Distance and dissimilarities

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knitr::opts_chunk$set(echo = TRUE)
install.packages("dplyr")
## Installing package into '/home/rstudio-user/R/x86_64-pc-linux-gnu-library/4.0'
## (as 'lib' is unspecified)
install.packages("stargazer")
## Installing package into '/home/rstudio-user/R/x86_64-pc-linux-gnu-library/4.0'
## (as 'lib' is unspecified)
```

Definition of a distance

- A distance function or a metric on \mathbb{R}^n , $n \geq 1$, is a function $d: \mathbb{R}^n \times \mathbb{R}^n \to \mathbb{R}$.
- A distance function must satisfy some required properties or axioms.
- There are three main axioms.
- A1. $d(\mathbf{x}, \mathbf{y}) = 0 \iff \mathbf{x} = \mathbf{y}$ (identity of indiscernibles);
- A2. $d(\mathbf{x}, \mathbf{y}) = d(\mathbf{y}, \mathbf{x})$ (symmetry);
- A3. $d(\mathbf{x}, \mathbf{z}) \leq d(\mathbf{x}, \mathbf{y}) + d(\mathbf{y}, \mathbf{z})$ (triangle inequality), where $\mathbf{x} = (x_1, \dots, x_n)$, $\mathbf{y} = (y_1, \dots, y_n)$ and $\mathbf{z} = (z_1, \dots, z_n)$ are all vectors of \mathbb{R}^n .
- We should use the term *dissimilarity* rather than *distance* when not all the three axioms A1-A3 are valid.
- Most of the time, we shall use, with some abuse of vocabulary, the term distance.

Exercice 1

• Prove that the three axioms A1-A3 imply the non-negativity condition:

$$d(\mathbf{x}, \mathbf{y}) \ge 0.$$

Euclidean distance

• It is defined by:

$$d(\mathbf{x}, \mathbf{y}) = \sqrt{\sum_{i=1}^{n} (x_i - y_i)^2}.$$

Manhattan distance

• The Manhattan distance also called taxi-cab metric or city-block metric is defined by:

$$d(\mathbf{x}, \mathbf{y}) = \sum_{i=1}^{n} |x_i - y_i|.$$

- A1-A2 hold.
- A3 also holds using the fact that $|a+b| \le |a| + |b|$ for any reals a, b.
- There exists also a weighted version of the Manhattan distance called the Canberra distance.

Manhattan distance vs Euclidean distance Graph

```
x = c(0, 0)

y = c(6,6)

dist(rbind(x, y), method = "euclidian")
```

^{*} A1A2 ae onbvious. * The proof of A3 is provided below.

```
## x
## y 8.485281
6*sqrt(2)
## [1] 8.485281
dist(rbind(x, y), method = "manhattan")
## x
## y 12
```

Canberra distance

• It is defined by:

$$d(\mathbf{x}, \mathbf{y}) = \sum_{i=1}^{n} \frac{|x_i - y_i|}{|x_i| + |y_i|}.$$

- Note that the term $|x_i y_i|/(|x_i| + |y_i|)$ is not properly defined when $x_i = y_i = 0$.
- By convention we set the ratio to be zero in that case.
- The Canberra distance is specially sensitive to small changes near zero.

```
x = c(0, 0)
y = c(6,6)
dist(rbind(x, y), method = "canberra")

