

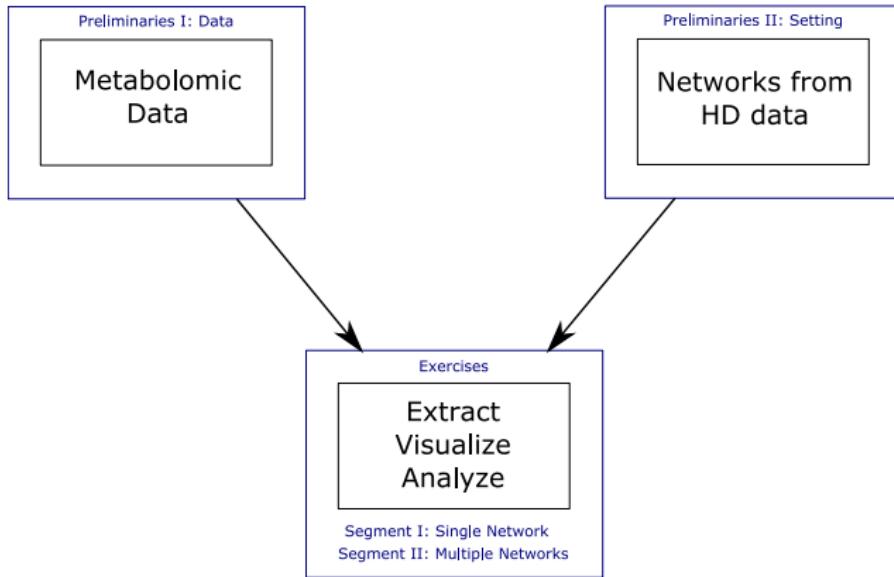
# rags2ridges

## Network Modeling of High-Dimensional Data

Carel F.W. Peeters  
Dept. of Epidemiology & Biostatistics  
VU University medical center, Amsterdam  
[cf.peeters@vumc.nl](mailto:cf.peeters@vumc.nl)

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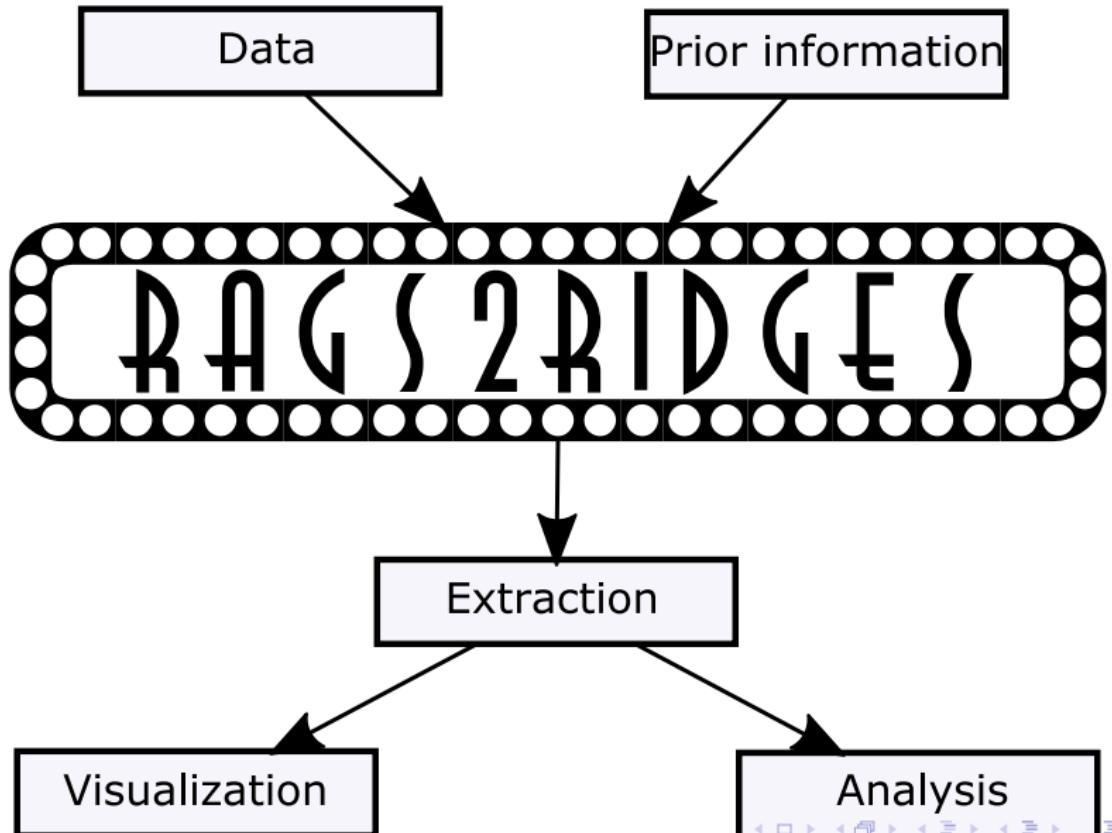
# Overview



## Materials

- These slides: Contain explanation theory/methodology and exercises
- `rags2ridges_Practical.HTML`: Contains solutions to exercises with additional explanations
- `NetworksExercises.R`: Contains bare solutions to exercises

## rags2ridges: One-Stop-Go



# Omics and omics data

## -ome

A totality of some (molecular biological) sort

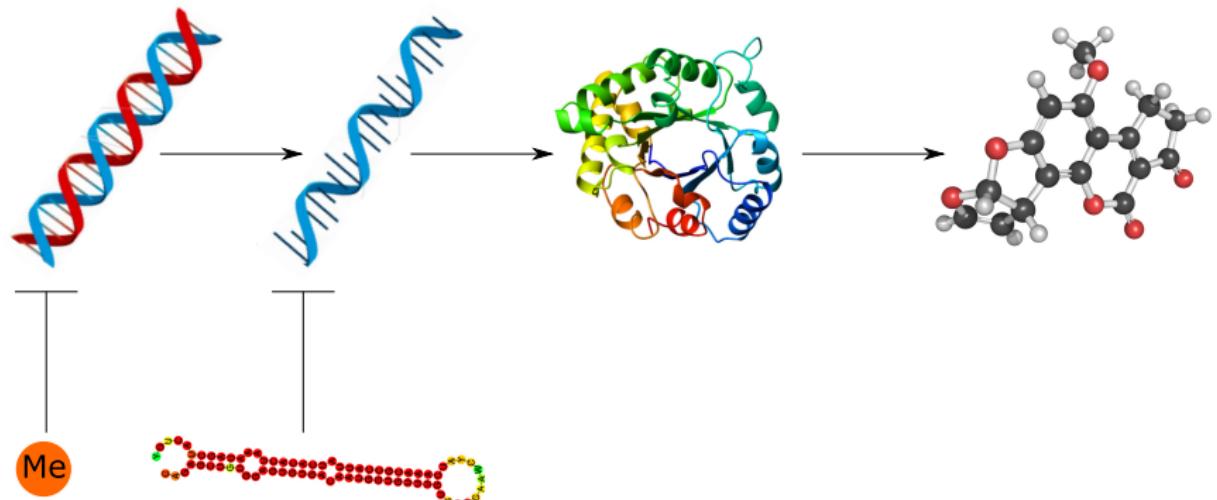
## -omics

Collective quantification of some pool of molecular molecules

## Genomics

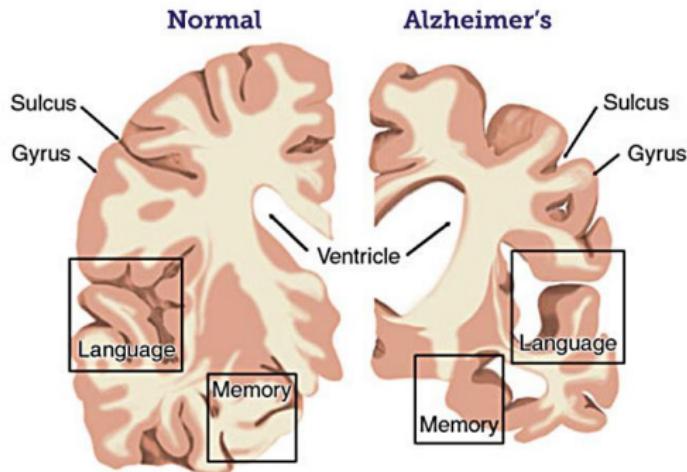
The omics of the genome (of some organism)

