

Lecture 14

Multidimensional Scaling

Reading: Izenman Chapter 13
The main reference for these slides is from Dr. Markus Kalisch's Lecture Notes at <https://stat.ethz.ch/education/semesters/ss2012/ams/slides/v4.1.1.pdf>

DSA 8070 Multivariate Analysis

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Multidimensional Scaling
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Main Idea
Classical Multidimensional Scaling
Non-metric Multidimensional Scaling
14.1

Notes

Agenda

- 1 Main Idea
- 2 Classical Multidimensional Scaling
- 3 Non-metric Multidimensional Scaling

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Main Idea
Classical Multidimensional Scaling
Non-metric Multidimensional Scaling
14.2

Notes

Principal Component Analysis and Multidimensional Scaling

- Principal Component Analysis (PCA):
In PCA, one starts with n data points $y_i \in \mathbb{R}^p$, then tries to find a low-dimensional projection of these points, e.g., $x_1, \dots, x_n \in \mathbb{R}^r$ with $r < p$, in such a way as to maximize the variance (thus minimizing the reconstruction error)
- Multidimensional Scaling (MDS):
In MDS, instead of being given the data $Y = \{y_i\}_{i=1}^n$, a matrix of distances or dissimilarities between the data points, $D = \{d_{ij}\}_{i,j=1}^n$ is provided. The goal of MDS is to find a set of points in a low-dimensional Euclidean space \mathbb{R}^r , usually $r = 2$, whose inter-point distances are as close as possible to the $\{d_{ij}\}$ distances

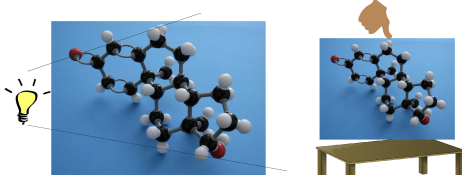
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Main Idea
Classical Multidimensional Scaling
Non-metric Multidimensional Scaling
14.3

Notes

Basic Idea of MDS

Represent a high-dimensional point cloud in a low (usually 2)-dimensional Euclidean space while *preserving, as closely as possible, the inter-point distances*. Commonly used MDS methods include classical/metric MDS and non-metric MDS:

- **Classical/Metric MDS:** Use a clever projection
- **Non-metric MDS:** Squeeze data on table



Source: Dr. Markus Kalisch's Lecture Notes on MDS

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 Main Idea
 Classical Multidimensional Scaling
 Non-metric Multidimensional Scaling
 144

Notes

Classical MDS (cMDS)

- **Goal:** Given pairwise distances among points, recover the position of the points!
- **Example:** Distance between 10 US major cities

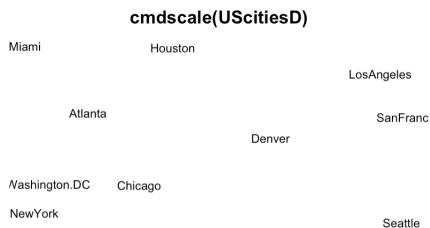
```
> UScitiesD
      Atlanta Chicago Denver Houston LosAngeles Miami NewYork SanFrancisco Seattle
Chicago      587
Denver      1212      920
Houston      701      940      879
LosAngeles  1936      1745      831      1374
Miami        604      1188      1726      968      2339
NewYork      748      713      1631      1420      2451      1092
SanFrancisco 2139      1858      949      1645      347      2594      2571
Seattle      2182      1737      1021      1891      959      2734      2408      678
Washington.DC 543      597      1494      1220      2300      923      205      2442      2329
```

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 Main Idea
 Classical Multidimensional Scaling
 Non-metric Multidimensional Scaling
 145

Notes

Classical MDS: First Try

```
loc <- cmdscale(UScitiesD)
x <- loc[, 1]; y <- loc[, 2]
plot(x, y, type = "n", xlab = "", ylab = "", asp = 1,
     axes = FALSE, main = "cmdscale(UScitiesD)")
text(x, y, rownames(loc), cex = 0.8)
''''
```

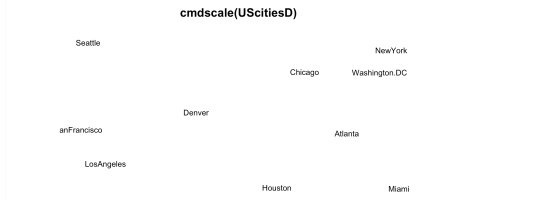


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 Classical Multidimensional Scaling
 Non-metric Multidimensional Scaling
 146

