

Statistics 5401

34. Multidimensional Scaling

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Multidimensional scaling tries to find low dimensional representations of points, based originally on a matrix of distances between the points.

It is a cousin of principal components, but the original distances do not have to be Euclidean.

We have a distance matrix \mathbf{D} $n \times n$. We want to choose a dimension k (typically $k = 2$) and construct an $(n \times k)$ matrix \mathbf{X} so that the distances between the rows of \mathbf{X} match the corresponding elements of \mathbf{D} .

Then distances we see when plotting points in \mathbf{X} reflect the more complex, presumably high-dimensional distances coming from \mathbf{D} .

What we will actually try to do is match the squares of the distances between the rows with the squares of the distances in \mathbf{D} . The squared distance between $\vec{\mathbf{X}}_i$ and $\vec{\mathbf{X}}_j$ (d_{ij}^2) is

$$\begin{aligned}d_{ij}^2 &= \sum_{\ell=1}^k (X_{i\ell} - X_{j\ell})^2 \\ &= \sum_{\ell=1}^k X_{i\ell}^2 + \sum_{\ell=1}^k X_{j\ell}^2 - 2 \sum_{\ell=1}^k X_{i\ell} X_{j\ell}\end{aligned}$$

If we make a matrix of the d_{ij}^2 s, the i row contains an additive term of $\sum_{\ell=1}^k X_{i\ell}^2$, the j column contains an additive term of $\sum_{\ell=1}^k X_{j\ell}^2$, and the i, j th element contains an additive term of $-2 \sum_{\ell=1}^k X_{i\ell} X_{j\ell}$.

If we take the matrix of squared distances d_{ij}^2 , subtract the row means, then subtract the column means, and then take $-.5$ times the difference, we are left with a matrix with elements

$$\tilde{d}_{ij} = \sum_{\ell=1}^k X_{i\ell} X_{j\ell}$$

Look at this again, we get

$$\tilde{d}_{ij} = \sum_{\ell=1}^k X_{i\ell} X_{j\ell}$$

This expresses our (centered and rescaled) matrix of squared distances as a sum of outer products of the columns of \mathbf{X} .

Thus we can “recover” \mathbf{X} from the (centered and rescaled) matrix of squared distances \tilde{d} via SVD or an eigenvalue decomposition.

$$\tilde{d} = \mathbf{H}\mathbf{\Lambda}\mathbf{H}'$$

so

$$\mathbf{X} = \mathbf{H}\mathbf{\Lambda}^{1/2}$$

and

$$\tilde{d} = \mathbf{X}\mathbf{X}'$$

where $\Lambda^{1/2}$ is a diagonal matrix of the square roots of the eigenvalues of \tilde{d} .

Hold on! We haven't *really* recovered \mathbf{X} . Let \mathbf{H}_k be any $k \times k$ orthogonal matrix. Note that

$$\mathbf{Y} = \mathbf{X}\mathbf{H}_k = \mathbf{H}\Lambda^{1/2}\mathbf{H}_k$$

also satisfies

$$\tilde{d} = \mathbf{Y}\mathbf{Y}'$$

Thus we only recover \mathbf{X} up to some rotation.

So how do we do multidimensional scaling. We're trying to find an \mathbf{X} with distances between rows that match \mathbf{D} . So pretend that \mathbf{D} really came from \mathbf{X} and "recover" \mathbf{X} .

Square the elements in \mathbf{D} , subtract row means, subtract column means, multiply the difference by $-.5$, and then do an eigenvector/eigenvalue decomposition of the result. Rescale the first k eigenvectors by the square roots of the corresponding eigenvalues, and voila, we have

