

Ordering Objects

What Heuristic Should We Use?

Michael Hahsler

Dept. of Engineering Management, Information, and Systems
SMU

November 4, 2015
INFORMS Annual Meeting 2015



- 1 Introduction and Motivation
- 2 Seriation
- 3 Loss Functions
- 4 Optimization Techniques
- 5 Experimental Comparison

Introduction and Motivation

Introduction

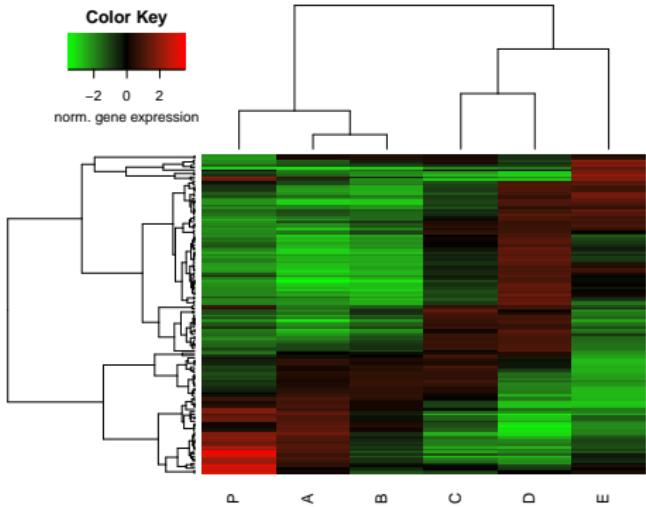
- Part of **Combinatorial Data Analysis** (P. Arabie and Hubert 1996)
- Ordering objects to reveal structural information
- Related to ranking

Some Applications

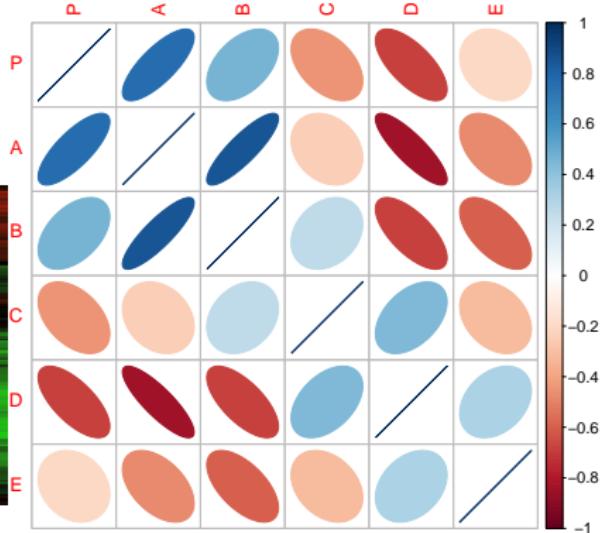
- **Sociology:** Find group structure in sociograms
- **Psychology:** Order subject-by-item response matrix
- **Ecology:** Analyze plant associations
- **Manufacturing:** Product flow analysis
- **Biology:**
 - ▶ Arrange gene expression data (heat maps)
 - ▶ Gene sequencing, read assembly using the consecutive ones problem (C1P)
- **Visualization:**
 - ▶ Reorder large data tables
 - ▶ Cluster tendency (visual assessment of cluster tendency, VAT)
 - ▶ Evaluation of clustering quality (dissimilarity plot)

Examples

Gene Expression Data

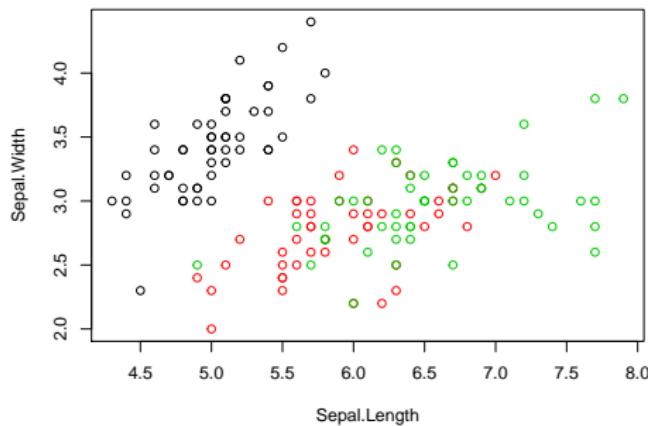


Reordered Correlation Plot

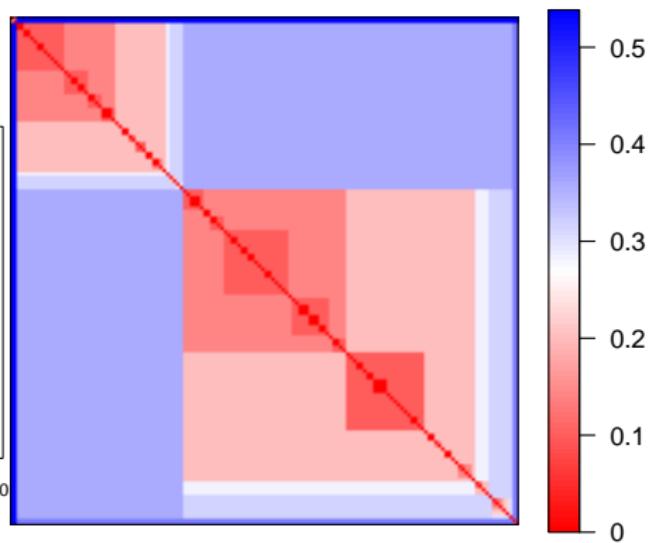


Examples

Iris Data Set



Clustering Tendency (iVAT)



Motivation

Many methods to order objects have been proposed.

How should we do it?

Seriation

Problem Definition

- Arrange a set of n objects

$$\mathcal{O} = \{O_1, O_2, \dots, O_n\}$$

in a **linear order** given available data and some loss function in order to reveal structural information.

- **Data:** A symmetric dissimilarity matrix $\mathbf{D} = [d_{ij}]_{n \times n}$, where d_{ij} represents the dissimilarity between O_i and O_j , and $d_{ii} = 0$ for all i .
- **Linear order:** A permutation function $\Psi(\mathbf{D}) = P_\pi \mathbf{D} P_\pi^T$, where π is the permutation vector and P_π is the permutation matrix for Ψ .

Problem Definition

- Arrange a set of n objects

$$\mathcal{O} = \{O_1, O_2, \dots, O_n\}$$

in a **linear order** given available data and some loss function in order to reveal structural information.

