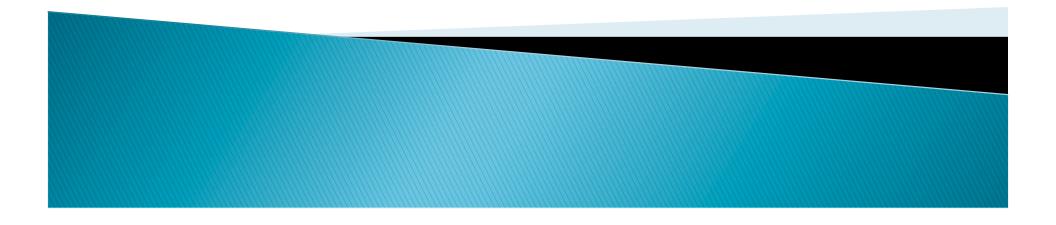
An Introduction of Some Methods in Learning Gene Association Networks Bo Li



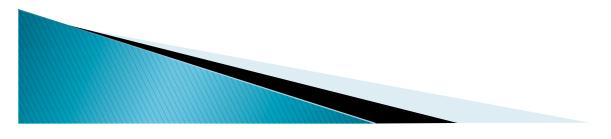
Overview

- Motivation
- Naïve Method
- Graphical Lasso



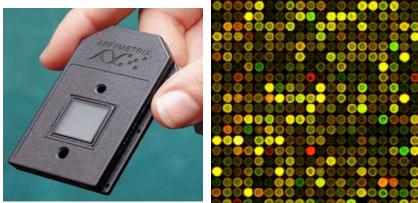
Motivation

- Genes are never isolated
- Want to study them in a systematic way
- A good way to look at their interactions : build gene association networks



Data

- Microarrays
 - Traditional technology
 - Data are noisy





- Appears recently
- Data are more accurate



*Pictures are found from internet, unauthorized

Data

• Data Matrix X $X \in R^{p \times n}$

- p : number of genes
- n : number of samples

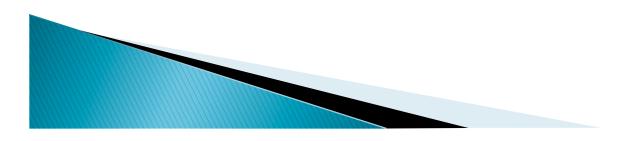


Goal

Build a undirected graph

Describing relationship between genes

As simple as possible



Naïve Method

For each pair of genes

Calculate Pearson Correlation Coefficient

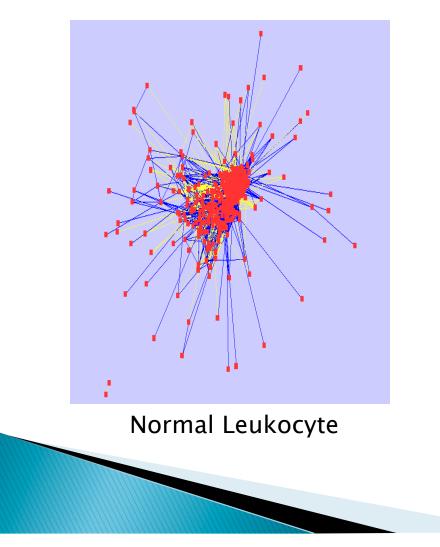
$$\rho(\mathbf{X},\mathbf{Y}) = \frac{\sum_{i=1}^{n} (x_i - \bar{\mathbf{x}}) (y_i - \bar{\mathbf{y}})}{\sqrt{\sum_{i=1}^{n} (x_i - \bar{\mathbf{x}})^2} \sqrt{\sum_{i=1}^{n} (y_i - \bar{\mathbf{y}})^2}}$$

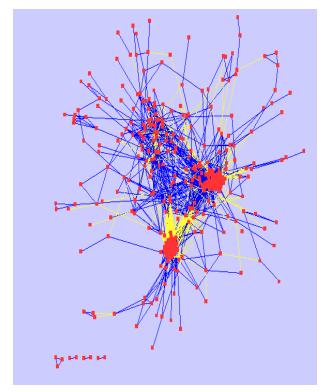
- Calculate P value
 - Permutation Tests
 - t-distribution
- Control False Discovery Rate (FDR)



