gene expression patterns.
 - We need to infer the subtypes based on the observed gene expressions.

- *Variables*: subtypes (s), gene expressions (e)
- *Relation*: subtypes define the patterns of expressions



- 1) *Circles* represent variables. *Shadow/white* circles represent observed/unobserved variables.
- 2) *Rectangle* indicates the variables in the same logic layer. The *number* (*N*) at the right-bottom corner represents the number of samples.

- *Variables*: subtypes (s), gene expressions (e)
- *Relation*: subtypes define the patterns of expressions
- Important: carefully check the independences in your model



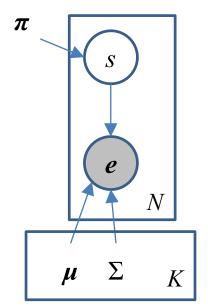
- Consider the local CPDs
- Different subtypes have different occurrence probability \rightarrow (s)

$$P(s=k) = \pi_i, \quad \sum_{k=1}^{K} \pi_i = 1$$

• Each expression pattern (indicated by i) independently follows different Gaussian distributions given the cancer subtype $\rightarrow (e|s)$

$$p(e | s = k) \sim N(\mu_k, \Sigma_k)$$

$$P(s=k) = \pi_i, \quad \sum_{k=1}^K \pi_i = 1$$



$$p(e | s = k) = N(\mu_k, \Sigma_k)$$

 How about the complete probabilistic model?

$$p(s = k, e) = P(s = k) p(e | s = k)$$

$$= \pi_k \times \frac{\exp\left(-\frac{1}{2}(e - \mu_k)^T \Sigma_k^{-1}(e - \mu_k)\right)}{\sqrt{(2\pi)^k |\Sigma_k|}}$$

Multiply all the CPDs in your model!!

$$p(S,E) = \prod_{n=1}^{N} p(s[n] = k, e[n])$$

For learning, you have N i.i.d. samples

- Cancer is usually caused by *independent* driving processes (factors), such as sustainable proliferation, resistance to cell death, immune escape and promoted vascular growth, etc.
- These driving processes have combined effects on the gene expressions patterns.
- We want to infer the driving processes based on large-scale gene expression datasets.

- *Variables*: driving factors (z_i) , gene expressions (e_i)
- *Relations*: *z* determines the distribution of *e*
- Local CPDs
 - Driving process z_i follows a standard Gaussian distribution

$$p(z_i) \sim N(0,1)$$
 $i = 1, \cdots$

Expression e_j also follows a
 Gaussian distribution but its
 mean is determined by a linear
 model of z

$$p(e_j | \mathbf{z}) \sim N\left(\alpha_0 + \sum_{i=1}^K \alpha_i z_i, \tau_j^2\right)$$

- Variables: independent driving processes (z_i) , gene expressions (e_i)
- *Relations*: *z* determines the distribution of *e*

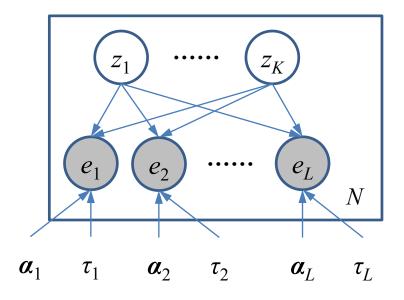
• Local CPDs

- Driving process z_i follows a standard Gaussian distribution

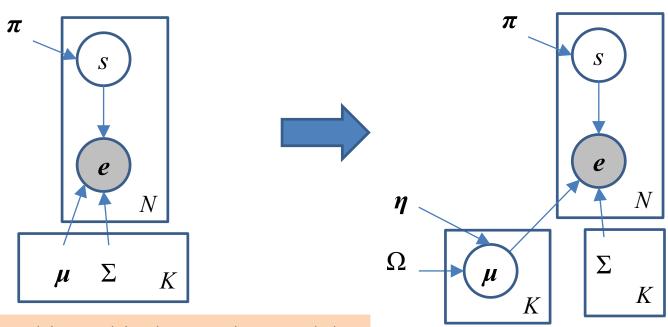
$$p(z_i) \sim N(0,1)$$
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- Expression e_j also follows a Gaussian distribution but its mean is determined by a linear model of z

$$p(e_j | \mathbf{z}) \sim N\left(\alpha_0 + \sum_{i=1}^K \alpha_i z_i, \tau_j^2\right)$$