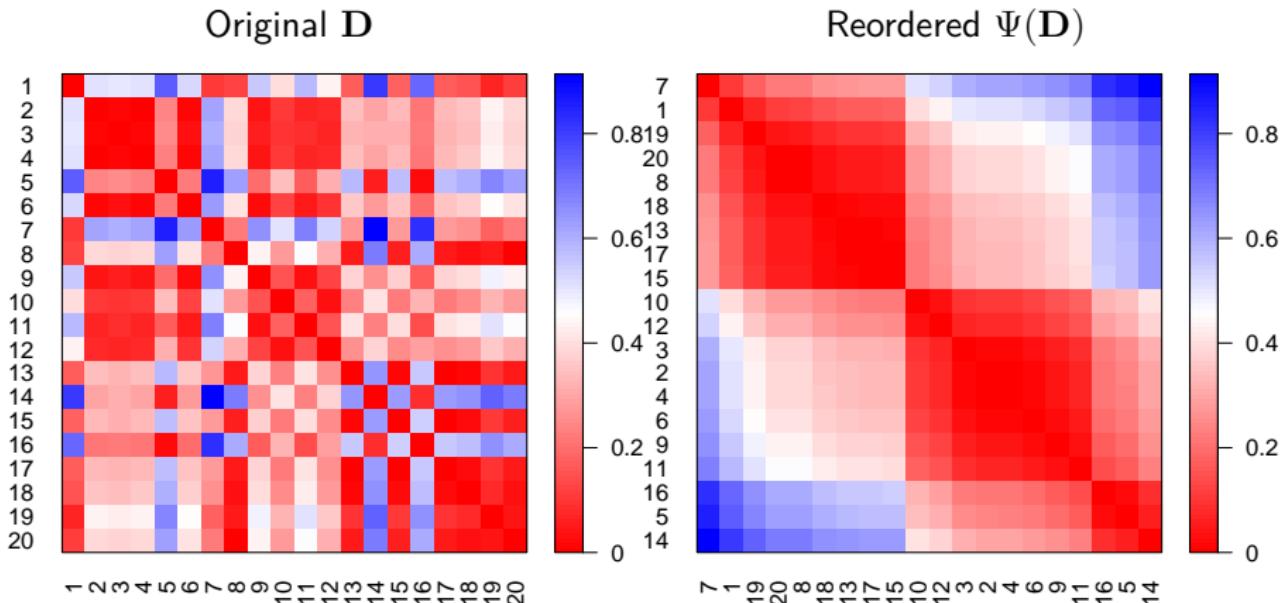
- **Data:** A symmetric dissimilarity matrix $\mathbf{D} = [d_{ij}]_{n \times n}$, where d_{ij} represents the dissimilarity between O_i and O_j , and $d_{ii} = 0$ for all i .
- **Linear order:** A permutation function $\Psi(\mathbf{D}) = P_\pi \mathbf{D} P_\pi^T$, where π is the permutation vector and P_π is the permutation matrix for Ψ .

Optimization problem

$$\Psi^* = \underset{\Psi \in \mathcal{P}_n}{\operatorname{argmin}} L(\Psi(\mathbf{D})),$$

where L is a loss function to evaluate how well a given permutation reveals structural information. \mathcal{P}_n is the set of all possible permutation functions.

Example of Seriation



Typical assumption: Structural information is revealed if more similar objects are presented closer together.

Issues and Techniques

Issues

- How do we define the loss function L ?
- Requires to solve a discrete optimization problem with solution space size of $O(n!)$.

Issues and Techniques

Issues

- How do we define the loss function L ?
- Requires to solve a discrete optimization problem with solution space size of $O(n!)$.

Techniques

- Partial enumeration methods (currently solve problems with $n \leq 40$):
 - ▶ dynamic programming
 - ▶ branch-and-bound
- Other discrete optimization methods
 - ▶ QAP formulation
 - ▶ spectral methods
- Heuristics for larger problems

Loss Functions

Column/Row Gradient Measures

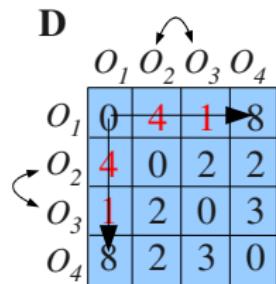
Perfect anti-Robinson matrix (Robinson 1951): A symmetric matrix where the values in all rows and columns only increase when moving away from the main diagonal. Gradient conditions (L. Hubert, Arabie, and Meulman 1987):

within rows: $d_{ik} \leq d_{ij}$ for $1 \leq i < k < j \leq n$;

within columns: $d_{kj} \leq d_{ij}$ for $1 \leq i < k < j \leq n$.

D

	O_1	O_2	O_3	O_4
O_1	0	4	1	8
O_2	4	0	2	2
O_3	1	2	0	3
O_4	8	2	3	0



$\Psi(D)$

	O_1	O_3	O_2	O_4
O_1	0	1	4	8
O_3	1	0	2	3
O_2	4	2	0	2
O_4	8	3	2	0

The closer objects are together in the order of the matrix, the higher their similarity.

Note: Most matrices can only be brought into a near anti-Robinson form.

Column/Row Gradient Measures (cont.)

Gradient measure (quantifies the divergence from anti-Robinson form)

$$L(\mathbf{D}) = \sum_{i < k < j} f(d_{ik}, d_{ij}) + \sum_{i < k < j} f(d_{kj}, d_{ij})$$

where $f(\cdot, \cdot)$ is a function which defines how a violation or satisfaction of a gradient condition for an object triple $(O_i, O_k \text{ and } O_j)$ is counted.

Column/Row Gradient Measures (cont.)

Gradient measure (quantifies the divergence from anti-Robinson form)

$$L(\mathbf{D}) = \sum_{i < k < j} f(d_{ik}, d_{ij}) + \sum_{i < k < j} f(d_{kj}, d_{ij})$$

where $f(\cdot, \cdot)$ is a function which defines how a violation or satisfaction of a gradient condition for an object triple $(O_i, O_k \text{ and } O_j)$ is counted.

Raw number of violations minus satisfactions:

$$f(z, y) = \text{sign}(y - z) = \begin{cases} -1 & \text{if } z > y; \\ 0 & \text{if } z = y; \\ +1 & \text{if } z < y. \end{cases}$$

Weight each satisfaction or violation by its magnitude (absolute difference between the values):

$$f(z, y) = |y - z| \text{ sign}(y - z) = y - z$$

Anti-Robinson Events/Deviation

An even simpler loss function can be created in the same way as the gradient measures above by concentrating on violations only.

AR Events

$$L(\mathbf{D}) = \sum_{i < k < j} f(d_{ik}, d_{ij}) + \sum_{i < k < j} f(d_{kj}, d_{ij})$$

To only count the violations (called events) we use

$$f(z, y) = I(z, y) = \begin{cases} 1 & \text{if } z < y \quad \text{and} \\ 0 & \text{otherwise.} \end{cases}$$

$I(\cdot)$ is an indicator function returning 1 only for violations.

(Chen 2002) also introduced a weighted versions of this loss function by using the absolute deviations as weights:

$$f(z, y) = |y - z| I(z, y)$$

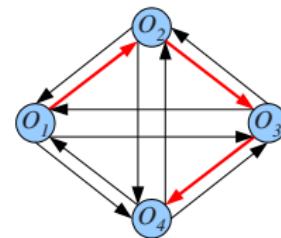
Hamiltonian Path Length

- \mathbf{D} is seen as a complete weighted graph $G = (\Omega, E)$ with $\Omega = \{O_1, O_2, \dots, O_n\}$ and the weight w_{ij} for edge $e_{ij} \in E$ represents d_{ij} .
- An order Ψ can be seen as a *Hamiltonian path* through the graph.
- **Minimizing the path length** results in a seriation optimal with respect to dissimilarities between neighboring objects (Hubert 1974, Caraux and Pinloche (2005)).