##  x
## y 2
6/6+6/6

## [1] 2
```

Exercice 2

• Prove that the Canberra distance is a true distance.

Minkowski distance

• Both the Euclidian and the Manattan distances are special cases of the Minkowski distance which is defined, for $p \ge 1$, by:

$$d(\mathbf{x}, \mathbf{y}) = \left[\sum_{i=1}^{n} |x_i - y_i|^p \right]^{1/p}.$$

- For p = 1, we get the Manhattan distance.
- For p = 2, we get the Euclidian distance.
- Let us also define:

$$\|\mathbf{x}\|_p \equiv \left[\sum_{i=1}^n |x_i|^p\right]^{1/p},$$

where $\|\cdot\|_p$ is known as the *p*-norm or Minkowski norm.

• Note that the Minkowski distance and norm are related by:

$$d(\mathbf{x}, \mathbf{y}) = \|\mathbf{x} - \mathbf{y}\|_{p}.$$

• Conversely, we have:

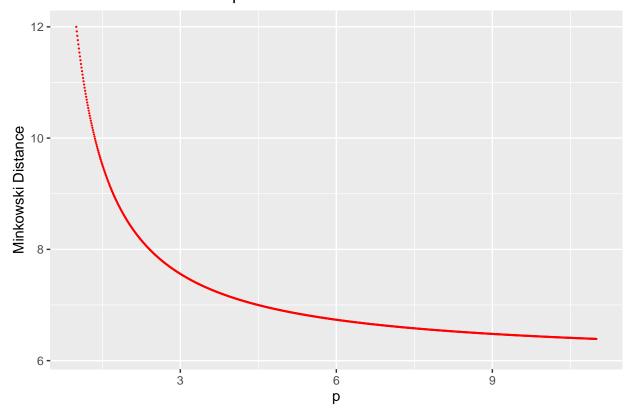
$$\|\mathbf{x}\|_p = d(\mathbf{x}, \mathbf{0}),$$

where **0** is the null-vetor of \mathbb{R}^n .

