# smacof Data Structures

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### Metric MDS

The input for a metric (ratio) smacof consists of four vectors ik, jk, sk, and wk of the same length.

Elements ik[k], jk[k], dk[k], and wk[k] give the row index, the column index, the dissimilarity value, and the weight value for observation k. There can be replications and data can be asymmetric.

#### We assume

- the vector dk of dissimilarities is in non-decreasing order,
- dk is non-negative,
- the vector wk of weights is strictly positive,
- for all  $k = 1, \dots, m$  index ik[k] is not equal to index jk[k].

If the dissimilarity data come in a square matrix

and all weights for non-missing data are one, then then a corresponding smacof data structure is

```
ik jk dk wk
     2
[1,]
            1
[2,]
     1
         2
            2
               1
[3,]
     1
         3
            2
               1
[4,] 3 1
            2
```

```
[5,] 1 4 3 1
[6,] 4 1 3 1
[7,] 3 2 4 1
[8,] 2 4 5 1
[9,] 4 2 5 1
[10,] 3 4 8 1
[11,] 4 3 8 1
```

Because there are ties in the dissimilarities the data structure corresponding to the matrix is not unique. From the point of view of minimizing stress an equivalent data structure is

$$\sigma(X) = \frac{1}{2} \sum_{k=1}^{m} w_k (\delta_k - d_k(X))^2$$
 
$$d_k(X) = \sqrt{\operatorname{tr} X' A_k X}$$

 $A_k := (e_{i_k} - e_{j_k})(e_{i_k} - e_{j_k})'$ 

The  ${\cal A}_k$  are intended to be used in formulas, not in actual computation.

$$V = \sum_{k=1}^{m} w_k A_k$$

```
smacofMakeVmat <- function(dat) {
  ndat <- nrow(dat)
  nobj <- max(max(dat[, 1], max(dat[, 2])))
  vmat <- matrix(0, nobj, nobj)
  for (k in 1:ndat) {
    i <- dat[k, 1]
    j <- dat[k, 2]
    w <- dat[k, 4]
    vmat[i, j] <- vmat[i, j] - w
    vmat[j, i] <- vmat[i, j]</pre>
```

```
diag(vmat) <- -rowSums(vmat)
return(vmat)
}

print(smacofMakeVmat(thedata1))</pre>
```

```
[,1] [,2] [,3] [,4]
[1,] 6 -2 -2 -2
[2,] -2 5 -1 -2
[3,] -2 -1 5 -2
[4,] -2 -2 -6
```

#### print(smacofMakeVmat(thedata2))

```
[,1] [,2] [,3] [,4]
[1,] 6 -2 -2 -2
[2,] -2 5 -1 -2
[3,] -2 -1 5 -2
[4,] -2 -2 -6
```

```
smacofDistance <- function(dat, x) {
  ndat <- nrow(dat)
  dmat <- rep(0, ndat)
  for (k in 1:ndat) {
    i <- dat[k, 1]
    j <- dat[k, 2]
    dmat[k] <- sqrt(sum((x[i, ] - x[j, ]) ^ 2))
  }
  return(dmat)
}</pre>
```

```
dmat1 <- smacofDistance(thedata1, x)
dmat2 <- smacofDistance(thedata2, x)
print(dmat2)</pre>
```

[1] 1.000000 1.000000 1.414214 1.414214 1.000000 1.000000

```
smacofMakeBmat <- function(dat, dmat) {</pre>
  ndat <- nrow(dat)</pre>
  nobj <- max(max(dat[, 1]), max(dat[, 2]))</pre>
  bmat <- matrix(0, nobj, nobj)</pre>
  for (k in 1:ndat) {
    i <- dat[k, 1]
    j <- dat[k, 2]
    w \leftarrow dat[k, 4]
    e \leftarrow dat[k, 3]
    d <- dmat[k]</pre>
    bmat[i, j] \leftarrow bmat[i, j] - w * (e / d)
    bmat[j, i] <- bmat[i, j]</pre>
  diag(bmat) <- -rowSums(bmat)</pre>
  return(bmat)
print(smacofMakeBmat(thedata2, dmat2))
                                  [,3]
           [,1]
                      [,2]
                                              [,4]
[1,] 11.242641 -3.000000 -4.000000 -4.242641
[2,] -3.000000 15.828427 -2.828427 -10.000000
[3,] -4.000000 -2.828427 22.828427 -16.000000
[4,] -4.242641 -10.000000 -16.000000 30.242641
print(smacofMakeBmat(thedata1, dmat1))
           [,1]
                      [,2]
                                  [,3]
                                              [,4]
[1,] 11.242641 -3.000000 -4.000000 -4.242641
[2,] -3.000000 15.828427 -2.828427 -10.000000
[3,] -4.000000 -2.828427 22.828427 -16.000000
[4,] -4.242641 -10.000000 -16.000000 30.242641
smacofStress <- function(dat, dmat) {</pre>
   return(sum(dat[, 4] * (dat[, 3] - dmat) ^ 2))
```

[1] 144.2157

print(smacofStress(thedata2, dmat2))

print(smacofStress(thedata1, dmat1))

[1] 144.7157

$$B(X) = \sum_{k=1}^m \{w_k \frac{\delta_k}{d_k(X)} A_k \mid d_k(X) > 0\}$$

# **Semimetric**

this has a numeric delta, and we add a column with tie blocks this is for splines etc and for the shepard plot

# **Nonmetric**

delta are rank numbers with ties getting the same rank number

# **Pairs and Triads**

## **Nominal**

# **Individual Differences**