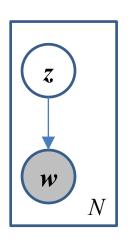
- Hierarchical Bayesian models
 - Key points: some parameters are random variables and further conditional on a few hyper-parameters



Comments: hierarchical Bayesian models have strong representation capability

- Topic Models in NLP
 - We have M documents
 - Each document has one or more topics (such as sport, finance, politics, entertainment, etc.)
 - Each topic has different word usage (such as Qian Yang, Ming Yao, Olympics, football for sport)
- Can we draw a model for learning the topics from the documents?

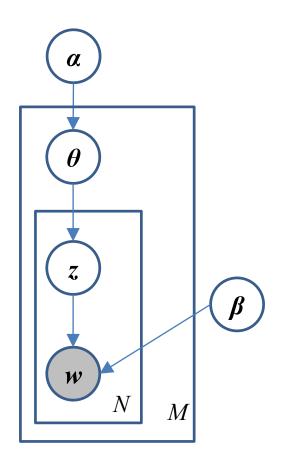
- Define variables for each document
 - Word (as binary vector): \mathbf{w}_n
 - Topic (as binary vector): \mathbf{z}_n
- Determine edges & local CPDs



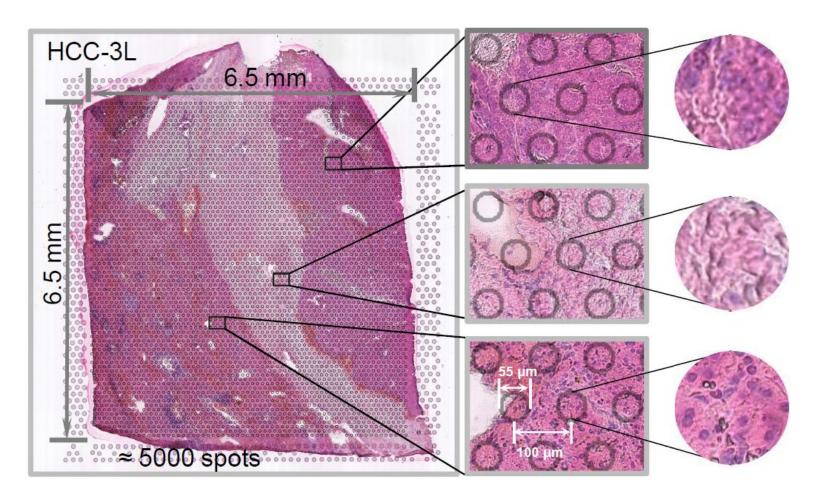
z follows a multinomial distribution

(the document generates one topic at a time)

w follows a multinomial distribution conditional on z (the topic generates one word, repeating N times)



Course Project: Multi-Model Spatial Information Modeling



Course Project: 2-Page Proposal

- Project selection
 - Assigned projects: segmentation of multi-model spatial information of tumor tissues
 - Task #1, use LDA or other models to cluster the spatial spots according to the expression data
 - Task #2, use MRF or other models to segment and partition the histological image
- Proposal deadline: November 8 (网络学堂)
- *Everyone* should submit his own proposal

Own Projects: Some Examples

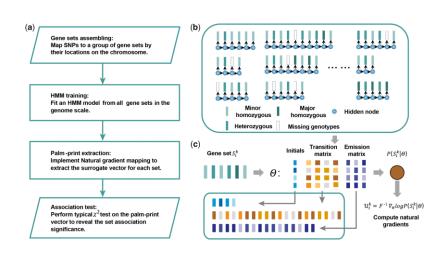


Fig. 4. Graphical representation of socoLDA. The Observed variables p, f, and r denote travel packages, co-travelers, and relationship types between users and co-travelers, separately. The hidden variables x and z denote topics assigned for travel packages and topics assigned for co-travelers, separately.

Hidden Markov models

Bao et al., Briefings in Bioinformatics, 2017

Extended Latent Dirichlet Allocation (LDA)

He et al., *Information & Management*, 2016

Methods & Theories

Project: Training deep generative models by MCEM

Image analysis

Project: 基于MRI图像的心脏功能辅助检测