Loss function:

$$L(\mathbf{D}) = \sum_{i=1}^{n-1} d_{i,i+1}$$

\mathbf{D}	O_1	O_2	O_3	O_4
O_1	0	4	1	8
O_2	4	0	2	2
O_3	1	2	0	3
O_4	8	2	3	0



This optimization problem is related to the *traveling salesperson problem* (Gutin and Punnen 2002) for which good solvers and efficient heuristics exist.

Other Measures

- Inertia criterion (Caraux and Pinloche 2005)
- Least squares criterion (Caraux and Pinloche 2005)
- Measure of Effectiveness (McCormick, Schweitzer, and White 1972)
- Moore and Neumann stress (Niermann 2005)
- Relative generalized Anti-Robinson events (RGAR) (Tien et al. 2008)

Optimization Techniques

Partial Enumeration

Directly optimize the column/row gradient measures using:

- Dynamic programming (L. Hubert, Arabie, and Meulman 1987)
- Branch-and-bound (Michael Brusco and Stahl 2005)

solves problems with $n \leq 40$

Heuristics for larger problems:

- Simulated annealing (M. Brusco, Köhn, and Stahl 2007)

QAP Formulations

Quadratic Assignment problem

The QAP is in general NP-hard. Methods include QIP, linearization, branch and bound and cutting planes as well as heuristics including Tabu search, simulated annealing, genetic algorithms, and ant systems (Burkhard 1998)

$$\text{QAP}(\mathbf{A}, \mathbf{B}) : \min_{\Psi \in \mathcal{P}_n} \sum_{i,j=1}^n \mathbf{A}_{ij} \Psi(\mathbf{B})_{ij}$$

QAP Formulations

Quadratic Assignment problem

The QAP is in general NP-hard. Methods include QIP, linearization, branch and bound and cutting planes as well as heuristics including Tabu search, simulated annealing, genetic algorithms, and ant systems (Burkhard 1998)

$$\text{QAP}(\mathbf{A}, \mathbf{B}) : \min_{\Psi \in \mathcal{P}_n} \sum_{i,j=1}^n \mathbf{A}_{ij} \Psi(\mathbf{B})_{ij}$$

Formulate as **2-Sum Problem** (Barnard, Pothen, and Simon 1993)

$$\min_{\Psi} \sum_{i,j=1}^n \Psi(\mathbf{S})_{ij} (i - j)^2 \rightarrow \text{QAP}([(i - j)^2]_{n \times n}, \mathbf{S})$$

where similarity matrix $\mathbf{S} = \frac{1}{1+\mathbf{D}}$

QAP Formulations

Quadratic Assignment problem

The QAP is in general NP-hard. Methods include QIP, linearization, branch and bound and cutting planes as well as heuristics including Tabu search, simulated annealing, genetic algorithms, and ant systems (Burkhard 1998)

$$\text{QAP}(\mathbf{A}, \mathbf{B}) : \min_{\Psi \in \mathcal{P}_n} \sum_{i,j=1}^n \mathbf{A}_{ij} \Psi(\mathbf{B})_{ij}$$

Formulate as **2-Sum Problem** (Barnard, Pothen, and Simon 1993)

$$\min_{\Psi} \sum_{i,j=1}^n \Psi(\mathbf{S})_{ij} (i - j)^2 \rightarrow \text{QAP}([(i - j)^2]_{n \times n}, \mathbf{S})$$

where similarity matrix $\mathbf{S} = \frac{1}{1+\mathbf{D}}$

Formulate as **Linear Seriation Problem** (L. Hubert and Schultz 1976)

$$\min_{\Psi} \sum_{i,j=1}^n \Psi(\mathbf{D})_{ij} (-|i - j|) \rightarrow \text{QAP}([-|i - j|]_{n \times n}, \mathbf{D})$$

Spectral Seriation

Minimizes the **2-Sum Problem** formulation:

$$\min_{\Psi} \sum_{i,j=1}^n \Psi(\mathbf{S})_{ij}(i-j)^2$$

Rewriting the minimization problem using a permutation vector π , its inverse, rescaling to \mathbf{q} and using a Lagrangian multiplier for the constraint on the permutation yields (Ding and He 2004) the following equivalent optimization problem:

$$\min_{\mathbf{q}} \frac{\mathbf{q}^T L_{\mathbf{S}} \mathbf{q}}{\mathbf{q}^T \mathbf{q}}$$

where $L_{\mathbf{S}}$ is the Laplacian of \mathbf{S} .

The optimal order can be recovered by the order of the Fielder vector (second smallest eigenvector of the Laplacian).

Hamiltonian Path

TSP Solver

- Concorde (fast cutting planes and tour separation)

Heuristic

- TSP Heuristics (Lin–Kernighan, Nearest neighbors, insertion heuristics, etc.)
- Hierarchical clustering with reordering:
 - ▶ Gruvaeus-Wainer heuristic (Gruvaeus and Wainer 1972),
 - ▶ Optimal leaf ordering (Bar-Joseph et al. 2001)

Other Heuristics

- Hierarchical clustering
- Visual Assessment of cluster Tendency (VAT based on a MST) (Bezdek and Hathaway 2002)
- Multidimensional scaling (metric, non-metric, angle)
- Rank-two ellipse seriation (series of correlation matrices) (Chen 2002).
- Sorting Points Into Neighborhoods (SPIN) (D. Tsafrir et al. 2005)

Experimental Comparison

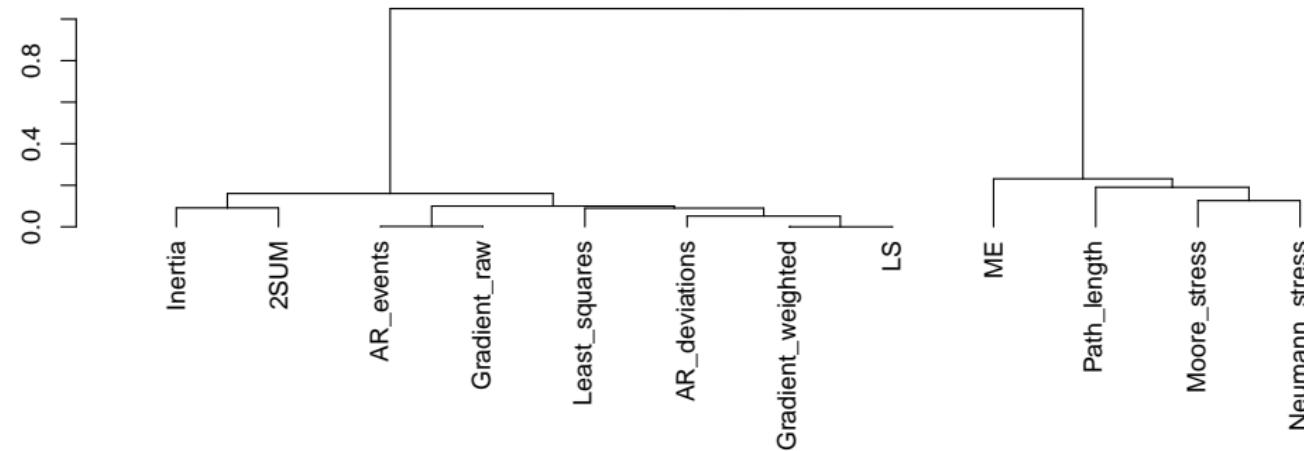
Data Sets

- **Pre-Robinson:** A pre-Anti-Robinson matrix with 100 objects.
- **Random:** 200 objects with 2 features independently and uniformly drawn from $[0, 1]$ (Euclidean dist.)
- **Iris:** 150 flowers with 4 features (scaled, Euclidean dist.)
- **Zoo:** 101 animals with 17 (mostly 0-1) features (Euclidean dist.)
- **Votes:** 1984 Congressional votes for 435 congress men on 16 key votes: yes, no, abstain (Jaccard index on 32 binary features)
- **Wood:** 136 poplar trees with normalized gene expression for 6 locations (Euclidean dist.)
- **Elutriation:** Ratios of gene expression levels for a sample of 100 genes of *Saccharomyces cerevisiae* with 14 eigengenes (from SVD) as features (Euclidean dist.)

