# Appendix: Cluster Analysis

More information can be found in Everitt, Landau & Leese, 2001. *Custer Analysis*. Oxford University Press

# Similarity and Dissimilarity Metrics

• The similarity metric has to satisfy the properties:

```
Symmetry: s_{ij} = s_{ji} for all i, j \in \{1, 2, ..., n\}
Ordering: s_{ij} \le s_{ii} similarity of itself larger than else Self-similarity: s_{ii} = s_{jj}
```

#### o Notes:

- Correlations can be considered similarities on the scale  $-1 \le r_{ij} \le 1$
- The topological spatial adjacency can be considered a similarity measure.
- Frequently it is assumed that the similarity  $s_{ij} \ge 0$ .
- There is no upper limit for the similarity unless the assumption is made that  $s_{11} = \cdots = s_{nn} = 1$ .
- The similarities among all *n*objects can be pooled into a similarity matrix  $S_{n\times n}$ .
- The dissimilarity metric has to satisfy the properties

Symmetry: 
$$d_{ij} = d_{ji}$$
 for all  $i, j \in \{1, 2, ..., n\}$ 

Ordering: 
$$d_{ij} \geq 0$$
 and  $d_{11} = \cdots = d_{nn} = 0$ 

- o Notes:
  - If a dissimilarity metric also satisfies the triangle equation  $d_{ij} \le d_{ik} + d_{jk}$  then it has a geometric interpretation as a distance measure.
  - A dissimilarity metric has a natural lower bound of zero.
  - The dissimilarities among all *n*objects can be pooled into a dissimilarity matrix  $\mathbf{D}_{n \times n}$ .
- Possible transformations between both metrics:
  - An inverse relationship between both metrics exists  $d_{ij} \approx \frac{1}{s_{ij}}$
  - For  $0 \le s_{ij} \le 1$  the transformation becomes:  $d_{ij} = 1 s_{ij}$
  - For  $-1 \le s_{ij} \le 1$  the transformation becomes  $d_{ij} = \frac{1}{2} \cdot (1 s_{ij})$
  - $\circ s_{ij} = 1 \frac{d_{ij}}{\max(d_{ij})}$

#### **Nominal Scaled Variables**

• Objects with *nominal* scaled features:

- Nominal scaled features with K factor levels must be encoded [a] using either the one-hot coding with K binary variables or the dummy coding with K − 1 binary variables (see Boehmke et al. pp 58-66). p should be k here
- o Let  $\mathbf{x}_i = (b_{i1}, ..., b_{iP})^T$  and  $\mathbf{x}_j = (b_{j1}, ..., b_{jP})^T$  where each object is characterized by P binary features  $b_{ip} = \begin{cases} 1 & \text{if feature } p \text{ is present} \\ 0 & \text{otherwise} \end{cases}$  and analogue for  $b_{jp}$
- Both feature vectors can be organized in a contingency table counting the concordant and discordant pairs of features:

|   | 1                          | 0                          |                            |
|---|----------------------------|----------------------------|----------------------------|
| 1 | $c_{11}$                   | $c_{10}$                   | $c_{11} + c_{10} = c_{1+}$ |
| 0 | c <sub>01</sub>            | c <sub>00</sub>            | $c_{01} + c_{00} = c_{0+}$ |
|   | $c_{11} + c_{01} = c_{+1}$ | $c_{10} + c_{00} = c_{+0}$ | P                          |

- The number of concordant pairs is  $c_{11} + c_{00}$  and the number of discordant pairs is  $c_{10} + c_{01}$
- Several similarity and dissimilarity metrics can be derived from this table. See
   help(dist) and the options binary:

```
> x <- c(0, 0, 1, 1, 1, 1)
> y <- c(1, 0, 1, 1, 0, 1)
> dist(rbind(x, y), method = "binary")
```

This is a *dissimilarity* measure because  $d_{ij} = \frac{c_{01} + c_{10}}{c_{01} + c_{10} + c_{11}}$ . Here missing feature pairs  $c_{00}$  are not counted because neither feature is at all present.

o A generic error rate similarity metric is

$$s_{ij} = \frac{\alpha \cdot (c_{11} + c_{00})}{\alpha \cdot (c_{11} + c_{00}) + (1 - \alpha) \cdot (c_{10} + c_{01})}$$

where  $0 < \alpha < 1$  weights concordant and discordant pairs differently for  $\alpha = 0.5$  we obtain the concordance rate  $s_{ij} = (c_{11} + c_{00})/P$ .

o A measure analog to a correlation coefficient between the pairs  $\mathbf{x}_i$  and  $\mathbf{x}_j$  can be obtained by

$$s_{ij} = \frac{c_{11} \cdot c_{00} - c_{10} \cdot c_{01}}{\sqrt{c_{1+} \cdot c_{0+} \cdot c_{+1} \cdot c_{+0}}}$$

### **Ordinal Scaled Variables**

• For *ordinal* scaled features p similar coefficients can be calculated by recognizing the ranking of features includes lower ranked features. Let  $p_1 < p_2 < p_3$  then

$$p_1 = \begin{pmatrix} 1 \\ 0 \\ 0 \end{pmatrix}$$
,  $p_2 = \begin{pmatrix} 1 \\ 1 \\ 0 \end{pmatrix}$  and  $p_3 = \begin{pmatrix} 1 \\ 1 \\ 1 \end{pmatrix}$ 