Classic (metric) Multidimensional Scaling.

```
Cmd> readdata("", school, x1, x2, x3, x4, x5, x6)
```

```
Read from file "~/JW5data/T12-9.DAT"
```

```
Column 1 saved as factor school
```

```
Column 2 saved as REAL vector x1
```

```
Column 3 saved as REAL vector x2
```

```
Column 4 saved as REAL vector x3
```

```
Column 5 saved as REAL vector x4
```

```
Column 6 saved as REAL vector x5
```

```
Column 7 saved as REAL vector x6
```

```
Cmd> X <- hconcat(x1, x2, x3, x4, x5, x6)
```

```
Cmd> X <- X/describe(X, stddev:T)'
```

```
Cmd> dim(X)
```

```
(1)          25          6
```

```
Cmd> D2 <- matrix(rep(0, 25*25), 25)
```

```
Cmd> for(i, run(6)) {  
D2 <- D2 + (X[,i]-X[,i]')^2  
;;  
}
```

```
Cmd> D2s <- D2
```

```
Cmd> D2s <- D2s - sum(D2s)/25
```

```
Cmd> D2s <- D2s - sum(D2s')'/25
```

```
Cmd> D2s <- -.5 * D2s
```

```
Cmd> eigenvals(D2s)
```

```
(1) 110.69 18.884 6.8775 3.9307  
(5) 2.9833 0.63482 lots of 0s
```

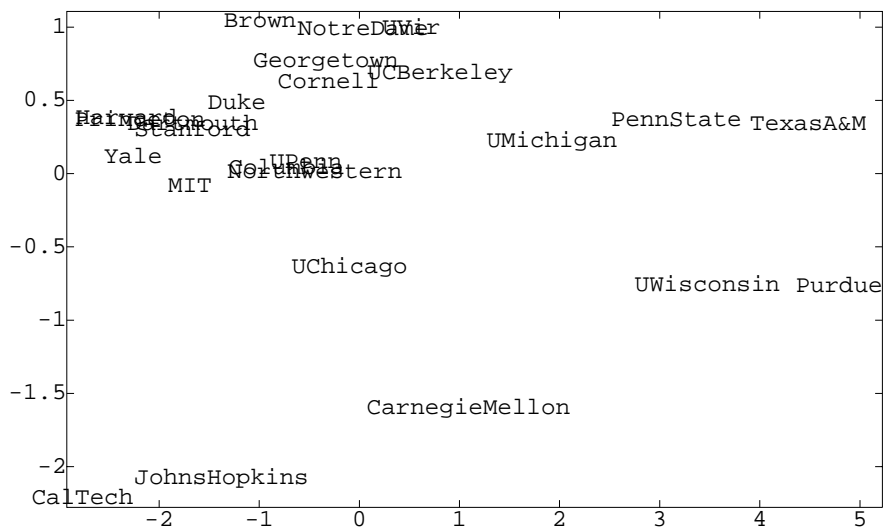
```
Cmd> (110.69+18.884)/sum(eigenvals(D2s))
```

```
(1) 0.89982
```

```
Cmd> Y <- eigen(D2s)$vectors[,run(2)]*\  
sqrt(eigenvals(D2s)[run(2)]')
```

```
Cmd> chplot(Y[,1],Y[,2]," ",xaxis:F,yaxis:F)
```

```
Cmd> addstrings(Y[,1],Y[,2],getlabels(school))
```



```
Cmd> s <- vecread("")
```

```
Read from file "~/JW5data/T12-4.DAT"
```

```
Cmd> S <- triunpack(s)
```

```
Cmd> D <- 10-S
```

```
Cmd> print(D,format:"f3.0",labels:F)
```

```
D:
```

```
0 2 2 7 6 6 6 6 7 9 9  
2 0 1 5 4 6 6 6 7 8 9  
2 1 0 6 5 6 5 5 6 8 9  
7 5 6 0 5 9 9 9 10 8 9
```

```

6  4  5  5  0  7  7  7  8  9  9
6  6  6  9  7  0  2  1  5 10  9
6  6  5  9  7  2  0  1  3 10  9
6  6  5  9  7  1  1  0  4 10  9
7  7  6 10  8  5  3  4  0 10  9
9  8  8  8  9 10 10 10 10  0  8
9  9  9  9  9  9  9  9  9  8  0