Comparison: Loss Functions

Compare how different loss functions rank the results of different seriation methods.

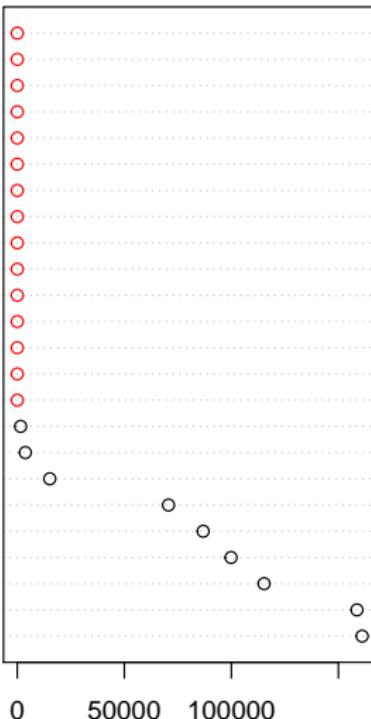
Consensus hierarchy for loss functions



Best least square ultrametric approximation over all methods and data sets.

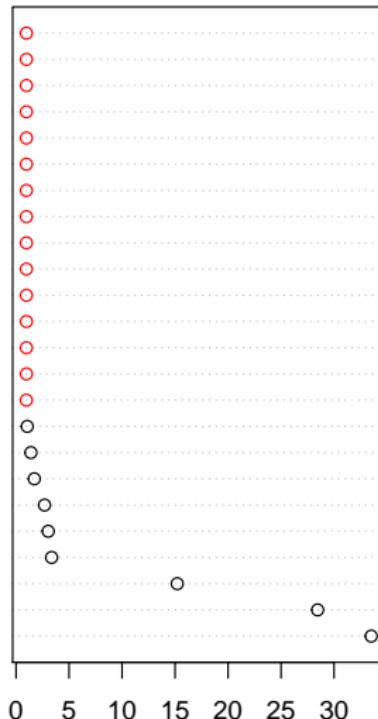
Method Comparison: Pre-Robinson Data

2SUM
LS
SPIN_STS
Spectral
VAT
OLO_complete
OLO_average
OLO_single
GW_complete
GW_average
GW_single
MDS_nonmetric
MDS_metric
R2E
TSP
ARSA
Spectral_norm
SPIN_NH
HC_single
HC_complete
HC_average
MDS_angle
Identity
Random



AR events

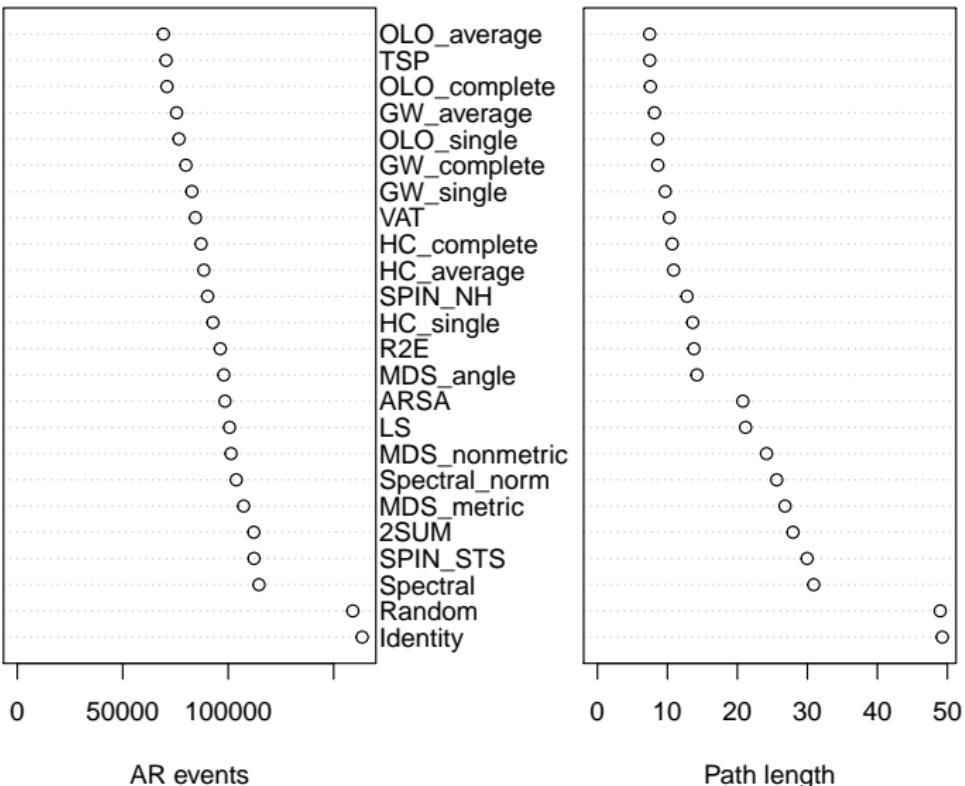
2SUM
LS
SPIN_STS
Spectral
VAT
OLO_complete
OLO_average
OLO_single
GW_complete
GW_average
GW_single
MDS_nonmetric
MDS_metric
R2E
TSP
ARSA
Spectral_norm
SPIN_NH
HC_single
HC_complete
HC_average
MDS_angle
Identity
Random