# The omic cascade



# Alzheimer's Disease

## Brain Cross-Sections



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# Metabolite quantification

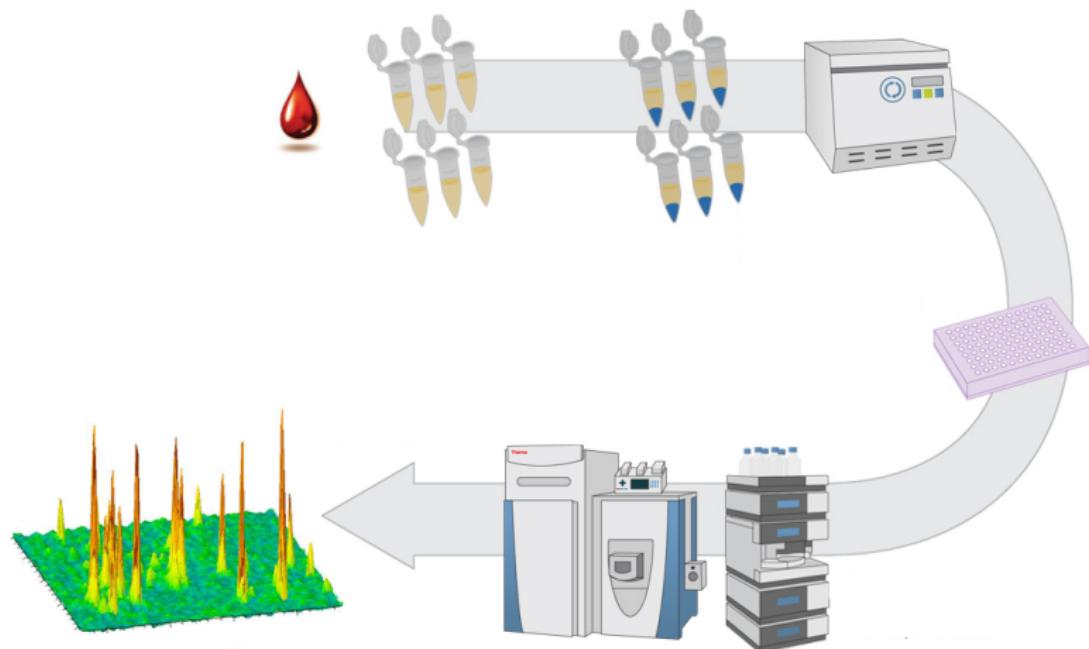


Illustration adapted from: <http://planetorbitrap.com/untargeted-metabolomics#.Vzw6yfmLRaQ> &  
<http://metabolomicsplatform.com/metabolomics-overview/>

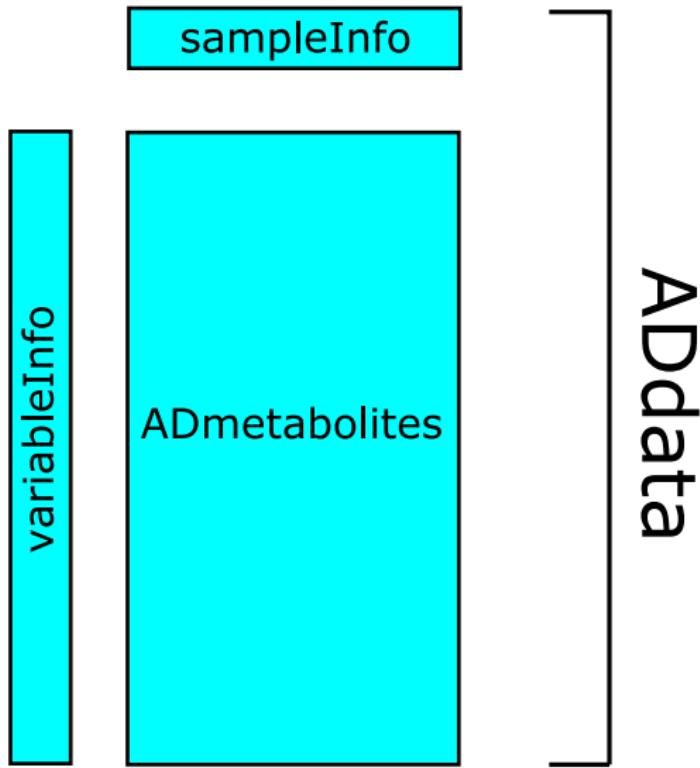
# Metabolomic Data

	Variables (features)						
	1	2	3	4	5	.....	p
1	$y_{11}$	$y_{12}$	$y_{13}$	$y_{14}$	$y_{15}$	.....	$y_{1p}$
2	$y_{21}$	$y_{22}$	$y_{23}$	$y_{24}$	$y_{25}$	.....	$y_{2p}$
3	$y_{31}$	$y_{32}$	$y_{33}$	$y_{34}$	$y_{35}$	.....	$y_{3p}$
⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮
⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮
⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮
n	$y_{n1}$	$y_{n2}$	$y_{n3}$	$y_{n4}$	$y_{n5}$	.....	$y_{np}$

- Amines
- Lipids
- Organic Acids
- Oxidative Stress

$p = 230$ : 53 Amines, 116 Lipids, 22 Organic acids, 39 Oxidative stress compounds  
 $n = 127$ : 40 AD class 1, 87 AD class 2

# Metabolomic Data Objects



## Exercise 1: Get acquainted with the data

```
## Set working directory  
setwd("")
```

```
## Needed libraries  
library(rags2ridges)
```

Invoke data

```
load()
```

Which objects?

```
objects()
```

Simple exploration objects

```
head()
```

# Gaussian graphical modeling

## Graphical modeling

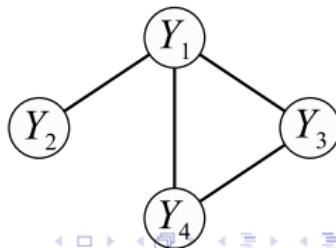
Class of models using graphs to express conditional (in)dependence relations between random variables

## Gaussian setting

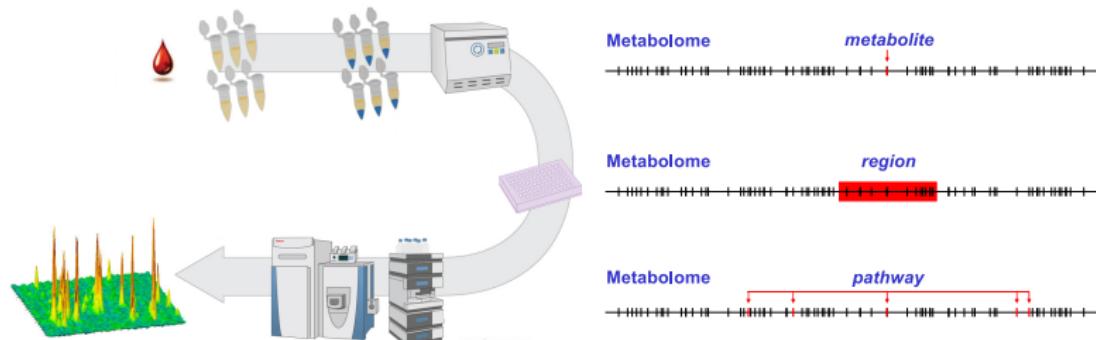
- Vertices: Correspond to random variables with normal distribution
- Edges: Correspond to the dependence structure
- Say  $\mathbf{y} \sim \mathcal{N}_p(\mathbf{0}, \boldsymbol{\Sigma})$ , and define  $\boldsymbol{\Sigma}^{-1} \equiv \boldsymbol{\Omega}$ . Then, for  $a, b \in$  vertex set  $V$ ,  $a \neq b$

$$-\frac{\omega_{ab}}{\sqrt{\omega_{aa}\omega_{bb}}} = 0 \iff \omega_{ab} = 0 \iff a \perp\!\!\!\perp b | V \setminus \{a, b\} \iff a \not\sim b$$

$$\begin{bmatrix} \omega_{11} & \omega_{12} & \omega_{13} & \omega_{14} \\ \omega_{21} & \omega_{22} & 0 & 0 \\ \omega_{31} & 0 & \omega_{33} & \omega_{34} \\ \omega_{41} & 0 & \omega_{43} & \omega_{44} \end{bmatrix}$$