```
library("ggplot2")
x = c(0, 0)
y = c(6,6)
MinkowDist=c()
for (p in seq(1,30,.01))
{
MinkowDist=c(MinkowDist,dist(rbind(x, y), method = "minkowski", p = p))
}
ggplot(data =data.frame(x = seq(1,30,.01), y=MinkowDist ) , mapping = aes(x = x, y = y))+geom_point(siz
```

Warning: Removed 1900 rows containing missing values (geom_point).

Minkowski distance wrt p



Chebyshev distance

• At the limit, we get the Chebyshev distance which is defined by:

$$d(\mathbf{x}, \mathbf{y}) = \max_{i=1,\dots,n} (|x_i - y_i|) = \lim_{p \to \infty} \left[\sum_{i=1} |x_i - y_i|^p \right]^{1/p}.$$

• The corresponding norm is:

$$\|\mathbf{x}|_{\infty} = \max_{i=1,\dots,n} (|x_i|).$$

Minkowski inequality

- The proof of the triangular inequality A3 is based on the Minkowski inequality:
- For any nonnegative real numbers $a_1, \dots, a_n; b_1, \dots, b_n$, and for any $p \ge 1$, we have:

$$\left[\sum_{i=1}^{n} (a_i + b_i)^p\right]^{1/p} \le \left[\sum_{i=1}^{n} a_i^p\right]^{1/p} + \left[\sum_{i=1}^{n} b_i^p\right]^{1/p}.$$

• To prove that the Minkowski distance satisfies A3, notice that

$$\sum_{i=1}^{n} |x_i - z_i|^p = \sum_{i=1}^{n} |(x_i - y_i) + (y_i - z_i)|^p.$$

• Since for any reals x, y, we have: $|x + y| \le |x| + |y|$, and using the fact that x^p is increasing in $x \ge 0$, we obtain:

$$\sum_{i=1}^{n} |x_i - z_i|^p \le \sum_{i=1}^{n} (|x_i - y_i| + |y_i - z_i|)^p.$$

• Applying the Minkowski inequality with $a_i = |x_i - y_i|$ and $b_i = |y_i - z_i|, i = 1, \dots, n$, we get:

$$\sum_{i=1}^{n} |x_i - z_i|^p \le \left(\sum_{i=1}^{n} |x_i - y_i|^p\right)^{1/p} + \left(\sum_{i=1}^{n} |y_i - z_i|^p\right)^{1/p}.$$

Hölder inequality

- The proof of the Minkowski inequality itself requires the Hölder inequality:
- For any nonnegative real numbers $a_1, \dots, a_n; b_1, \dots, b_n$, and any p, q > 1 with 1/p + 1/q = 1, we have:

$$\sum_{i=1}^{n} a_i b_i \le \left[\sum_{i=1}^{n} a_i^p \right]^{1/p} \left[\sum_{i=1}^{n} b_i^q \right]^{1/q}$$

- The proof of the Hölder inequality relies on the Young inequality:
- For any a, b > 0, we have

$$ab \leq \frac{a^p}{p} + \frac{b^q}{q},$$

with equality occurring iff: $a^p = b^q$.

- To prove the Young inequality, one can use the (strict) convexity of the exponential function.
- For any reals x, y, we have:

$$e^{\frac{x}{p} + \frac{y}{q}} \le \frac{e^x}{p} + \frac{e^y}{q}.$$

- We then set: $x = p \ln a$ and $y = q \ln b$ to get the Young inequality.
- A good reference on inequalities is: Z. Cvetkovski, Inequalities: theorems, techniques and selected problems, 2012, Springer Science & Business Media. # Cauchy-Schwartz inequality

• Note that the triangular inequality for the Minkowski distance implies:

$$\sum_{i=1}^{n} |x_i| \le \left[\sum_{i=1}^{n} |x_i|^p \right]^{1/p}.$$

• Note that for p=2, we have q=2. The Hölder inequality implies for that special case

$$\sum_{i=1}^{n} |x_i y_i| \le \sqrt{\sum_{i=1}^{n} x_i^2} \sqrt{\sum_{i=1}^{n} y_i^2}.$$

• Since the LHS od thes above inequality is greater then $|\sum_{i=1}^n x_i y_i|$, we get the Cauchy-Schwartz inequality

$$\left|\sum_{i=1}^{n} x_i y_i\right| \le \sqrt{\sum_{i=1}^{n} x_i^2} \sqrt{\sum_{i=1}^{n} y_i^2}.$$

* Using the dot product notation called also scalar product notation: $\mathbf{x} \cdot \mathbf{y} = \sum_{i=1}^{n} x_i y_i$, and the norm notation $\|\cdot\|_2\|$, the Cauchy-Schwart inequality is:

$$|\mathbf{x} \cdot \mathbf{y}| \le ||\mathbf{x}||_2 ||\mathbf{y}||_2.$$

Pearson correlation distance

• The Pearson correlation coefficient is a similarity measure on \mathbb{R}^n defined by:

$$\rho(\mathbf{x}, \mathbf{y}) = \frac{\sum_{i=1}^{n} (x_i - \bar{\mathbf{x}})(y_i - \bar{\mathbf{y}})}{\sqrt{\sum_{i=1}^{n} (x_i - \bar{\mathbf{x}})^2 \sum_{i=1}^{n} (y_i - \bar{\mathbf{y}})^2}},$$

where $\bar{\mathbf{x}}$ is the mean of the vector \mathbf{x} defined by:

$$\bar{\mathbf{x}} = \frac{1}{n} \sum_{i=1}^{n} x_i,$$

• Note that the Pearson correlation coefficient satisfies P2 and is invariant to any positive linear transformation, i.e.:

$$\rho(\alpha \mathbf{x}, \mathbf{y}) = \rho(\mathbf{x}, \mathbf{y}),$$

for any $\alpha > 0$.

• The Pearson distance (or correlation distance) is defined by:

$$d(\mathbf{x}, \mathbf{y}) = 1 - \rho(\mathbf{x}, \mathbf{y}).$$

• Note that the Pearson distance does not satisfy A1 since $d(\mathbf{x}, \mathbf{x}) = 0$ for any non-zero vector \mathbf{x} . It neither satisfies the triangle inequality. However, the symmetry property is fullfilled.

Cosine correlation distance

• The cosine of the angle θ between two vectors **x** and **y** is a measure of similarity given by:

$$\cos(\theta) = \frac{\mathbf{x} \cdot \mathbf{y}}{\|\mathbf{x}\|_2 \|\mathbf{y}\|_2} = \frac{\sum_{i=1}^n x_i y_i}{\sqrt{\sum_{i=1}^n x_i^2 \sum_{i=1}^n y_i^2}}.$$

- Note that the cosine of the angle between the two centred vectors $\mathbf{x} \bar{\mathbf{x}}\mathbf{1}$ and $\mathbf{y} \bar{\mathbf{y}}\mathbf{1}$ coincides with the Pearson correlation coefficient of \mathbf{x} and \mathbf{y} , where $\mathbf{1}$ is a vector of units of \mathbb{R}^n .
- The cosine correlation distance is defined by:

$$d(\mathbf{x}, \mathbf{y}) = 1 - \cos(\theta).$$

 It shares similar properties than the Pearson correlation distance. Likewise, Axioms A1 and A3 are not satisfied.

Spearman correlation distance

• To calculate the Spearman's rank-order correlation, we need to map seperately each of the vectors to ranked data values:

$$\mathbf{x} \to \operatorname{rank}(\mathbf{x}) = (x_1^r, \cdots, x_n^r).$$

- Here, x_i^r is the rank of x_i among the set of values of \mathbf{x} .
- We illustrate this transformation with a simple example:
- If $\mathbf{x} = (3, 1, 4, 15, 92)$, then the rank-order vector is rank(\mathbf{x}) = (2, 1, 3, 4, 5).