Path length

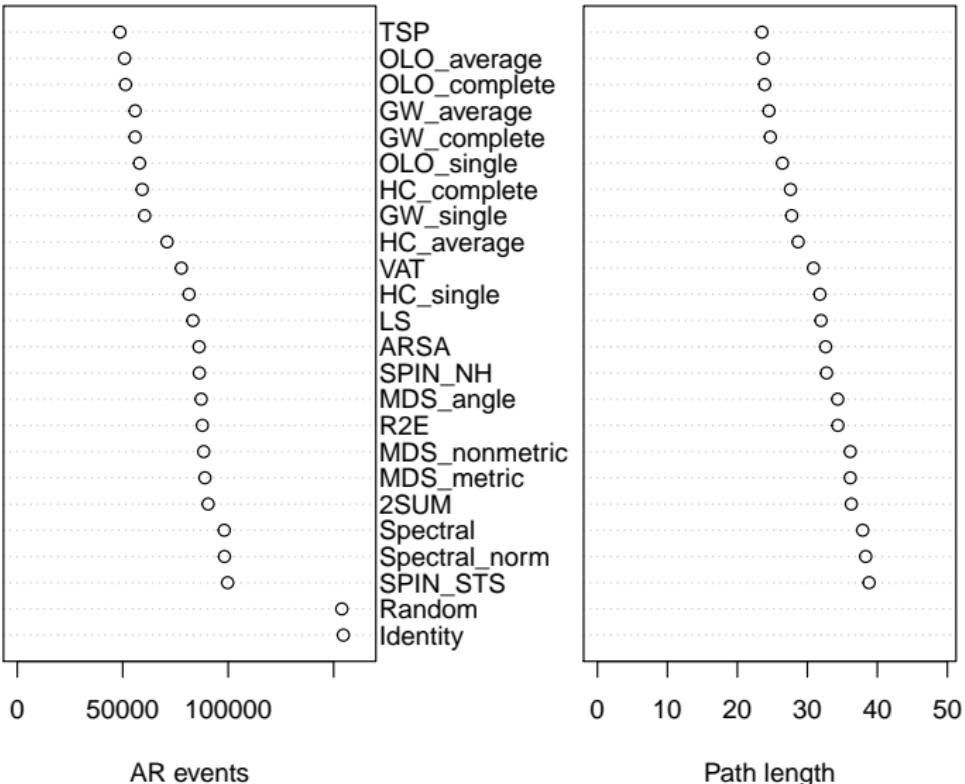
Method Comparison: Random Data

LS
2SUM
ARSA
MDS_nonmetric
Spectral
MDS_metric
SPIN_STS
Spectral_norm
OLO_complete
OLO_single
GW_average
R2E
MDS_angle
SPIN_NH
GW_single
GW_complete
OLO_average
HC_single
HC_average
VAT
HC_complete
TSP
Identity
Random

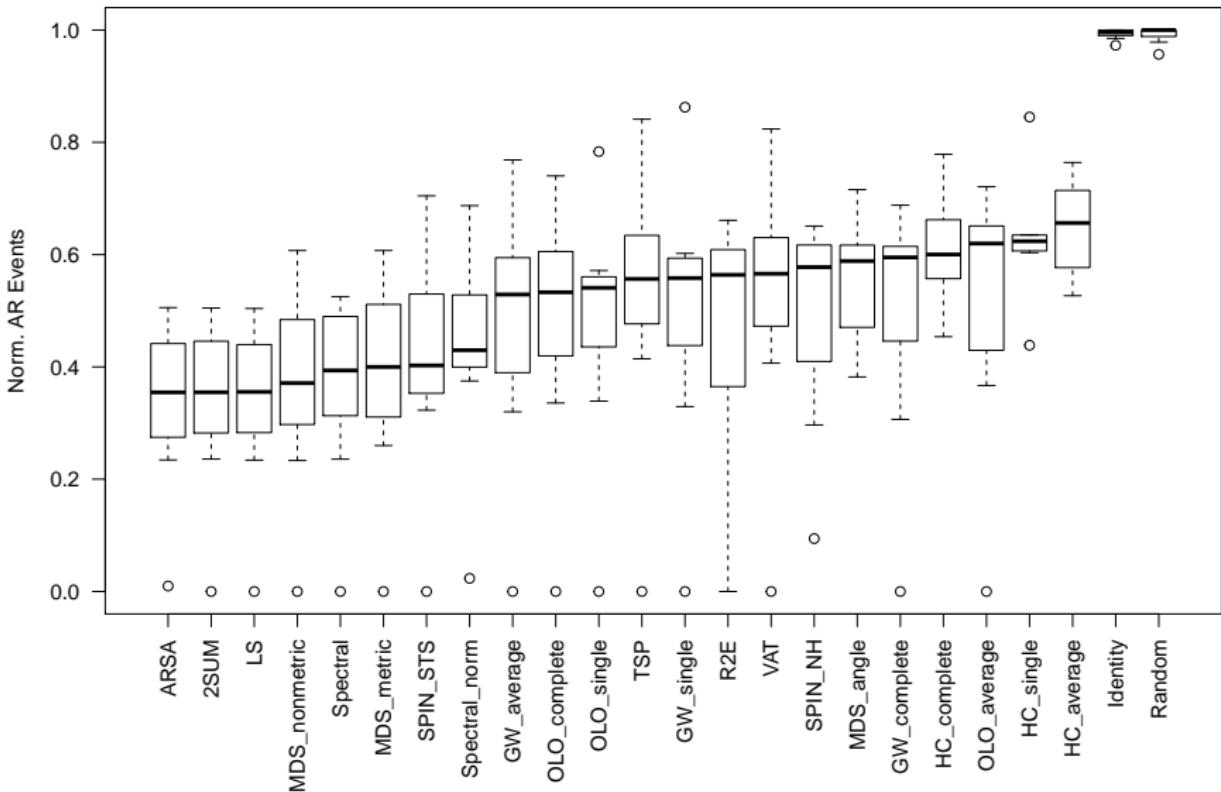


Method Comparison: Votes

ARSA
2SUM
LS
MDS_nonmetric
MDS_metric
Spectral_norm
SPIN_STS
Spectral
GW_average
MDS_angle
HC_average
VAT
OLO_complete
GW_single
R2E
TSP
OLO_single
HC_complete
GW_complete
HC_single
SPIN_NH
OLO_average
Identity
Random