# Challenge



## Problem

- Let  $\mathbf{S} \equiv \hat{\Sigma}$  denote the sample covariance matrix on  $\mathbf{y}_i$
- When  $p \approx n$  or  $p > n$ ,  $\mathbf{S}$  is ill-behaved or singular
- The precision  $\mathbf{S}^{-1} \equiv \hat{\Omega}$  is then undefined

## $\ell_2$ -Penalization: Proper ridge

Maximize

$$\underbrace{\ln |\Omega| - \text{tr}(\mathbf{S}\Omega)}_{\text{log-likelihood}} - \underbrace{\frac{\lambda}{2} \|\Omega - \mathbf{T}\|_2^2}_{\ell_2\text{-penalty}}$$

- $\mathbf{T}$  denotes a p.d. symmetric target matrix
- $\lambda \in (0, \infty)$  denotes a penalty parameter

Analytic penalized ML estimator

$$\hat{\Omega}(\lambda) = \left\{ \left[ \lambda \mathbf{I}_p + \frac{1}{4} (\mathbf{S} - \lambda \mathbf{T})^2 \right]^{1/2} + \frac{1}{2} (\mathbf{S} - \lambda \mathbf{T}) \right\}^{-1}$$

# Properties

## Behavior

- i.  $\hat{\Omega}(\lambda) \succ 0$ , for all  $\lambda \in (0, \infty)$ ;
- ii.  $\lim_{\lambda \rightarrow 0^+} \hat{\Omega}(\lambda) = \mathbf{S}^{-1}$  if  $p < n$ ;
- iii.  $\lim_{\lambda \rightarrow \infty} \hat{\Omega}(\lambda) = \mathbf{T}$ .

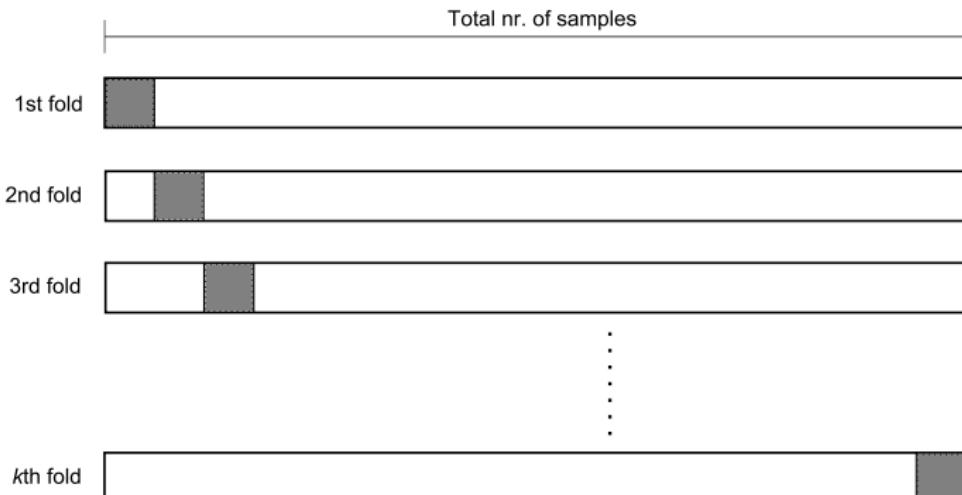
## Consistency

- i.  $\lim_{n \rightarrow \infty} \mathbb{E} [\hat{\Omega}_n(\lambda_n)] \longrightarrow \lim_{n \rightarrow \infty} \mathbb{E} (\mathbf{S}_n^{-1}) = \Omega$ ;
- ii.  $\lim_{n \rightarrow \infty} \mathbb{E} (\|\hat{\Omega}_n(\lambda_n) - \Omega\|_F^2) = 0$ .

# Choosing the penalty value

## K-fold cross-validation (CV)

### Single iteration of K-fold CV



Test data



Training data

## Choosing the penalty value

*K*-fold CV score

$$\varphi^K(\lambda) = \sum_{k=1}^K n_k \left\{ -\ln |\hat{\Omega}(\lambda)_{-k}| + \text{tr}[\hat{\Omega}(\lambda)_{-k} \mathbf{S}_k] \right\},$$

$n_k$  is the size of subset  $k$ , for  $k = 1, \dots, K$  disjoint subsets;

$\mathbf{S}_k$  denotes the sample covariance matrix on  $k$ th test set;

$\hat{\Omega}(\lambda)_{-k}$  denotes the estimated regularized precision matrix on  $k$ th training set

Choose

$$\lambda^* = \arg \min_{\lambda \in \mathbb{R}^+} \varphi^K(\lambda)$$

Efficiency

- Implementation uses C++ at its core
- Makes use of a root-finding (Brent) algorithm
- Utilizes rotational equivariance property when possible

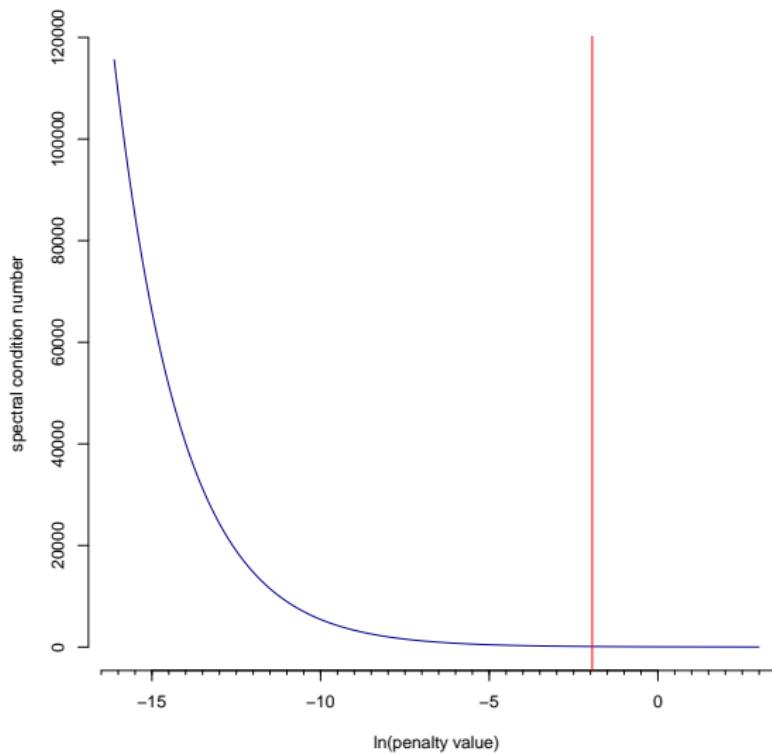
## Exercise 2: Find an optimal precision matrix for the Class 2 AD data

```
optPenalty.kCVauto(Y,           ← data (matrix)
                     lambdaMin,   ← min. λ
                     lambdaMax,   ← max. λ
                     target)      ← T (use default.target())
```

Returns list object

- \$optLambda: Optimal penalty parameter
- \$optPrec: Precision estimate under optimal penalty parameter

# Assessing the Conditioning of the Estimate: Condition Number Plot

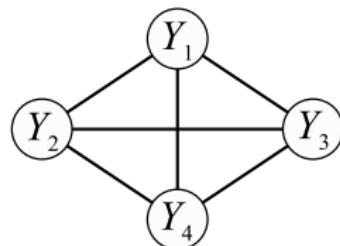


## Exercise 3: Assess the conditioning of the optimal precision matrix

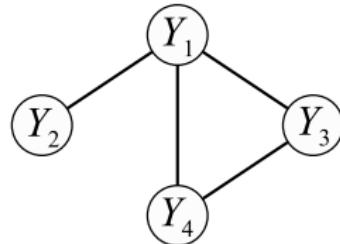
```
CNplot(S,           ← covariance matrix data
       lambdaMin,   ← min. λ
       lambdaMax,   ← max. λ
       step,        ← coarseness grid
       target,      ← T
       vertical,    ← logical
       value)
```

