MDS: Link Squared Distance Matrix and Gram Matrix in Practice

Lieven Clement

statOmics, Ghent University (https://statomics.github.io)

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1 Data

Part of the iris dataset

```
X <- iris[1:5,1:4] %>% as.matrix
X
```

```
Sepal.Length Sepal.Width Petal.Length Petal.Width
##
## 1
              5.1
                           3.5
                                        1.4
                                                     0.2
              4.9
                           3.0
                                        1.4
                                                     0.2
              4.7
                           3.2
                                        1.3
                                                     0.2
## 3
## 4
              4.6
                           3.1
                                        1.5
                                                     0.2
## 5
              5.0
                           3.6
                                        1.4
                                                     0.2
```

2 Centering

$$\begin{aligned} \mathbf{H} &= \mathbf{I} - \frac{1}{n}\mathbf{1}\mathbf{1}^T, \\ \mathbf{X}_c &= \mathbf{H}\mathbf{X} \\ \mathbf{H}\mathbf{X}_c &= \mathbf{X}_c \end{aligned}$$

```
H <- diag(nrow(X)) - matrix(1/nrow(X),nrow=nrow(X),ncol=nrow(X))
Xc <- H%*%X
colMeans(Xc)

## Sepal.Length Sepal.Width Petal.Length Petal.Width
## 1.776357e-16 -8.881784e-17 -4.440892e-17 0.0000000e+00

H%*%Xc-Xc

## Sepal.Length Sepal.Width Petal.Length Petal.Width
## [1,] -1.665335e-16 8.326673e-17 4.440892e-17 0
## [2,] -1.804112e-16 1.110223e-16 4.440892e-17 0
## [3,] -1.665335e-16 8.326673e-17 4.163336e-17 0
## [4,] -1.665335e-16 8.326673e-17 4.163336e-17 0
## [5,] -1.665335e-16 1.110223e-16 4.440892e-17 0</pre>
```

3 Gram matrix

Gram matrix: $\mathbf{G} = \mathbf{X}\mathbf{X}^T$ Here we work on centered data so $\mathbf{G} = \mathbf{X}_c\mathbf{X}_c^T$

```
G <- Xc%*%t(Xc)
```

4 Squared Distance matrix

[3,] 0.26 0.09 0.00 0.06 0.26 ## [4,] 0.42 0.11 0.06 0.00 0.42 ## [5,] 0.02 0.37 0.26 0.42 0.00

$$\mathbf{D}_X = \mathbf{N} - 2\mathbf{X}\mathbf{X}^T + \mathbf{N}^T,$$

```
N <- matrix(diag(G),nrow(Xc),nrow(Xc))

## [,1] [,2] [,3] [,4] [,5]

## [1,] 0.106 0.106 0.106 0.106 0.106
## [2,] 0.080 0.080 0.080 0.080 0.080
## [3,] 0.042 0.042 0.042 0.042
## [4,] 0.110 0.110 0.110 0.110 0.110
## [5,] 0.122 0.122 0.122 0.122 0.122

dist2 <- N-2*G+t(N)
dist2

## [,1] [,2] [,3] [,4] [,5]
## [1,] 0.00 0.29 0.26 0.42 0.02
## [2,] 0.29 0.00 0.09 0.11 0.37</pre>
```

dist(Xc)^2

```
## 1 2 3 4
## 2 0.29
## 3 0.26 0.09
## 4 0.42 0.11 0.06
## 5 0.02 0.37 0.26 0.42
```

5 Link Gram Matrix and Squared Distance Matrix

$$\mathbf{G} = \mathbf{X}_c \mathbf{X}_c^T = -\frac{1}{2} \mathbf{H} \mathbf{D}_X \mathbf{H}$$

G

```
## [,1] [,2] [,3] [,4] [,5]

## [1,] 0.106 -0.052 -0.056 -0.102 0.104

## [2,] -0.052 0.080 0.016 0.040 -0.084

## [3,] -0.056 0.016 0.042 0.046 -0.048

## [4,] -0.102 0.040 0.046 0.110 -0.094

## [5,] 0.104 -0.084 -0.048 -0.094 0.122
```

-1/2*H%*%dist2%*%H

```
## [,1] [,2] [,3] [,4] [,5]

## [1,] 0.106 -0.052 -0.056 -0.102 0.104

## [2,] -0.052 0.080 0.016 0.040 -0.084

## [3,] -0.056 0.016 0.042 0.046 -0.048

## [4,] -0.102 0.040 0.046 0.110 -0.094

## [5,] 0.104 -0.084 -0.048 -0.094 0.122
```

-1/2*H%*%dist2%*%H - G

```
## [,1] [,2] [,3] [,4] [,5]

## [1,] -2.775558e-17 -7.632783e-17 -2.775558e-17 1.387779e-17 0.000000e+00

## [2,] -7.632783e-17 -5.551115e-17 -1.387779e-17 1.387779e-17 -2.775558e-17

## [3,] -2.081668e-17 -1.387779e-17 2.775558e-17 6.245005e-17 2.775558e-17

## [4,] 1.387779e-17 6.938894e-18 5.551115e-17 8.326673e-17 0.000000e+00

## [5,] 0.000000e+00 -4.163336e-17 2.081668e-17 0.000000e+00 4.163336e-17
```

6 Show that N cancels out when multiplying with H

$$HNH = 0$$

$$\mathbf{HN}^T\mathbf{H} = \mathbf{0}$$

N**%*%**H

```
## [,1] [,2] [,3] [,4] [,5]
## [1,] 0.000000e+00 0.000000e+00 0 0.000000e+00 0
## [2,] -6.938894e-18 -6.938894e-18 0 3.469447e-18 0
## [4,] -6.938894e-18 -6.938894e-18 0 6.938894e-18 0
## [5,] -6.938894e-18 -6.938894e-18 0 3.469447e-18 0
```

H%*%t(N)