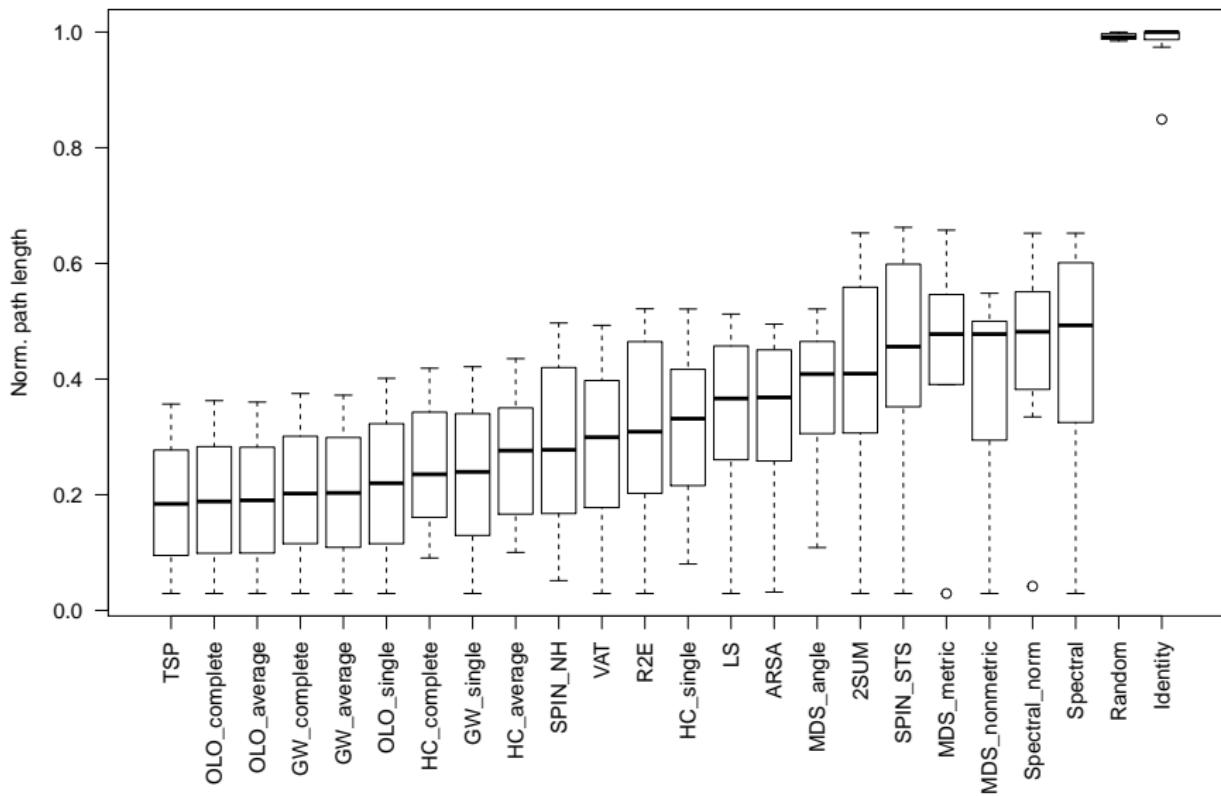


Method Comparison: Anti-Robinson Events



Distribution over all data sets; normalized by largest AR.

Method Comparison: Path length



Distribution over all data sets; normalized by longest path.

Compare Resulting Orders

	Identity	Random	ARSA	TSP	R2E	MDS_metric	MDS_nonmetric	MDS_angle	HC_single	HC_complete
1	47	2	1	36	36	1	1	60	12	1
2	5	56	2	71	71	2	2	47	84	58
3	25	55	3	41	41	3	3	74	82	90
4	69	85	4	31	31	33	33	72	28	62
5	29	26	5	43	43	5	5	88	6	6
6	26	54	6	68	68	6	6	31	5	5
7	97	32	7	64	64	7	7	97	9	68
8	98	59	8	73	73	8	8	4	65	52
9	94	11	9	90	90	9	9	46	96	78
10	65	24	10	60	60	10	10	9	95	77
11	1	43	11	26	26	11	11	2	11	3
12	53	28	12	55	55	12	12	12	55	95
13	32	86	13	33	33	31	31	16	30	31
14	87	5	14	82	82	62	62	15	86	51
15	64	69	15	44	44	15	15	86	70	42
16	10	7	16	42	42	20	20	91	91	82
17	13	79	17	14	14	17	17	70	19	57
18	19	89	18	34	34	18	18	1	39	84
19	22	95	19	76	76	22	22	20	57	20
20	8	46	20	65	65	16	16	84	3	39
21	63	58	21	93	93	21	21	71	27	65
22	21	53	22	23	23	19	19	83	79	83
23	80	63	23	19	19	23	23	43	25	69
24	6	51	24	96	96	24	24	90	24	19
25	72	42	25	5	5	43	43	76	62	30
26	100	35	26	11	11	26	26	37	77	43
27	45	60	27	35	35	27	27	17	36	56
28	85	10	28	59	59	28	28	8	41	34
29	83	82	29	32	32	29	29	28	76	99
30	67	66	30	50	50	30	30	26	53	87

Shown are the object ranks for the pre-Robinson data (first 30 objects sorted by ARSA and first 10 methods).

Compare Orders (cont.)

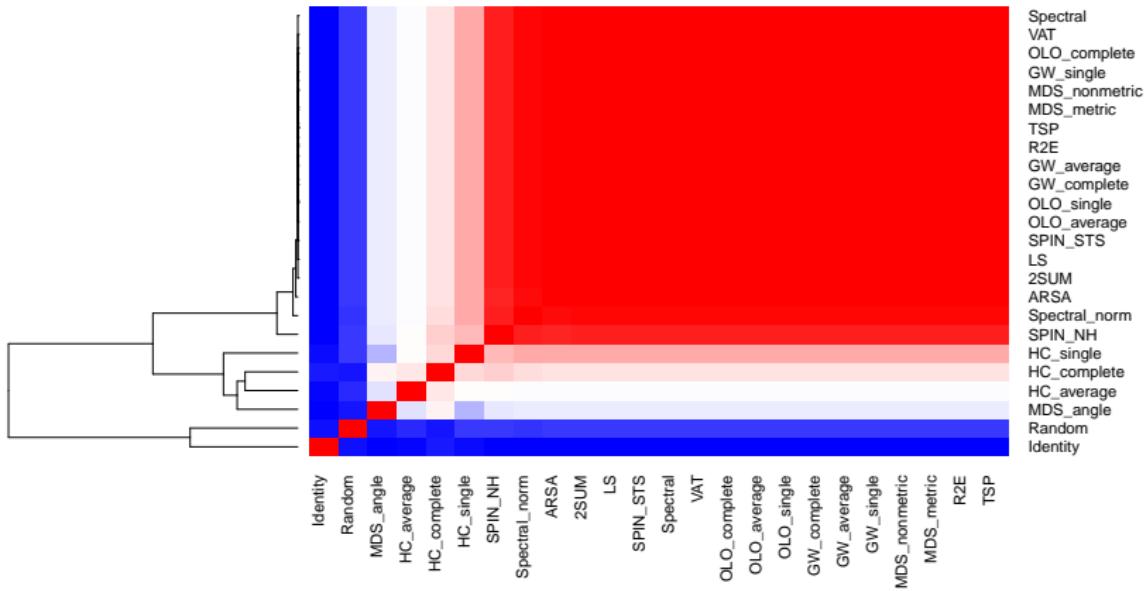
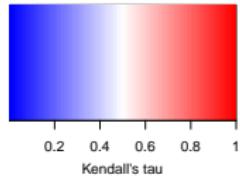
Compare the seriation order produced by two methods using