# Support Determination

$$\begin{bmatrix} \omega_{11} & \omega_{12} & \omega_{13} & \omega_{14} \\ \omega_{21} & \omega_{22} & \omega_{23} & \omega_{24} \\ \omega_{31} & \omega_{32} & \omega_{33} & \omega_{34} \\ \omega_{41} & \omega_{42} & \omega_{43} & \omega_{44} \end{bmatrix}$$



$$\begin{bmatrix} \omega_{11} & \omega_{12} & \omega_{13} & \omega_{14} \\ \omega_{21} & \omega_{22} & 0 & 0 \\ \omega_{31} & 0 & \omega_{33} & \omega_{34} \\ \omega_{41} & 0 & \omega_{43} & \omega_{44} \end{bmatrix}$$



## Support determination

### Scaling

$\hat{\mathbf{P}}(\lambda)$ : Regularized precision estimate scaled to partial correlation form

### Assume

Nonredundant coefficients (indexed by  $j < j'$ ) follow a mixture distribution:

$$f \left\{ [\hat{\mathbf{P}}(\lambda^*)]_{jj'} \right\} = \eta_0 f_0 \left\{ [\hat{\mathbf{P}}(\lambda^*)]_{jj'}; \kappa \right\} + (1 - \eta_0) f_{\mathcal{E}} \left\{ [\hat{\mathbf{P}}(\lambda^*)]_{jj'} \right\}$$

- $\eta_0 \in [0, 1]$  is the mixture weight
- $f_0\{\cdot\}$  denotes the distribution of a null-edge
- $f_{\mathcal{E}}\{\cdot\}$  denotes the distribution of a present edge
- $\kappa$  denotes degrees of freedom

### Determine

$$P(Y_j \neq Y_{j'} | [\hat{\mathbf{P}}(\lambda^*)]_{jj'}) = \frac{\hat{\eta}_0 f_0 \left\{ [\hat{\mathbf{P}}(\lambda^*)]_{jj'}; \hat{\kappa} \right\}}{\hat{\eta}_0 f_0 \left\{ [\hat{\mathbf{P}}(\lambda^*)]_{jj'}; \hat{\kappa} \right\} + (1 - \hat{\eta}_0) \hat{f}_{\mathcal{E}} \left\{ [\hat{\mathbf{P}}(\lambda^*)]_{jj'} \right\}}$$

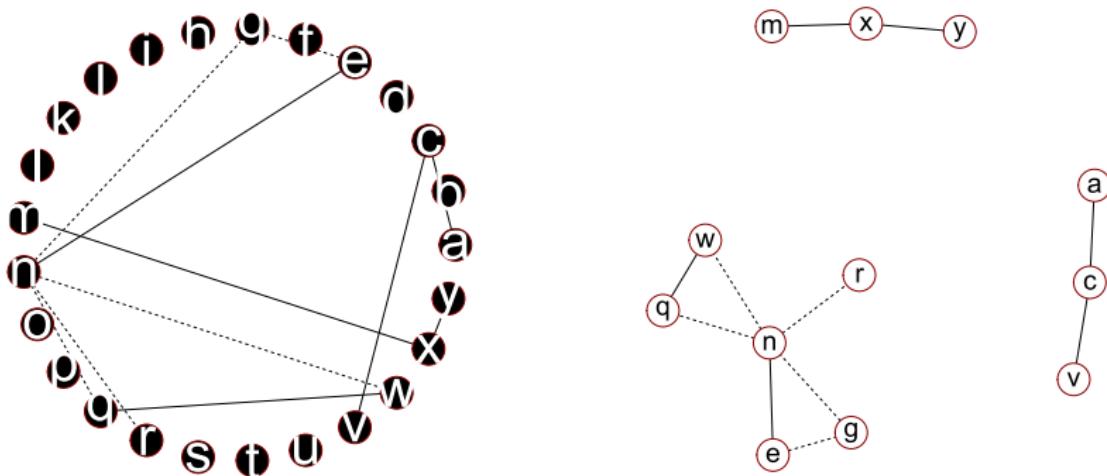
## Exercise 4: Extract a network from the optimal precision matrix by retaining elements whose posterior probability of being present $\geq .999$

```
sparsify(P,           ← estimated precision matrix  
        threshold, ← type: "localFDR", "top"  
        FDRcut)    ← cut-off for 1 - IFDR
```

Returns list object

- \$sparsePrecision: Sparsified precision matrix
- \$sparseParCor: Sparsified partial correlation matrix

# Visualization



## Things to Consider

- Layout
- Size of vertices
- colorings
- ...

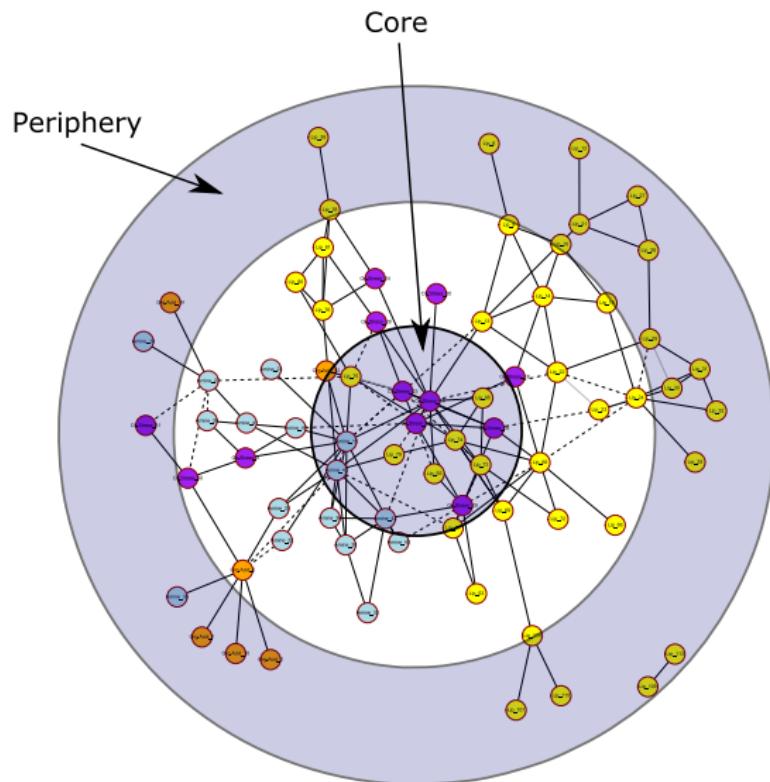
## Exercise 5: Visualize the extracted network

```
Ugraph(M,  
       type, ← "plain", "fancy", "weighted"  
       lay, ← specifies layout  
       Vsize, ← size of vertices  
       Vcex, ← size vertex labels  
       Vcolor, ← vertex color  
       VBcolor, ← vertex boundary color  
       VLcolor, ← color vertex labels  
       prune, ← removes unconnected vertices?  
       ...)
```

