Disease Prediction using Bayesian Networks

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Introduction

- ▶ Disease risk prediction
- Using diagnosis history
- Applications
 - ▶ Treatment and intervention
 - ► Health insurance risk management
 - ▶ Public Health research

Data

- ► Electronic Health records: Laboratory results, physician notes, medication, diagnosis codes
- ▶ ICD-9
 - ▶ 5 digit diagnosis codes
 - ► Primary and Secondary diagnosis
- ▶ ~9000 documented diseases

Challenges

- ▶ Limited publicly available datasets
- ▶ High Dimensional data
- ▶ 9000 variables
- ▶ Sparse data

Approach

- Clustering
 - ► Multiple specialized models
 - ▶ K-Means for categorical data
- ▶ Learning
 - ▶ Structure
 - Parameters
- Prediction
 - ▶ Risk for a disease
 - ▶ Most likely disease

Structure Learning

- ► Search Optimization
- ▶ Model selection
 - ▶ BIC criteria
 - ► Prefer simpler models
- ▶ Hill Climbing
 - ► Greedy Search
 - ▶ Adjacent space: Add, delete, reverse edges
 - ► Random restarts

Structure learning

- Sparse Candidate algorithm
- ► Each node has at most k parents
- Mutual Information

$$I(X;Y) = \sum_{x,y} \hat{P}(x,y) \log \frac{\hat{P}(x,y)}{\hat{P}(x)\hat{P}(y)}$$

Prediction

- Assign a test patient to a cluster
- Approximate inference of conditional probabilities
- ▶ Monte Carlo simulations
- Generate samples from distributions
- Calculate conditional probabilities by counting

References

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