- **Kendall's rank-order correlation**

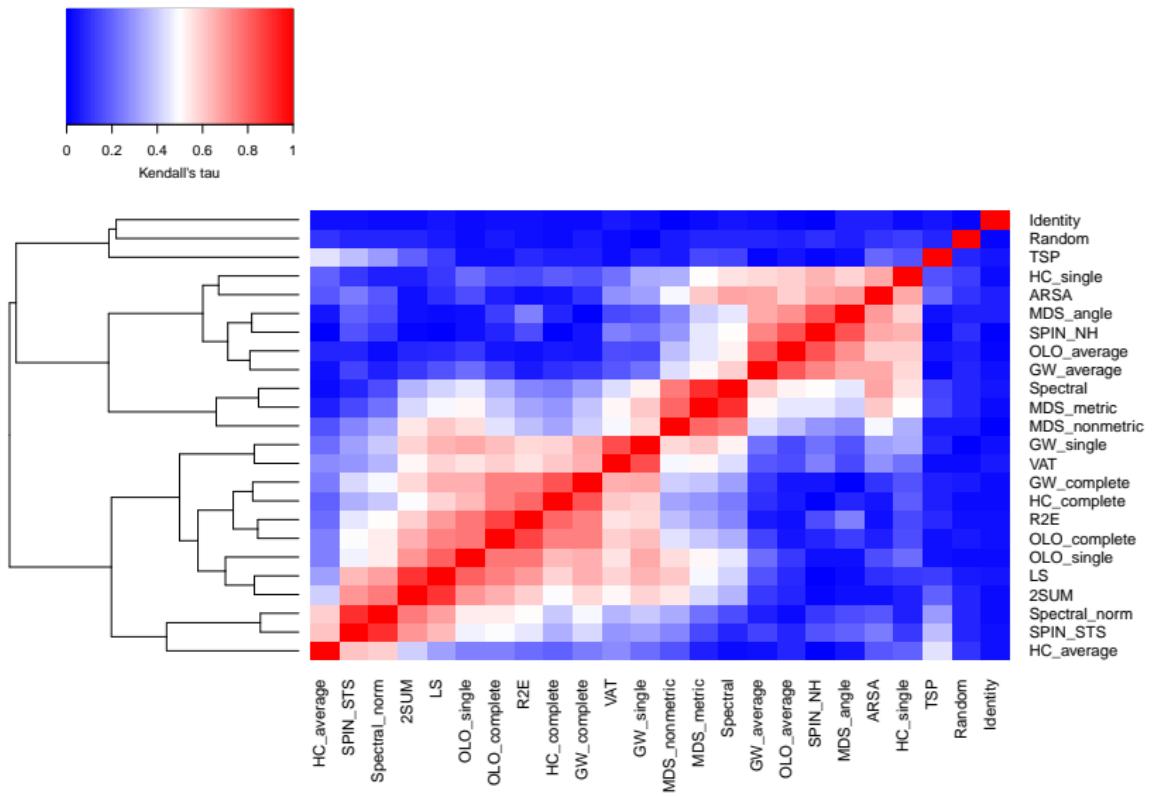
Other options would be

- Spearman's rank-order correlation
- Spearman's footrule metric (i.e., Manhattan distance between ranks) (Diaconis 1988)
- Positional proximity coefficient (Goulermas et al 2015)