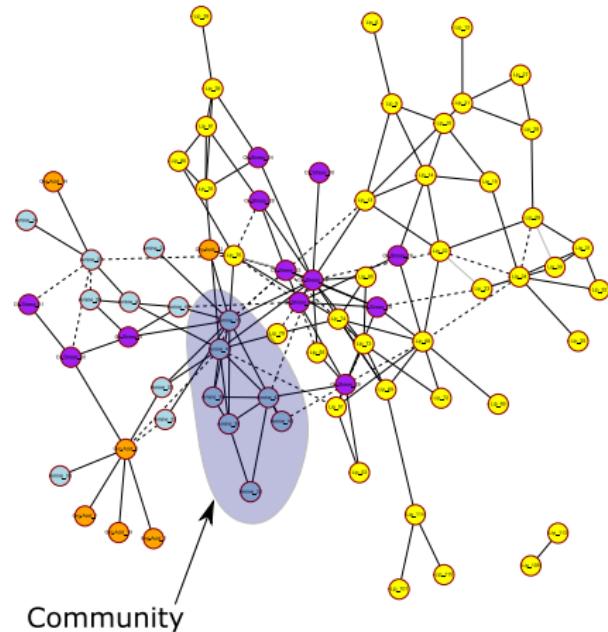
Returns matrix object

Containing the coordinates of the vertices in the given graph/network

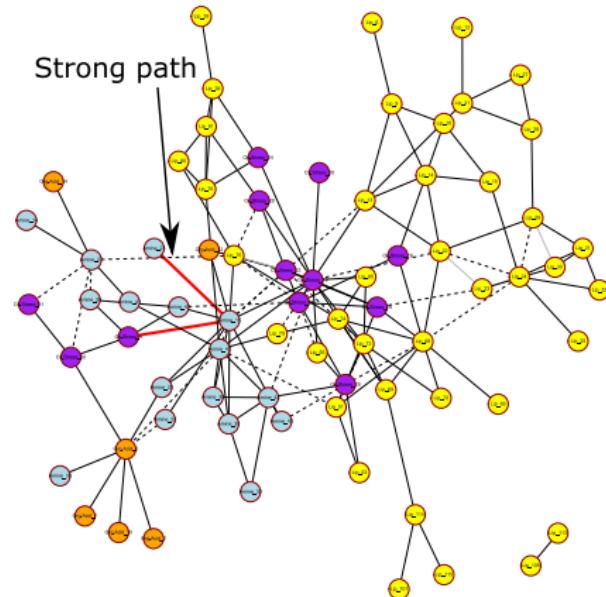
## Analysis: Global Level



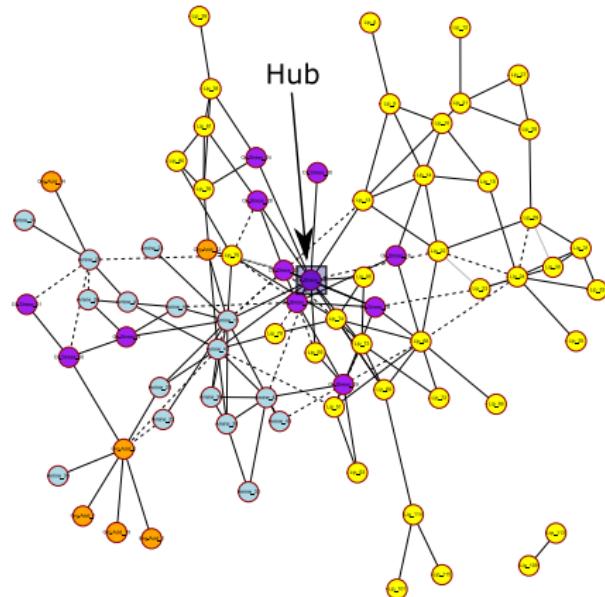
## Analysis: Group Level



## Analysis: Path Level



## Analysis: Node Level



## Node-Level Analysis: Centrality

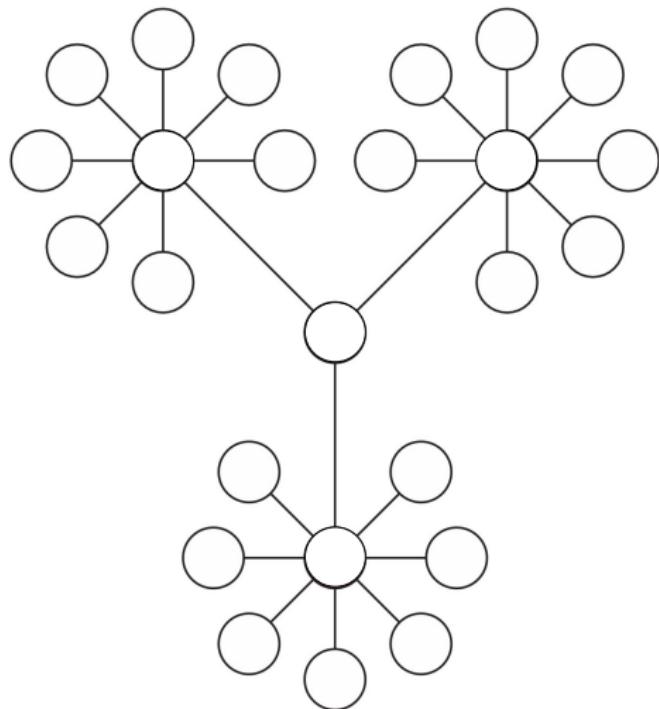


Illustration adapted from: <http://shareablespaces.blogspot.nl/2010/03/social-networking-tools.html>

## Node-Level Analysis: Degree Centrality

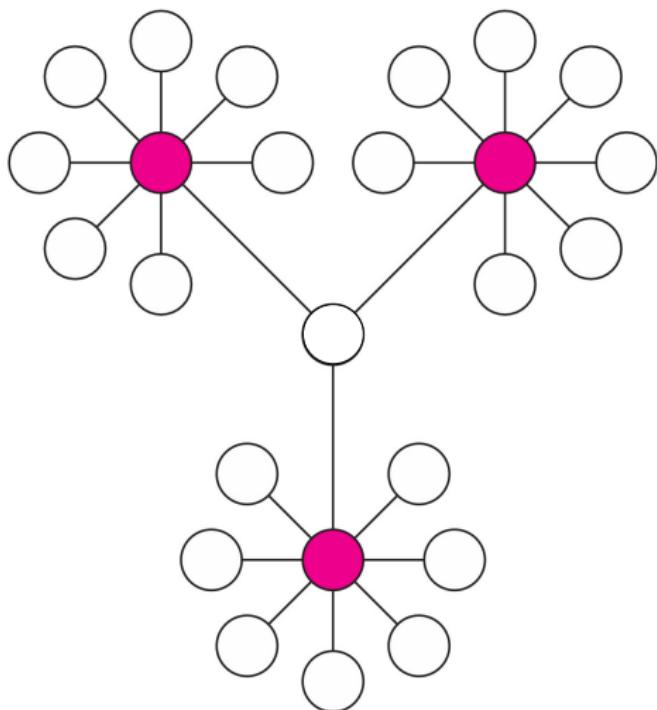


Illustration adapted from: <http://shareablespace.blogspot.nl/2010/03/social-networking-tools.html>

## Node-Level Analysis: Betweenness Centrality

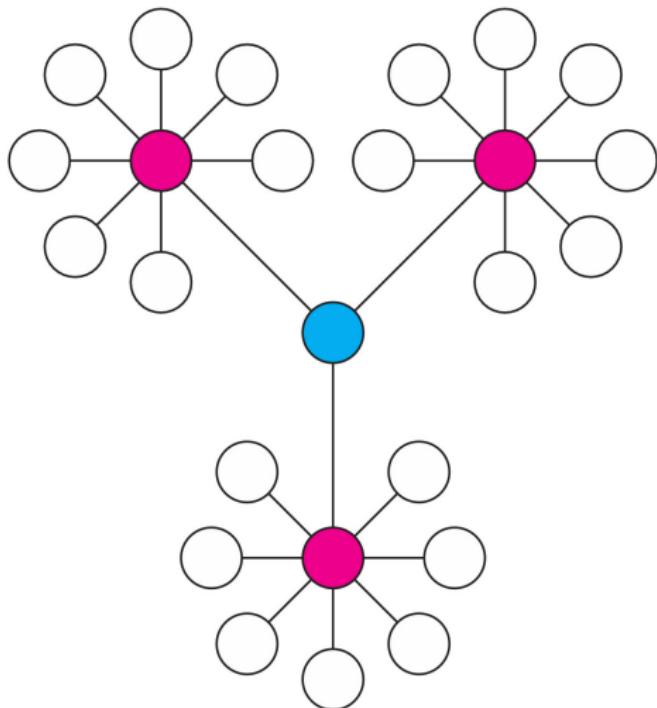


Illustration adapted from: <http://shareablespace.blogspot.nl/2010/03/social-networking-tools.html>