Notes

Classical MDS: Flip Axes

```
# Flip Axes
x1 <- -loc[, 1]; y1 <- -loc[, 2]
plot(x1, y1, type = "n", xlab = "", ylab = "", asp = 1,
     axes = FALSE, main = "cmdscale(UScitiesD)")
text(x1, y1, rownames(loc), cex = 0.8)
...
```



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14.7

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Another Example: Air Pollution in US Cities

> summary(dat)

S02	temp	manu	popul
Min. : 8.00	Min. :43.50	Min. : 35.0	Min. : 71.0
1st Qu.: 13.00	1st Qu.:50.60	1st Qu.: 181.0	1st Qu.: 299.0
Median : 26.00	Median :54.60	Median : 347.0	Median : 515.0
Mean : 30.05	Mean :55.76	Mean : 463.1	Mean : 608.6
3rd Qu.: 35.00	3rd Qu.:59.30	3rd Qu.: 462.0	3rd Qu.: 717.0
Max. :110.00	Max. :75.50	Max. :3344.0	Max. :3369.0

wind	precip	predays
Min. : 6.000	Min. : 7.05	Min. : 36.0
1st Qu.: 8.700	1st Qu.:30.96	1st Qu.:103.0
Median : 9.300	Median :38.74	Median :115.0
Mean : 9.444	Mean :36.77	Mean :113.9
3rd Qu.:10.600	3rd Qu.:43.11	3rd Qu.:128.0
Max. :12.700	Max. :59.80	Max. :166.0

- Range of manu and popul is much bigger than range of wind
- Need to standardize to give every variable equal weight

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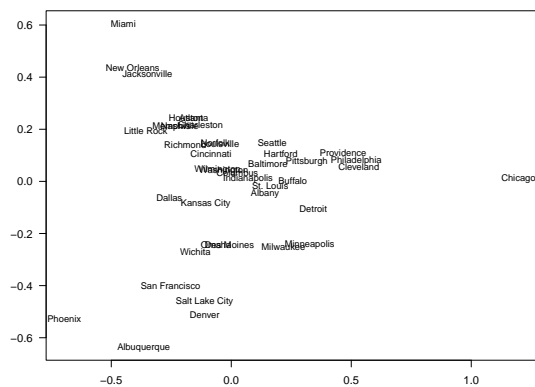
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14.8

Notes

Air Pollution in US Cities Example



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14.9

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Classical MDS: Technical Details

- **Input:** $D = \{d_{ij}\}_{i,j=1}^n$, the Euclidean distances between n objects in p dimensions
- **Output:** $X = \{x_i\}_{i=1}^n$, the “position” of points up to rotation, reflection, shift
- Two steps:
 - Compute inner products matrix $B = XX^T$ from distance

$$b_{ij} = -\frac{1}{2}(d_{ij}^2 - d_{i.}^2 - d_{.j}^2 + d_{..}^2)$$
 - Perform spectral decomposition to compute positions from B (see next slide)

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 Non-metric Multidimensional Scaling
 14.10

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Classical MDS: Technical Details

- Since $B = XX^T$, we need the “square root” of B
- Since B is a symmetric and positive definite $n \times n$ matrix $\Rightarrow B$ can be diagonalized:

$$B = V\Lambda V^T$$

Λ is a diagonal matrix with $\lambda_1 \leq \lambda_2 \leq \dots \leq \lambda_n$ on diagonal
- Assuming the rank of $B = p$, so that the last $n - p$ of its eigenvalues will be zero $\Rightarrow B$ can be written as

$$B = V_1\Lambda_1V_1^T,$$

where V_1 contains the first p eigenvectors and Λ_1 the p non-zero eigenvalues. Take “square root”:
 $X = V_1\Lambda_1^{-\frac{1}{2}}$

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 Non-metric Multidimensional Scaling
 14.11

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Classical MDS: Low-Dimensional Representation

- Keep only few (e.g. 2) largest eigenvalues and corresponding eigenvectors
- The resulting X will be the low-dimensional representation we were looking for
- “Goodness of fit” (GOF) if we reduce to r dimensions:

$$\text{GOF} = \frac{\sum_{i=1}^r \lambda_i}{\sum_{i=1}^n \lambda_i}$$
- Finds “optimal” low-dim representation:

Find $x_1, \dots, x_n \in \mathbb{R}^r$
 to minimize $\sum_{i=1}^n \sum_{j=1}^n (d_{ij} - d(x_i, x_j))^2$

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Classical MDS: Pros and Cons

- + Optimal for Euclidean input data
- + Still optimal, if B has non-negative eigenvalues
- + Very fast to compute
- - There is no guarantee it will be optimal if B has negative eigenvalues

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14.13

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Non-metric MDS: Idea

- Sometimes, there is no well-defined metric on original points
- Absolute values are not as meaningful, but the ranking is important, for example, in ordinal data and survey data (subjective preferences)
- Non-metric MDS finds a low-dimensional representation, which respects the ranking of distances

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14.14

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Non-metric MDS: Theory

- δ_{ij} is the true dissimilarity, d_{ij} is the distance of representation

- Minimize STRESS:

$$S = \frac{\sum_{i < j} (\theta(\delta_{ij}) - d_{ij})^2}{\sum_{i < j} d_{ij}^2},$$

where $\theta(\cdot)$ is an increasing function

- Optimize over both position of points and θ
- $\hat{d}_{ij} = \theta(\delta_{ij})$ is called "disparity"
- Solved numerically (isotonic regression); Classical MDS as starting value; very time consuming

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14.15

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Non-metric MDS: Pros and Cons

- +: Fulfills a clear objective (minimize STRESS) without many assumptions
- +: Results don't change with rescaling or monotonic variable transformation
- +: Works even if you only have rank information
- -: computation can be slow in "large" problems
- -: Usually only local (not global) optimum found
- -: Only gets ranks of distances right

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Non-metric Multidimensional Scaling

14.16

Notes

House of Representatives Voting Data

Romesburg (1984) gives a set of data that shows the number of times 15 congressmen from New Jersey voted differently in the House of Representatives on 19 environmental bills

```
> voting[1:6, 1:6]
      Hunt(R) Sandman(R) Howard(D) Thompson(D) Freylinghuysen(R) Forsythe(R)
Hunt(R)      0         8       15         15         10         9
Sandman(R)    8         0       17         12         13        13
Howard(D)    15        17        0         9         16        12
Thompson(D)  15        12        9         0         14        12
Freylinghuysen(R) 10        13        16        14         0         8
Forsythe(R)   9         13        12        12         8         0
```

Question: Do people in the same party vote alike?

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Classical Multidimensional Scaling

Non-metric Multidimensional Scaling

14.17

Notes

Kruskal's Non-metric Multidimensional Scaling in R

Usage

```
isoMDS(d, y = cmdscale(d, k), k = 2, maxit = 50, trace = TRUE, tol = 1e-3, p = 2)
```

Voting Example

```
library(MASS)
voting_mds <- isoMDS(voting, k = 2)
str(voting_mds)
par(las = 1, mar = c(2, 2, 0.5, 0.5))
plot(voting_mds$points, type = "n", xlim = c(-12, 8),
      xlab = "", ylab = "")
text(voting_mds$points, labels = rownames(voting_mds$points),
      cex = 0.7, col = col)
```

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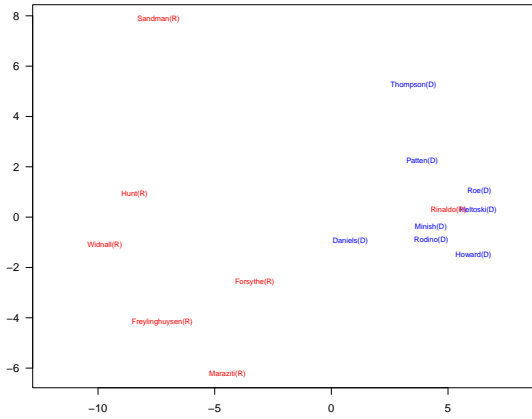
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14.18

Notes

Non-metric MDS: Voting Example



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Non-metric Multidimensional Scaling

14.19

Notes

Summary

- **Classical MDS:**
 - Finds low-dim projection that respects distances
 - Optimal for euclidean distances
 - No clear guarantees for other distances
 - Fast to compute (can use `cmdscale` in R)
- **Non-metric MDS:**
 - Squeezes data points on table
 - Respects only rankings of distances
 - (Locally) solves clear objective
 - Computation can be slow (can use `isoMDS` from the R package "MASS")

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Classical Multidimensional Scaling
Non-metric Multidimensional Scaling

14.20

Notes

Notes