Example Networks

Based on 431 mitosis cell cycle related microarray probe sets

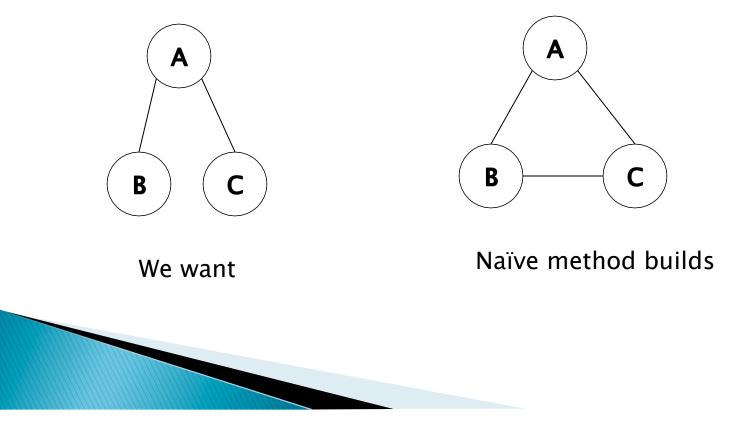




Acute myeloid leukemia

Drawback

- If B, C depend on A, but are conditionally independent given A
 - Say gene A regulates gene B and C
 - Naïve method tends to connect B and C, too!



Gaussian Graphical Model

• Assume each sample is sampled from a multivariate Gaussian distribution $N(\mu, \Sigma)$.

$$N(x|\mu, \Sigma) = \frac{1}{(2\pi)^{p/2}} \frac{1}{|\Sigma|^{1/2}} \exp\left\{-\frac{1}{2}(x-\mu)^{T}\Sigma^{-1}(x-\mu)\right\}$$

- Let Θ = Σ⁻¹. This is precision matrix
 If θ_{ij} is 0, variable i & j are conditionally independent given all other variables
 - Otherwise, connect i and j in the graph



Graphical Lasso [Friedman et al. 2008]

- Friedman, J., Hastie, T. and Tibshirani, R. (2008). Sparse inverse covariance estimation with the graphical lasso. *Biostatistics* 9, 432–441.
- Assume μ is fixed, say $\mu = 0$ Let $S = -\frac{1}{n}XX^{T}$
- Want to maximize penalized loglikelihood

 $\log \det \Theta - \operatorname{tr}(S\Theta) - \rho \|\Theta\|_1$

- Define $W = \Theta^{-1}$
- The subgradient equation is

$$W - S - \rho \cdot \Gamma = 0$$

$$\Gamma_{ij} = \begin{cases} sgn(\theta_{ij}), & \theta_{ij} \neq 0 \\ \gamma_{ij} \in [-1,1], & \theta_{ij} = 0 \end{cases}$$

Partition W as follows:

$$\begin{pmatrix} W_{11} & W_{12} \\ W_{12}^T & W_{22} \end{pmatrix} \begin{pmatrix} \Theta_{11} & \theta_{12} \\ \theta_{12}^T & \theta_{22} \end{pmatrix} = \begin{pmatrix} I & 0 \\ 0^T & 1 \end{pmatrix}$$

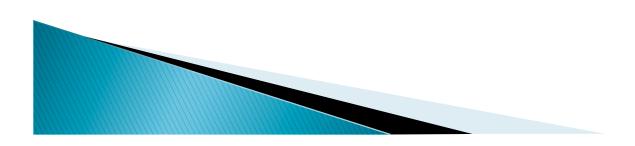
• Assume W_{11} is known, we can find W_{12} by solving

$$\min_{\beta} \{ \frac{1}{2} \left\| W_{11}^{1/2} \beta - W_{11}^{-1/2} s_{12} \right\|^2 + \rho \|\beta\|_1 \} \quad (*)$$

• w_{12} is given by

 $w_{12} = W_{11}\hat{eta}, \qquad \hat{eta}$ solves (*).

(*) is a lasso problem!



 To show it, the upper right block of penalized log-likelihood's subgradient equation is

$$w_{12} - s_{12} - \rho \cdot \gamma_{12} = 0$$

The subgradient equation of (*) is

$$W_{11}\beta - s_{12} + \rho \cdot \nu = 0$$

$$v_i = \begin{cases} sgn(\beta_i), & \beta_i \neq 0\\ v_i \in [-1,1], & \beta_i = 0 \end{cases}$$

- Algorithm
 - Start with $W = S + \rho I$
 - For j = 1,2,...,p,1,2,...,p
 - Permute jth diagonal item to the lower right corner of W
 - Solve the lasso problem (*) and get $W_{12} = W_{11}\hat{eta}$
 - Recover the corresponding row and column with w_{12}
 - Continue until convergence



Some Related References

P. Ravikumar, G. Raskutti, M. Wainwright, B. Yu (2008) Model selection in Gaussian graphical models: highdimensional consistency of 11-regularized MLE. *In Advances in Neural Information Processing Systems* (NIPS) 21, 2008.





Thanks!