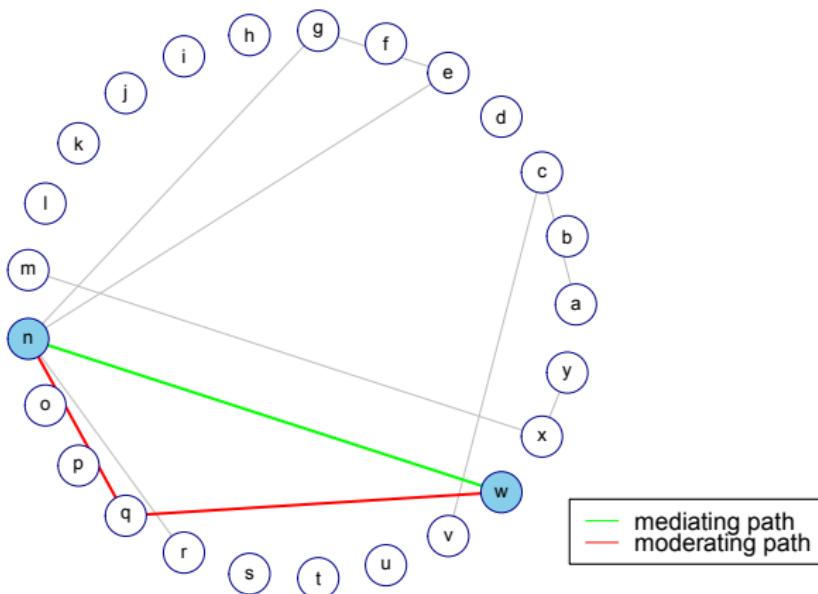
## Exercise 6: Find the nodes with the top degree and betweenness centralities

```
GGMnetworkStats(sparseP)  
↑  
sparse matrix
```

Returns list object

- \$degree: Degree centrality
- \$betweenness: Betweenness centrality
- ...

## Path-Level Analysis: Mediating and Moderating Paths



## Exercise 7: Find 2 strongest paths between Amines 1 and 2

```
GGMpathStats(P0,      ← sparse precision matrix  
            node1, ← endpoint 1  
            node2, ← endpoint 2  
            graph, ← logical, should graph be produced?  
            ... ) ← arguments passed to Ugraph
```

Returns list object

- \$pathStats: Matrix specifying paths
- ...

## Group-Level Analysis: Finding Communities

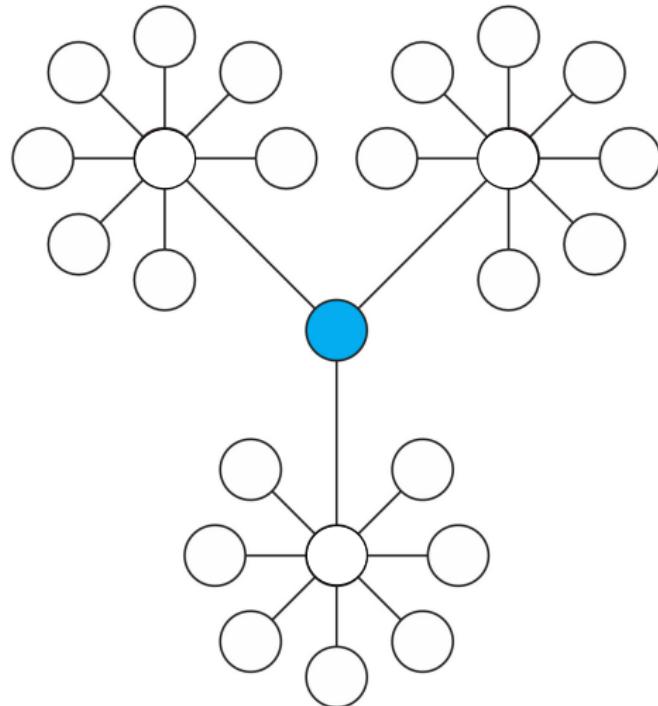


Illustration adapted from: <http://shareablespace.blogspot.nl/2010/03/social-networking-tools.html>

## Group-Level Analysis: Finding Communities

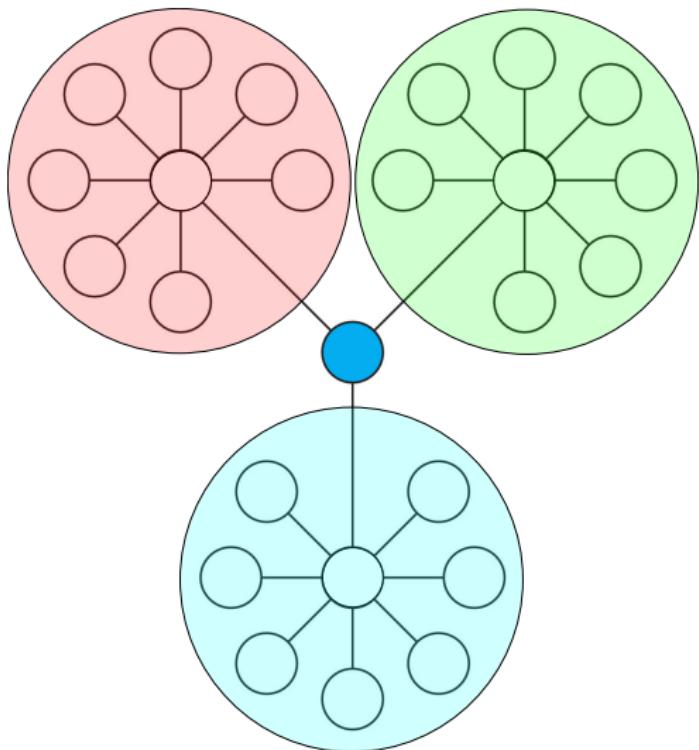


Illustration adapted from: <http://shareablesplaces.blogspot.nl/2010/03/social-networking-tools.html>

## Exercise 8: Find and visualize communities for the extracted network

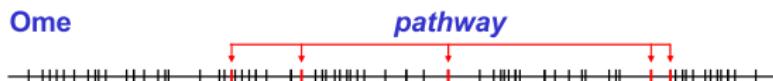
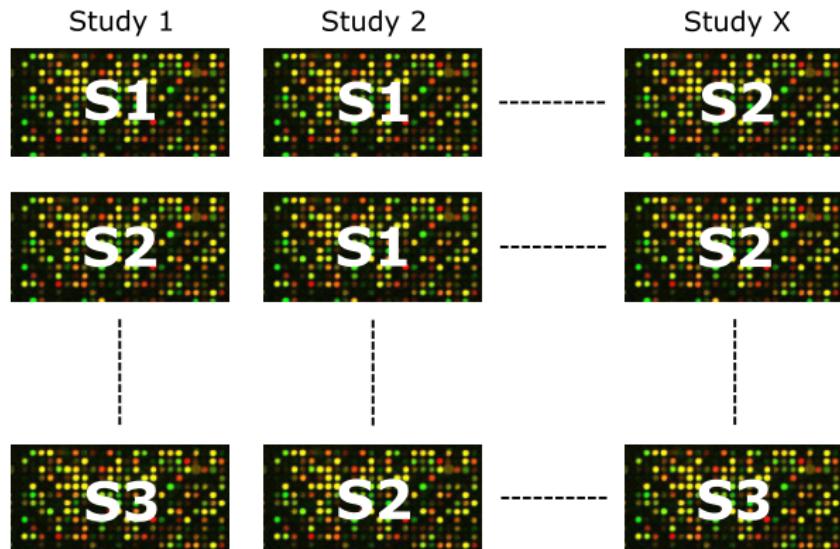
```
Communities(P,      ← sparse matrix  
graph,    ← logical, if TRUE, then graph is given  
lay,  
Vsize,  
Vcex,  
Vcolor,  
VBcolor,  
VLcolor,  
main)
```

arguments Ugraph

Returns list object

- \$membership: Community membership for each feature
- \$modularityscore: Modularity score

## Multiple data classes



# Situation

## Data

- G classes of  $(n_g \times p)$ -dimensional data
- Classes defined by data sets and/or (subtypes of) diseases

## Assumption

Precision matrices of constituent classes chiefly share the same structure but potentially differ in a number of locations of interest