```
x=c(3, 1, 4, 15, 92)
rank(x)
```

[1] 2 1 3 4 5

- The Spearman's rank correlation of two numerical variables \mathbf{x} and \mathbf{y} is simply the Pearson correlation of the two corresponding rank-order variables rank(\mathbf{x}) and rank(\mathbf{y}), i.e. $\rho(\text{rank}(\mathbf{x}), \text{rank}(\mathbf{y}))$. This measure is is useful because it is more robust against outliers than the Pearson correlation.
- If all the n ranks are distinct, it can be computed using the following formula:

$$\rho(\operatorname{rank}(\mathbf{x}), \operatorname{rank}(\mathbf{y})) = 1 - \frac{6\sum_{i=1}^{n} d_i^2}{n(n^2 - 1)},$$

where $d_i = x_i^r - y_i^r$, $i = 1, \dots, n$.

• The spearman distance is then defined by:

$$d(\mathbf{x}, \mathbf{y}) = 1 - \rho(\text{rank}(\mathbf{x}), \text{rank}(\mathbf{y})).$$

- It can be shown that easaly that it is not a proper distance.
- If all the n ranks are distinct, we get:

$$d(\mathbf{x}, \mathbf{y}) = \frac{6\sum_{i=1}^{n} d_i^2}{n(n^2 - 1)}.$$

```
x=c(3, 1, 4, 15, 92)
rank(x)
```

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

```
y=c(30,2, 9, 20, 48)
rank(y)
```

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

```
d=rank(x)-rank(y)
d
```

```
cor(rank(x),rank(y))
## [1] 0.7
1-6*sum(d^2)/(5*(5^2-1))
## [1] 0.7
```

Kendall tau distance

- The Kendall rank correlation coefficient is calculated from the number of correspondances between the rankings of \mathbf{x} and the rankings of \mathbf{y} .
- The number of pairs of observations among n observations or values is:

$$\binom{n}{2} = \frac{n(n-1)}{2}.$$

• The pairs of observations (x_i, x_j) and (y_i, y_j) are said to be concordant if:

$$sign(x_j - x_i) = sign(y_j - y_i),$$

and to be discordant if:

$$sign(x_j - x_i) = -sign(y_j - y_i),$$

where $sign(\cdot)$ returns 1 for positive numbers and -1 negative numbers and 0 otherwise.

- If $x_i = x_j$ or $y_i = y_j$ (or both), there is a tie.
- The Kendall τ coefficient is defined by (neglecting ties):

$$\tau = \frac{1}{n(n-1)} \sum_{i=1}^{n} \sum_{j=1}^{n} \text{sign}(x_j - x_i) \text{sign}(y_j - y_i).$$

• Let n_c (resp. n_d) be the number of concordant (resp. discordant) pairs, we have

$$\tau = \frac{2(n_c - n_d)}{n(n-1)}.$$

• The Kendall tau distance is then:

$$d(\mathbf{x}, \mathbf{v}) = 1 - \tau.$$

• Remark: the triangular inequality may fail in cases where there are ties.