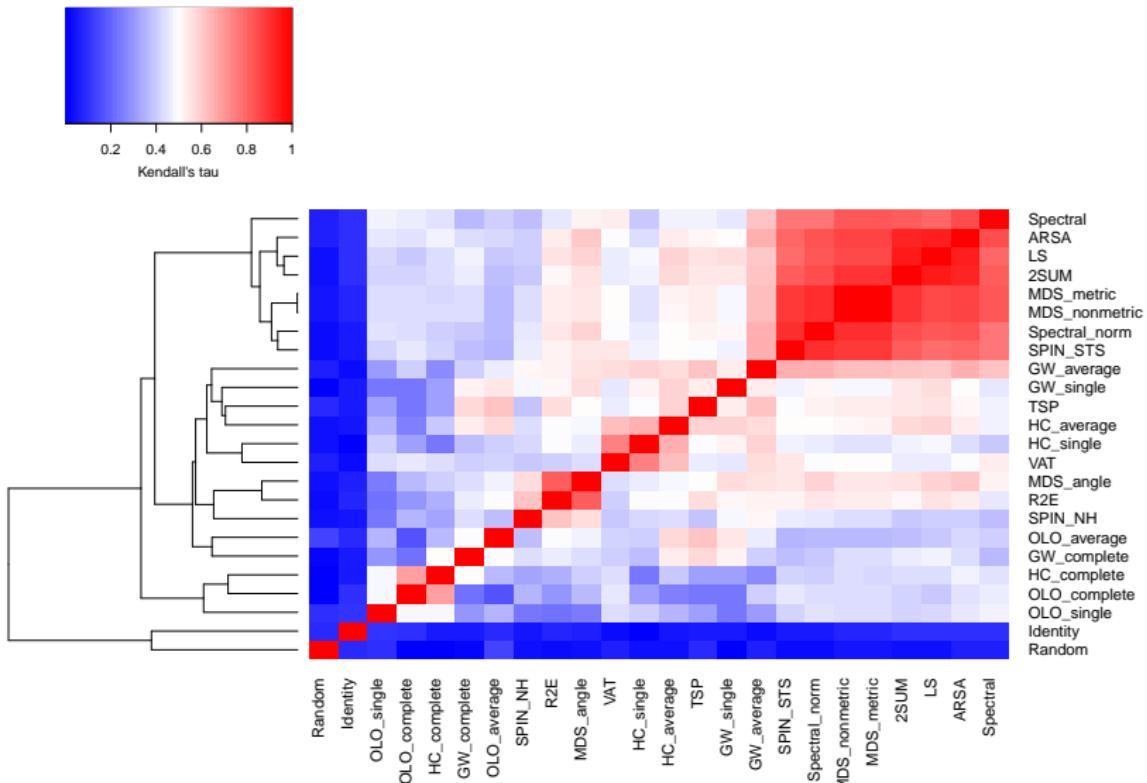
Comparison: Pre-Robinson



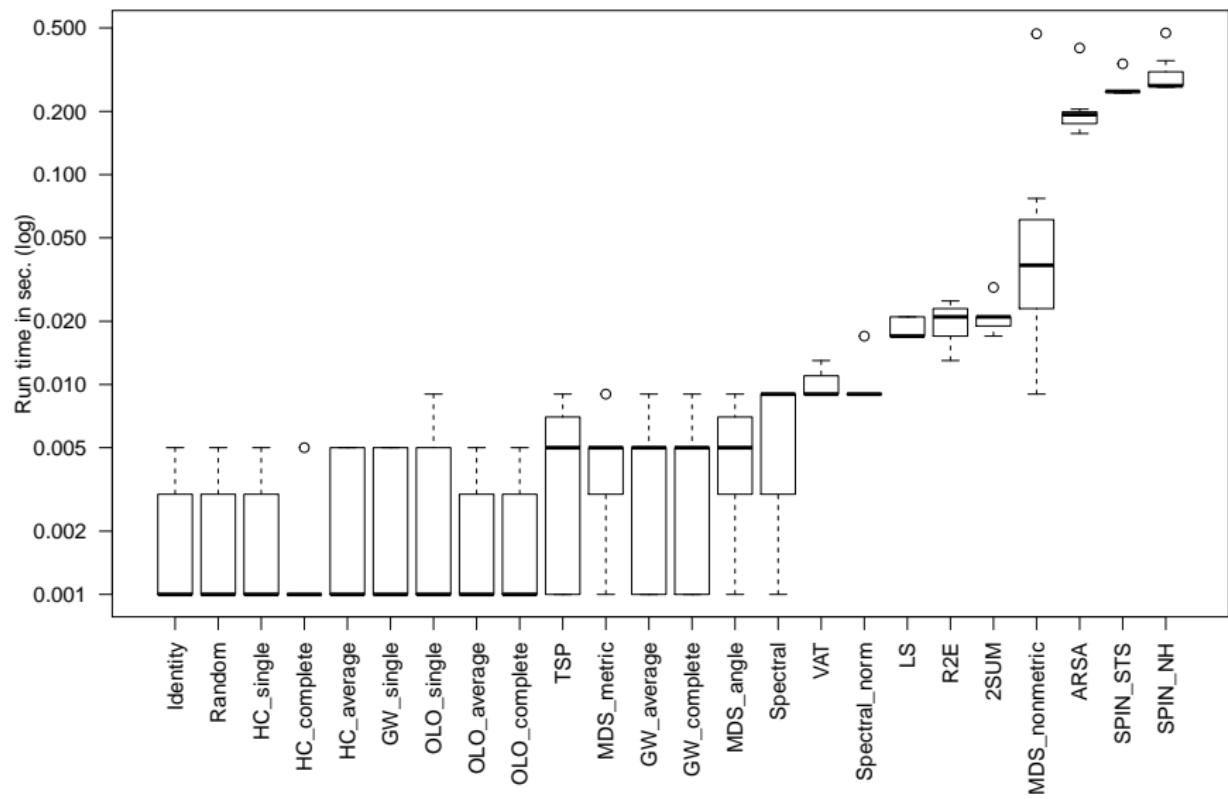
Comparison: Random Data



Comparison: Votes

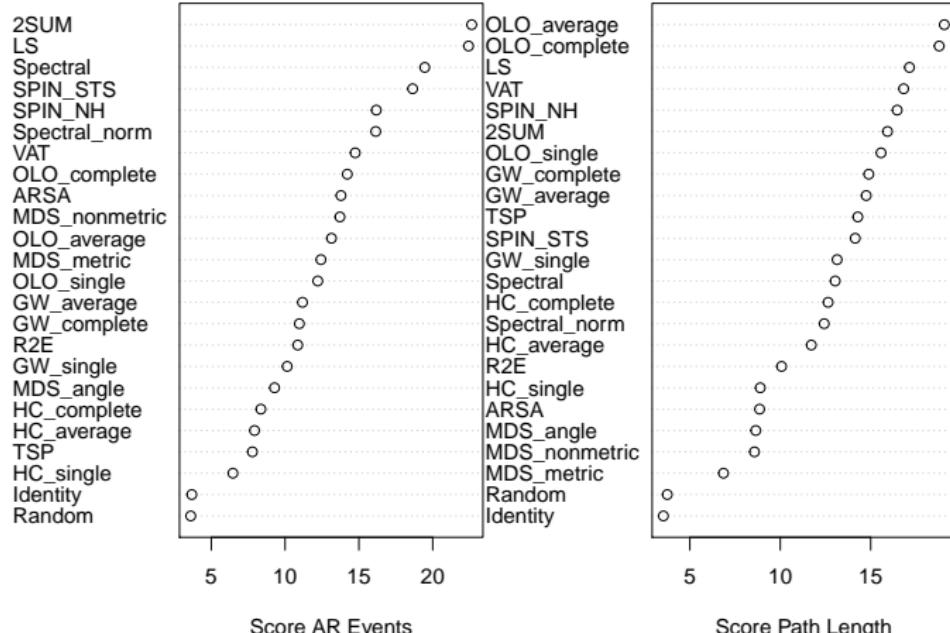


Computational Time



Comparison: Quality and Speed

- ① Rank each algorithm for quality and speed (1 point for the worse, 2 for the next, etc.)
- ② Average points over all data sets.



Conclusion

In this limited study we empirically found:

- There are two big groups of objective functions
 - ① **Anti Robinson Events (AR) / Gradient Measure**
 - ② **Path length**
- For group 1 (AR) **QAP formulations** provide a good tradeoff between quality and speed.
- For group 2 (path length) **Hierachical Clustering with Optimal Leaf Ordering** is fast and also provides quality comparable to TSP solvers.

Future work

- Even the heuristics (except for the TSP) are poorly scalable to applications with thousands of objects. Faster seriation heuristics are needed.

Thank you!

All seriation methods are available in the R package
seriation.

You can contact me at mhahsler@lyle.smu.edu

References I

- Arabie, P., and L. J. Hubert. 1996. "An Overview of Combinatorial Data Analysis." In *Clustering and Classification*, edited by P. Arabie, L. J. Hubert, and G. De Soete, 5–63. River Edge, NJ: World Scientific.
- Bar-Joseph, Z., E. D. Demaine, D. K. Gifford, and T. Jaakkola. 2001. "Fast Optimal Leaf Ordering for Hierarchical Clustering." *Bioinformatics* 17 (1): 22–29.
- Barnard, S. T., A. Pothen, and H. D. Simon. 1993. "A Spectral Algorithm for Envelope Reduction of Sparse Matrices." In *Proceedings of the 1993 ACM/IEEE Conference on Supercomputing*, 493–502. Supercomputing '93. New York, NY, USA: ACM.
- Bezdek, J.C., and R.J. Hathaway. 2002. "VAT: A Tool for Visual Assessment of (Cluster) Tendency." In *Proceedings of the 2002 International Joint Conference on Neural Networks (IJCNN '02)*, 3:2225–30.
- Brusco, M., H. F. Köhn, and S. Stahl. 2007. "Heuristic Implementation of Dynamic Programming for Matrix Permutation Problems in Combinatorial Data Analysis." *Psychometrika*.
- Brusco, Michael, and Stephanie Stahl. 2005. *Branch-and-Bound Applications in Combinatorial Data Analysis*. Springer.
- Caraux, G., and S. Pinloche. 2005. "Permutmatrix: A Graphical Environment to Arrange Gene Expression Profiles in Optimal Linear Order." *Bioinformatics* 21 (7): 1280–81.
- Chen, Chun-Houh. 2002. "Generalized Association Plots: Information Visualization via Iteratively Generated Correlation Matrices." *Statistica Sinica* 12 (1): 7–29.
- Ding, Chris, and Xiaofeng He. 2004. "Linearized Cluster Assignment via Spectral Ordering." In *In ICML '04: Proceedings of the Twenty-First International Conference on Machine Learning*, 30. ACM Press.
- Gruvaeus, G., and H. Wainer. 1972. "Two Additions to Hierarchical Cluster Analysis." *British Journal of Mathematical and Statistical Psychology* 25: 200–206.
- Gutin, G., and A. P. Punnen, eds. 2002. *The Traveling Salesman Problem and Its Variations*. Vol. 12. Combinatorial Optimization. Dordrecht: Kluwer.
- Hubert, L. J. 1974. "Some Applications of Graph Theory and Related Nonmetric Techniques to Problems of Approximate Seriation: The Case of Symmetric Proximity Measures." *British Journal of Mathematical Statistics and Psychology* 27: 133–53.
- Hubert, Lawrence, and James Schultz. 1976. "Quadratic Assignment as a General Data Analysis Strategy." *British Journal of Mathematical and Statistical Psychology* 29 (2). Blackwell Publishing Ltd: 190–241.

References II

Hubert, Lawrence, Phipps Arabie, and Jacqueline Meulman. 1987. *Combinatorial Data Analysis: Optimization by Dynamic Programming*. Society for Industrial Mathematics.

McCormick, William T., Paul J. Schweitzer, and Thomas W. White. 1972. "Problem Decomposition and Data Reorganization by a Clustering Technique." *Operations Research* 20 (5): 993–1009.

Niermann, Stefan. 2005. "Optimizing the Ordering of Tables with Evolutionary Computation." *The American Statistician* 59 (1): 41–46.

Robinson, W. S. 1951. "A Method for Chronologically Ordering Archaeological Deposits." *American Antiquity* 16: 293–301.

Tien, Yin-Jing, Yun-Shien Lee, Han-Ming Wu, and Chun-Hou Chen. 2008. "Methods for Simultaneously Identifying Coherent Local Clusters with Smooth Global Patterns in Gene Expression Profiles." *BMC Bioinformatics* 9 (155): 1–6.

Tsafrir, Dan, Ilan Tsafrir, Liat Ein-Dor, Or Zuk, Daniel A. Notterman, and Eytan Domany. 2005. *Bioinformatics* 21 (10): 2301–8.