## Desire

Integrative or meta-analytic Gaussian graphical modeling

## Targeted fused ridge estimation: General Formulation

Maximize

$$\underbrace{\mathcal{L}(\{\Omega_g\}; \{S_g\})}_{\text{log-likelihood}} - \sum_g \underbrace{\frac{\lambda_{gg}}{2} \|\Omega_g - T_g\|_F^2}_{\text{ridge-penalty}} - \sum_{g1,g2} \underbrace{\frac{\lambda_{g1g2}}{4} \|(\Omega_{g1} - T_{g1}) - (\Omega_{g2} - T_{g2})\|_F^2}_{\text{fusion-penalty}}$$

- $T_g$  indicate class-specific target matrices
- $\lambda_{gg} \in (0, \infty)$  denote class-specific ridge penalty parameters
- $\lambda_{g1g2} \in [0, \infty)$  denote pair-specific fusion penalty parameters,  $\lambda_{g1g2} = \lambda_{g2g1}$

Penalty matrix

All penalties can be collected into a non-negative symmetric matrix  $\Lambda = [\lambda_{g1g2}]$

# Targeted fused ridge estimation

Maximizing argument for class  $g_0$

$$\hat{\Omega}_{g_0}(\Lambda, \{\Omega_g\}_{g \neq g_0}) = \left\{ \left[ \bar{\lambda}_{g_0} \mathbf{I}_p + \frac{1}{4} (\bar{\mathbf{S}}_{g_0} - \bar{\lambda}_{g_0} \mathbf{T}_{g_0})^2 \right]^{1/2} + \frac{1}{2} (\bar{\mathbf{S}}_{g_0} - \bar{\lambda}_{g_0} \mathbf{T}_{g_0}) \right\}^{-1},$$

where

$$\bar{\mathbf{S}}_{g_0} = \mathbf{S}_{g_0} - \sum_{g \neq g_0} \frac{\lambda_{gg_0}}{n_{g_0}} (\Omega_g - \mathbf{T}_g), \quad \text{and} \quad \bar{\lambda}_{g_0} = \frac{\sum_g \lambda_{gg_0}}{n_{g_0}}$$

# Properties

## Behavior

- i.  $\hat{\Omega}_g \succ \mathbf{0}$  for all  $\lambda_{gg} \in (0, \infty)$ ;
- ii.  $\lim_{\lambda_{gg} \rightarrow 0^+} \hat{\Omega}_g = \mathbf{S}_g^{-1}$  if  $\sum_{g' \neq g} \lambda_{gg'} = 0$  and  $p \leq n_g$ ;
- iii.  $\lim_{\lambda_{gg} \rightarrow \infty} \hat{\Omega}_g = \mathbf{T}_g$  if  $\lambda_{gg'} < \infty$  for all  $g' \neq g$ ;
- iv.  $\lim_{\lambda_{g_1 g_2} \rightarrow \infty} (\hat{\Omega}_{g_1} - \mathbf{T}_{g_1}) = \lim_{\lambda_{g_1 g_2} \rightarrow \infty} (\hat{\Omega}_{g_2} - \mathbf{T}_{g_2})$  if  $\lambda_{g'_1 g'_2} < \infty$  for all  $\{g'_1, g'_2\} \neq \{g_1, g_2\}$ .

## Block coordinate ascent

```

1: Input:
2: Sufficient data:  $(\mathbf{S}_1, n_1), \dots, (\mathbf{S}_G, n_G)$ 
3: Penalty matrix:  $\Lambda$ 
4: Convergence criterion:  $\varepsilon > 0$ 
5: Output:
6: Estimates:  $\hat{\Omega}_1, \dots, \hat{\Omega}_G$ 
7: procedure RIDGE.P.FUSED( $\mathbf{S}_1, \dots, \mathbf{S}_G, n_1, \dots, n_G, \Lambda, \varepsilon$ )
8:   Initialize:  $\hat{\Omega}_g^{(0)}$  for all  $g$ .
9:   for  $c = 1, 2, 3, \dots$  do
10:    for  $g = 1, 2, \dots, G$  do
11:      Update  $\hat{\Omega}_g^{(c)} := \hat{\Omega}_g(\Lambda, \hat{\Omega}_1^{(c)}, \dots, \hat{\Omega}_{g-1}^{(c)}, \hat{\Omega}_{g+1}^{(c-1)}, \dots, \hat{\Omega}_G^{(c-1)})$ 
12:    end for
13:    if  $\max_g \left\{ \frac{\|\hat{\Omega}_g^{(c)} - \hat{\Omega}_g^{(c-1)}\|_F^2}{\|\hat{\Omega}_g^{(c)}\|_F^2} \right\} < \varepsilon$  then
14:      return  $(\hat{\Omega}_1^{(c)}, \dots, \hat{\Omega}_G^{(c)})$ 
15:    end if
16:  end for
17: end procedure

```

## Functional analogues

Core	Fused
ridgeP	ridgeP.fused
optPenalty	optPenalty.fused
sparsify	sparsify.fused
GGMnetworkStats	GGMnetworkStats.fused
GGMpathStats	GGMpathStats.fused

## Exercise 9: Construct lists of class-specific target and data matrices

```
## Subset
ADclass1 <- ADmetabolites[, sampleInfo$ApoEClass == "Class 1"]
ADclass2 <- ADmetabolites[, sampleInfo$ApoEClass == "Class 2"]

## Transpose and scale data
ADclass1 <- scale(t(ADclass1))
ADclass2 <- scale(t(ADclass2))

## Correlations for subsets
rAD1 <- cor(ADclass1)
rAD2 <- cor(ADclass2)

## Constructing list of correlation matrices
Rlist = list(rAD1 = rAD1, rAD2 = rAD2)
samps = c(dim(ADclass1)[1], dim(ADclass2)[1])

## Constructing list of target matrices and data
Tlist <- default.target.fused(Slist = Rlist, ns = samps, type = "DUPV")
Ylist <- list(AD1data = ADclass1, AD2data = ADclass2)
```

## Exercise 10: Find optimal precision matrices for Class 1 and Class 2 AD data

```
optPenalty.fused(Ylist,      ← list of data matrices  
                    Tlist,      ← list of target matrices  
                    cv.method) ← CV method: choose LOOCV
```

Returns list object

- \$lambda.unique: Optimal penalty parameters
- \$Plist: List of precision matrices under optimal penalty parameters
- ...

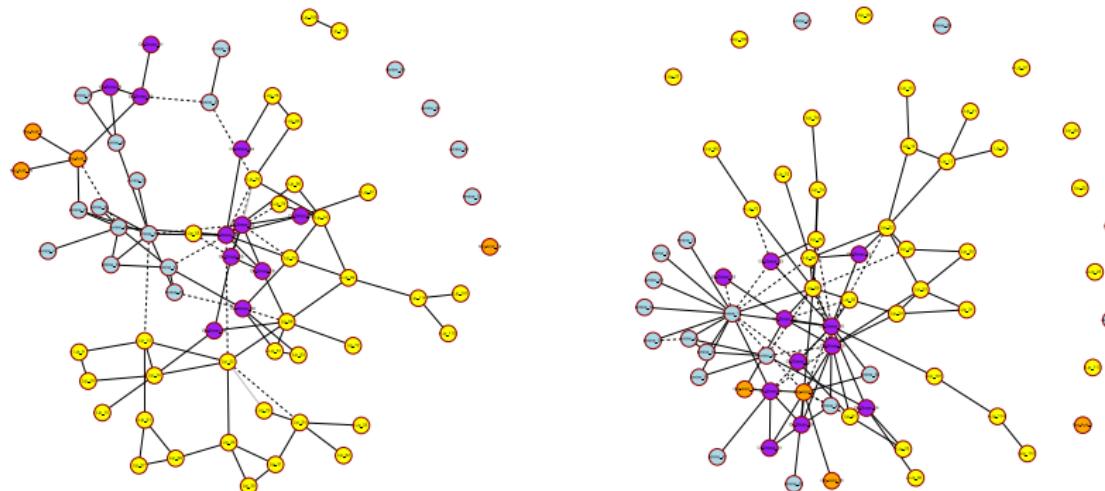
## Exercise 11: Extract class-specific networks on the basis of local FDR thresholding

```
sparsify.fused(Plist,      ← list of estimated precision matrices
                threshold, ← type: "localFDR", "top"
                FDRcut)    ← cut-off for 1 - IFDR
```