```
x=c(3, 1, 4, 15, 92)
y=c(30,2, 9, 20, 48)
tau=0
for (i in 1:5)
{
    tau=tau+sign(x -x[i])%*%sign(y -y[i])
}
tau=tau/(5*4)
tau

## [,1]
## [1,] 0.6
cor(x,y, method="kendall")
```

[1] 0.6

Variables standardization

- Variables are often standardized before measuring dissimilarities.
- Standardization converts the original variables into uniteless variables.
- A well known method is the z-score transformation:

$$\mathbf{x} \to (\frac{x_1 - \bar{\mathbf{x}}}{s_{\mathbf{x}}}, \cdots, \frac{x_n - \bar{\mathbf{x}}}{s_{\mathbf{x}}}),$$

where $s_{\mathbf{x}}$ is the sample standard deviation given by:

$$s_{\mathbf{x}} = \frac{1}{n-1} \sum_{i=1}^{n} (x_i - \bar{\mathbf{x}})^2.$$

- The transformed variable will have a mean of 0 and a variance of 1.
- The result obtained with Pearson correlation measures and standardized Euclidean distances are comparable.
- For other methods, see: Milligan, G. W., & Cooper, M. C. (1988). A study of standardization of variables in cluster analysis. *Journal of classification*, 5(2), 181-204.

```
x=c(3, 1, 4, 15, 92)
y=c(30,2,9,20,48)
(x-mean(x))/sd(x)
## [1] -0.5134116 -0.5647527 -0.4877410 -0.2053646 1.7712699
scale(x)
##
              [,1]
## [1,] -0.5134116
## [2,] -0.5647527
## [3,] -0.4877410
## [4,] -0.2053646
## [5,] 1.7712699
## attr(,"scaled:center")
## [1] 23
## attr(,"scaled:scale")
## [1] 38.9551
(y-mean(y))/sd(y)
## [1] 0.45263128 -1.09293895 -0.70654639 -0.09935809 1.44621214
scale(y)
##
               [,1]
## [1,] 0.45263128
## [2,] -1.09293895
## [3,] -0.70654639
## [4,] -0.09935809
## [5,] 1.44621214
## attr(,"scaled:center")
## [1] 21.8
## attr(,"scaled:scale")
## [1] 18.11629
```

Similarity measures for binary data

- A common simple situation occurs when all information is of the presence/absence of 2-level qualitative characters.
- We assume there are n measured characters.
- *The presence of the character or type is coded 1 and the absence by 0.
- We have have at our disposal two vectors.
- **x** is observed for first individal (or object).
- y is observed for a second individual.
- We can then calculate the following statistics:

$$a = \mathbf{x} \cdot \mathbf{y} = \sum_{i=1}^{n} x_i y_i.$$

$$b = \mathbf{x} \cdot (\mathbf{1} - \mathbf{y}) = \sum_{i=1}^{n} x_i (1 - y_i).$$

$$c = (\mathbf{1} - \mathbf{x}) \cdot \mathbf{y} = \sum_{i=1}^{n} (1 - x_i) y_i.$$

$$d = (\mathbf{1} - \mathbf{x}) \cdot (\mathbf{1} - \mathbf{y}) = \sum_{i=1}^{n} (1 - x_i) (1 - y_i).$$

- The counts of matches are a for (1,1) and d for (0,0);
- The counts of mismatches are b for (1,0) and c for (0,1).
- Obviously, a + b + c + d = n.
- This gives a very useful 2×2 association table.

		Second individual		
		1	0	\overline{Totals}
First individual	1	a	b	a + b
	0	c	d	c+d
Totals		a+c	b+d	n

- The data shows 8 people (individuals) and 10 binary variables:
- Sex, Married, Fair Hair, Blue Eyes, Wears Glasses, Round Face, Pessimist, Evening Type, Is an Only Child, Left-Handed.

• We are comparing the records for Ilan with Talia.

Table 9 Binary Variables for Eight People

Person	Sex (Male = 1, Female = 0)	Married (Yes = 1, No ≈ 0)	Fair Hair = 1, Dark Hair = 0	Blue Eyes = 1, Brown Eyes = 0	Wears Glasses (Yes = 1, No = 0)	Round Face = 1, Oval Face = 0	Pessimist = 1, Optimist = 0	Evening Type = 1, Morning Type = 0	Is an Only Child (Yes = 1, No = 0)	Left-Handed = 1, Right-Handed = 0
Ilan	1	0	1	1	0	0	1	0	0	0
Jacqueline	0	1	0	0	1	0	0	0	0	0
Kim	0	0	1	0	0	0	1	0	0	1
Lieve	0	1	0	0	0	0	0	1	1	0
Leon	1	1	0	0	1	1	0	1	1	0
Peter	1	1	0	0	1	0	1	1	0	0
Talia	0	0	0	1	0	1	0	0	0	0
Tina	0	0	0	1	0	1	0	0	0	0

Figure 1: Table from Kaufman, L., & Rousseeuw, P. J. (2009). Finding groups in data: an introduction to cluster analysis (Vol. 344). John Wiley & Sons

library(dplyr)