#### **Metric Variables**

• For metrically scaled variables the generic *Minkowski* dissimilarity metric is

$$d_{ij}^{[q]} = \left(\sum_{p=1}^{P} |x_{ip} - x_{jp}|^{q}\right)^{1/q}$$

- $\circ$  For q = 1 the Manhattan (city block) distance is obtained
- $\circ$  For q = 2 the Euclidian distance is obtained
- For  $q = \infty$  the maximum distance  $\max_{p} |x_{ip} x_{jp}|$  is obtained.
- Only the Euclidian distance are rotation invariant.
- $\circ$  The squared Euclidian distance  $d_{ij}^2$  is closely related to variance around a centroid, such the within class heterogeneity measure.
- The *Mahalanobis* measures the Euclidian distance between objects based on their uniformly scaled principle component scores.

$$d_{ij}^{[M]} = (\mathbf{x}_i - \mathbf{x}_j)^T \cdot \mathbf{S}^{-1} \cdot (\mathbf{x}_i - \mathbf{x}_j)$$

with  $S^{-1}$  being the inverse covariance matrix an  $\mathbf{x}_i = (x_{i1}, x_{i2}, ..., x_{iP})^T$  being the vector of the P features for the i<sup>th</sup> observation and analogue for the j<sup>th</sup> observation, or

$$d_{ij}^{[M]} = (\mathbf{z}_i - \mathbf{z}_j)^T \cdot \mathbf{R}^{-1} \cdot (\mathbf{z}_i - \mathbf{z}_j)$$

where  $\mathbf{z}_i$  and  $\mathbf{z}_j$  are based on z-transformed variables with  $\mathbf{z}_i = (z(x_{i1}), z(x_{i2}), ..., z(x_{iP}))^T$  and  $\mathbf{R}^{-1}$  is the inverse correlation matrix among all variables (see script **Mahalanobis**.**R**).

- o Notes:
  - It standardized the variance of each variable.
  - It controls for the correlation among the features and removes data redundancy.
  - The Mahalanobis distance is based on all observations. Therefore, within a cluster the features may very well be correlated.

#### **Mixture of Measurement Levels**

• For objects based on a *mixture* of feature types, a similarity metric based on a linear combination can be derived

$$s_{ij} = \frac{\left| p^{nominal} \right| \cdot s_{ij}^{nominal} + \left| p^{ordinal} \right| \cdot s_{ij}^{ordinal} + \left| p^{metric} \right| \cdot s_{ij}^{metric}}{P}$$

with  $P = |p^{nominal}| + |p^{ordinal}| + |p^{metric}|$  where | | is the number of features contributing to each similarity measure.

- The Gower option in **cluster::daisy( , metric='gower')** calculates a mixture dissimilarity measure for metric and nominal scaled features:
  - o For nominal scaled features it reports discordant pairs as 1 and concordant pairs as zero.

- $\circ$  For metric or ordinal scaled features it reports the absolute difference for each feature and then it scales each  $d_{ij}^p$  for the feature p to  $d_{ij}^p$  in 0 to 1.
- o Finally, it adds the distances of all features together either as a mean or weighted mean:

$$\mathbf{D} = \frac{\sum_{p=1}^{P} w_p \cdot \mathbf{D}_p}{\sum_{p=1}^{P} w_p}$$

- Notes:
  - Hierarchical cluster analysis has the tendency to group objects with similar levels of nominal scaled features together.
  - o Principal component analysis and k nearest neighbors, kMeans are not well tailored using binary features. For instance, performing a *z*-transformation on binary features does not dissolve the binning around just two values.

# **Generic Hierarchically Cluster Analysis Equation**

• If two classes  $C_k \cup C_l$  are merged then the heterogeneity increase to any of the remaining classes  $C_s$  can be calculated *recursively* by

$$d(C_k \cup C_l, C_s) = \alpha_k \cdot d(C_k, C_s) + \alpha_l \cdot d(C_l, C_s) + \beta \cdot d(C_k, C_l) + \gamma \cdot |d(C_k, C_s) - d(C_l, C_s)|$$

• Depending on the selected parameters values  $\alpha_k$ ,  $\alpha_l$ ,  $\beta$  and  $\gamma$  the different heterogeneity update methods can be derived.

• Only if  $\alpha_k$ ,  $\alpha_l \ge 0$  and  $\alpha_k + \alpha_l + \beta \ge 1$  as well as  $|\gamma| \le \alpha_k$ ,  $\alpha_l$  then the heterogeneity of the clustering method is monotonically increasing and does not exhibit inversions.

• Parameters of recursive agglomerative cluster algorithms

| Linkage  | $\alpha_k$        | $\alpha_l$        | β                            | γ    |
|----------|-------------------|-------------------|------------------------------|------|
| Methods  |                   |                   |                              |      |
| Single   | 1/2               | 1/2               | 0                            | -1/2 |
| Complete | 1/2               | 1/2               | 0                            | 1/2  |
| Average  | $n_k/(n_k+n_s)$   | $n_l/(n_l+n_s)$   | 0                            | 0    |
| Centroid | $n_k/(n_k+n_s)$   | $n_l/(n_l+n_s)$   | $-\frac{n_k \cdot n_l}{n_l}$ | 0    |
|          |                   |                   | $(n_k + n_l)^2$              |      |
| Ward     | $n_k + n_s$       | $n_l + n_s$       | $-\frac{n_s}{}$              | 0    |
|          | $n_k + n_l + n_s$ | $n_k + n_l + n_s$ | $n_k + n_l + n_s$            |      |
| Median   | 1/2               | 1/2               | -1/4                         | 0    |
| Flexible | $\varphi > 0$     | $\varphi$         | $1-2\cdot \varphi$           | 0    |
| strategy |                   |                   |                              |      |