Returns list object for each class

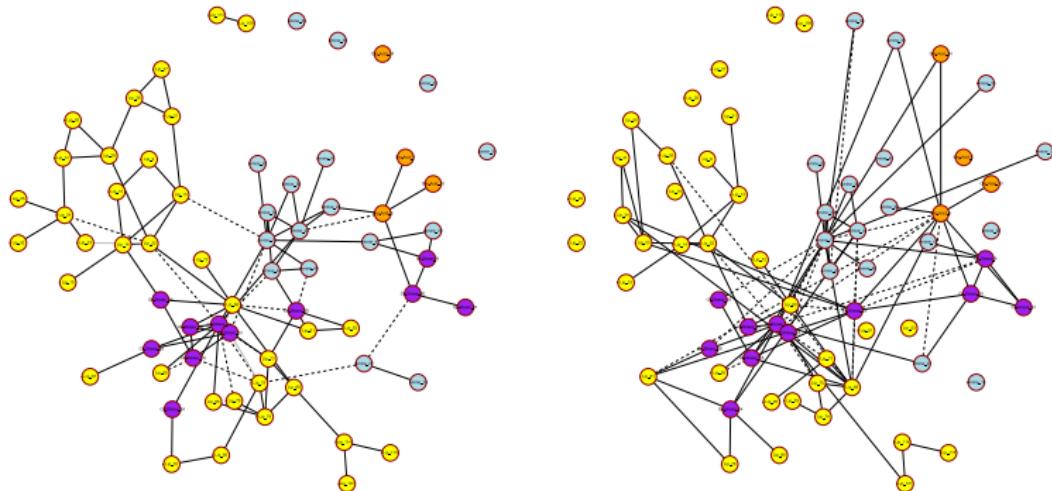
- \$sparsePrecision: Sparsified precision matrix
- \$sparseParCor: Sparsified partial correlation matrix

## Visualization



Two class-specific networks, independently visualized with the FR algorithm

# Visualization



Coordinate refinement supports visual comparison

Same two class-specific networks, visualized with the same node-coordinates

## Exercise 12: Visualize the retained networks in the same coordinates

```
Ugraph(M,  
       type,  
       lay,  
       coords, ← if lay = NULL, then  
       Vsize,      layout according to coordinates  
       Vcex,  
       Vcolor,  
       VBcolor,  
       VLcolor,  
       prune  
       ...)
```

Returns matrix object

- Containing coordinates of layout

### Tips

- Use the union function.
- Use coordinates return object.

## Exercise 13: Compare the top node-degrees for the class-specific networks

```
GGMnetworkStats.fused(Plist)
```

A list of sparse matrices



Returns `data.frame`

Names of `Plist` are prefixed to column-names

## Exercise 14: Find and visualize communities for the class-specific networks

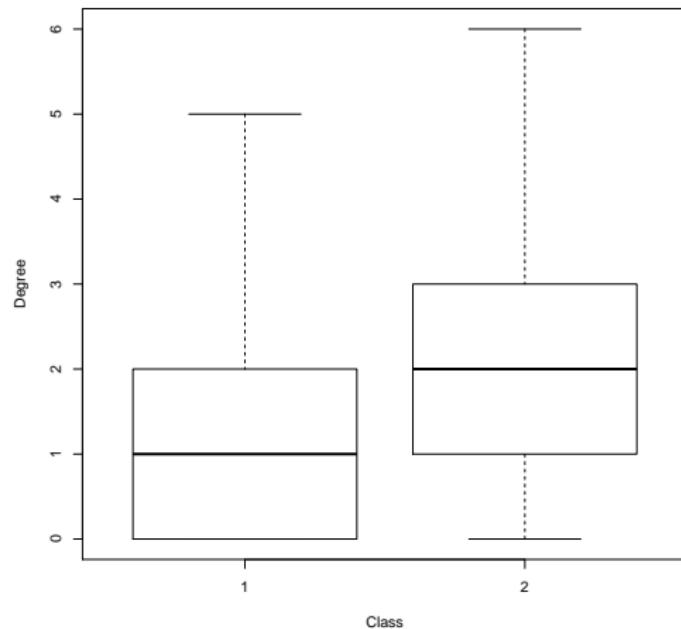
```
Communities(P,      ← sparse matrix  
graph,    ← logical, if TRUE, then graph is given  
lay,  
Vsize,  
Vcex,  
Vcolor,  
VBcolor,  
VLcolor,  
main)
```

arguments Ugraph

Returns list object

- \$membership: Community membership for each feature
- \$modularityscore: Modularity score

# Global Analysis: Degree Distributions



## Global Analysis: Entropy

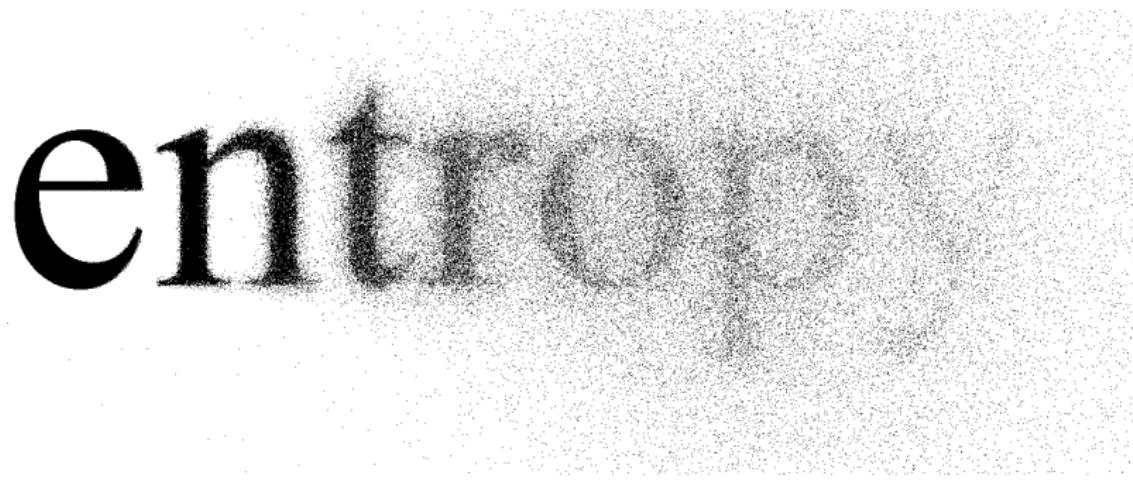


Illustration adapted from: <http://creepypasta.wikia.com/wiki/Entropy>

## Exercise 15: Compare either the degree distributions or the entropies for the class-specific networks

### Comparing degree distributions

Can be done with simple test, such as the Wilcoxon Signed Rank Test

### Comparing entropies

```
library(sigar)

entropyTest(Y,      ← data matrix, samples in rows
            id,      ← class indicator (coded 0,1)
            nPerm,   ← number of permutations
            method) ← distributional assumption: use "knn"
```

### Entropy test returns test information

```
summary()
```

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## Manual/Download

- Peeters, C.F.W., Bilgrau, A.E., & van Wieringen, W.N. (2017). "rags2ridges: Ridge Estimation of Precision Matrices from High-Dimensional Data". R package, version 2.2. URL: <https://cran.r-project.org/package=rags2ridges>.

## Theory/Methodology

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## Software

- Peeters, C.F.W., van de Wiel, M.A., & van Wieringen, W.N. (2016) "The Spectral Condition Number Plot for Regularization Parameter Determination". [arXiv:1608.04123v1 \[stat.CO\]](https://arxiv.org/abs/1608.04123v1).
- van Wieringen, W.N. & Peeters, C.F.W. (2015). "Application of a New Ridge Estimator of the Inverse Covariance Matrix to the Reconstruction of Gene-Gene Interaction Networks". In: di Serio, C., Lio, P., Nonis, A., and Tagliaferri, R. (Eds.) 'Computational Intelligence Methods for Bioinformatics and Biostatistics'. Lecture Notes in Computer Science, vol. 8623. Springer, pp. 170–179.