```
##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
## filter, lag
## The following objects are masked from 'package:base':
##
## intersect, setdiff, setequal, union
x=data["Ilan",]
y=data["Talia",]
knitr::kable(table(x, y)[2:1,2:1],"pipe")
```

	1	0
1	1	3
0	1	5

- Therefore: a = 1, b = 3, c = 1, d = 5.
- Note that interchanging Ilan and Talia would permute b and c while leaving a and d unchanged.
- A good similarity or dissimilarity coefficient must treat b and c symmetrically.
- A list of some of the similarity measures that have been suggested for binary data is shown below.
- a more extensive list can be found in Gower and Legendre (1986).

Measure	Formula
Matching coefficient	(a+d)/(a+b+c+d)
Jaccard coefficient	a/(a+b+c)
Rogers and Tanimoto (1960)	(a+d)/[a+2(b+c)+d]
Gower and Legendre (1986)	$(a+d)/[a+\frac{1}{2}(b+c)+d]$
Gower and Legendre (1986)	$a/[a+\frac{1}{2}(b+c)]$

- The reason for such a large number of possible measures has to do with the apparent uncertainty as to how to deal with the count of zero–zero matches (d in Table 3.2).
- In some cases, of course, zero–zero matches are completely equivalent to one–one matches, and therefore should be included in the calculated similarity measure.
- An example is gender, where there is no preference as to which of the two categories should be coded zero or one.
- But in other cases the inclusion or otherwise of d is more problematic; for example, when the zero category corresponds to the genuine absence of some property, such as wings in a study of insects.

Distance matrix computation

- We'll use a subset of the data USArrests
- We'll use only a by taking 15 random rows among the 50 rows in the data set.
- Next, we standardize the data using the function scale():

```
install.packages("FactoMineR")
```

Installing package into '/home/rstudio-user/R/x86_64-pc-linux-gnu-library/4.0'

```
## (as 'lib' is unspecified)
library("FactoMineR")
data("USArrests") # Loading
head(USArrests, 3) # Print the first 3 rows
          Murder Assault UrbanPop Rape
## Alabama 13.2
                     236
                              58 21.2
## Alaska 10.0
                     263
                              48 44.5
## Arizona 8.1
                     294
                              80 31.0
set.seed(123)
ss <- sample(1:50, 15) # Take 15 random rows
df <- USArrests[ss, ] # Subset the 15 rows</pre>
df.scaled <- scale(df) # Standardize the variables</pre>
```