```

```
Cmd> D2 <- D^2
```

```
Cmd> D2s <- D2
```

```
Cmd> D2s <- D2s - sum(D2s)/11
```

```
Cmd> D2s <- D2s - sum(D2s')'/11
```

```
Cmd> D2s <- -.5 * D2s
```

```
Cmd> eigenvals(D2s)
```

```

(1)  110.8  71.209  31.683  21.895
(5)  13.598  8.5499  2.3585  0
(9) -0.06506 -1.0985 -3.1124

```

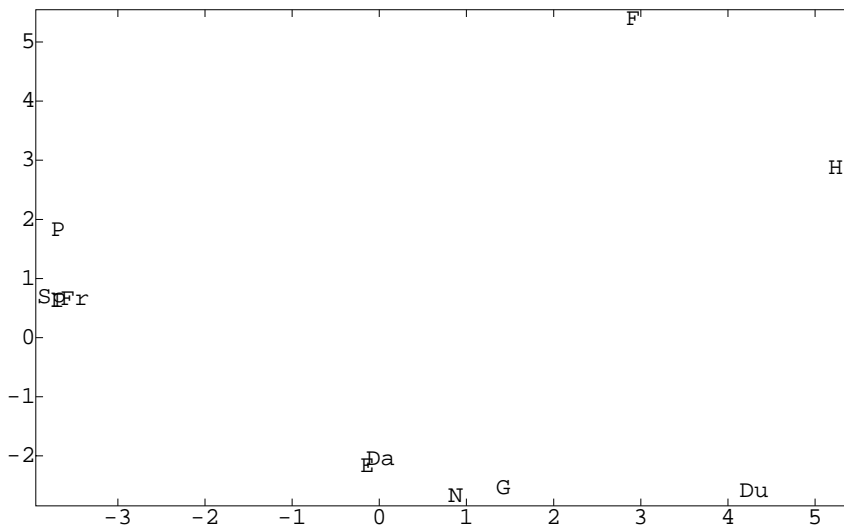
```
Cmd> (110.8+71.2)/sum(abs(eigenvals(D2s)))
```

```
(1)  0.68843
```

```
Cmd> Y <- eigen(D2s)$vectors[,run(2)]*\
sqrt(eigenvals(D2s)[run(2)]')
```

```
Cmd> lang <- vector("E", "N", "Da", "Du", \
"G", "Fr", "Sp", "I", "P", "H", "F")
```

```
Cmd> chplot(Y[,1],Y[,2],lang,xaxis:F,yaxis:F)
```



In some instances, we have dissimilarities, but not really distances. In particular, the difference of 1 between dissimilarities of 0 and 1 may not have any relation to the difference of 1 between dissimilarities of 9 and 10. In such a case, we want points \mathbf{X} such that the distances between the rows of \mathbf{X} have the same order as the dissimilarities, but the actual distances don't matter.

This is *Nonmetric Multidimensional Scaling*.

For any set of points \mathbf{X} , compute the distances d_{ij} .

Let \hat{d}_{ij} be an isotonic fit of these distances to the ordering from \mathbf{D} . This means that the \hat{d}_{ij} s are the closest numbers to the d_{ij} s that obey the correct ordering from \mathbf{D} . (Use the pool adjacent violators algorithm to get the isotonic fit.)

Define the *stress* to be

$$\text{Stress} = \left[\frac{\sum_{i < k} (d_{ik} - \hat{d}_{ik})^2}{\sum_{i < k} d_{ik}^2} \right]^{1/2}$$

Nonmetric MDS finds a matrix of points \mathbf{X} to minimize the stress. \mathbf{X} is not unique; rotations don't change the stress, and rescaling all the variables by the same factor doesn't change the stress.

Some people prefer to minimize the *SStress*

$$\text{SStress} = \left[\frac{\sum_{i < k} (d_{ik}^2 - \hat{d}_{ik}^2)^2}{\sum_{i < k} d_{ik}^4} \right]^{1/2}$$