



Signed
projections

V. Batagelj

Signed
two-mode
networks

Salton's cosine

Jaccard's
similarity

Remarks

References

Projections of signed two-mode networks

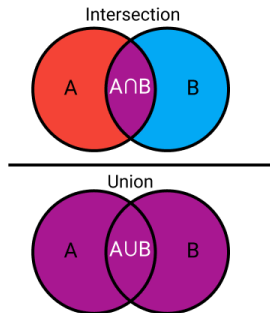
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IMFM Ljubljana, IAM UP Koper, and NRU HSE Moscow

1317 and 1318. sredin seminar
on Zoom, March 2 and 16, 2022

- 1 Signed two-mode networks
- 2 Salton's cosine
- 3 Jaccard's similarity
- 4 Remarks
- 5 References

Jaccard



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Current version of slides (March 16, 2022 at 17:47): [slides PDF](#)

<https://github.com/bavla/NormNet/tree/main/docs>



Signed two-mode networks

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A *signed two-mode network* is described by an affiliation matrix A with rows corresponding to *respondents*/actors U (primary mode) and columns corresponding to different *cases*/choices V (secondary mode).

The matrix entry $A[u, v]$ contains the decision/opinion of the respondent u about the case v and can take one of the values: p (Yes, Positive, Support), n (No, Negative, Oppose), a (Ambivalent, No opinion), r (Not valid), and z (NA, Not available, Absent) [3, 7].

For example, UN or parliament votings (U – countries / representatives; V – resolutions / bills); pre-election pools (U – respondents, V – selected politicians), etc.

The network matrix A depends on values p, n, a, r, z – we write $A(p, n, a, r, z)$. The notation $A(1, -1, 0, 0, 0)$ denotes a matrix obtained from A by substitutions $p \rightarrow 1, n \rightarrow -1, a \rightarrow 0, r \rightarrow 0, z \rightarrow 0$.



Signed two-mode networks in R

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```
> wdir <- ".../sreda/1317/SC"
> setwd(wdir)

> recode <- function(T,p,n,a,r,z){
+   A <- matrix(NA,nrow=nrow(T),ncol=ncol(T))
+   dimnames(A) <- list(rownames(T),colnames(T))
+   A[T=="p"] <- p; A[T=="n"] <- n; A[T=="a"] <- a;
+   A[T=="r"] <- r; A[T=="z"] <- z
+   return(A)
+ }

> T <- read.csv("SCa.txt",sep=";",head=TRUE,row.names=1,skip=3,
+   strip.white=TRUE,stringsAsFactors=FALSE)
> dim(T)
> L <- c("z","r","a","n","p")
> for(i in 1:9) T[,i] <- factor(T[,i],levels=L)
> head(T)

> A <- recode(T,1,2,3,4,5)
> head(A)
> A <- recode(T,1,-1,0,0,0)
> head(A)
```

1. US senate and congress

Bradley Robinson US Senate Voting Records for 1990-2016 / Sessions 101-114.

<https://data.world/bradrobinson/us-senate-voting-records>
Bavla

2. Bundestag

Data from <https://github.com/jmbh/bundestag> used by Jonas Haslbeck in [Analyzing voting pattern of German parliament](#).

Data for the time period 26.11.2014 - 14.04.2016 were obtained from: <https://www.bundestag.de/bundestag/plenum/abstimmung>
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3. Parlameter

Voting in Slovenian parliament <https://parlamer.si/>
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Let us have data on the survey collected in the matrix (table) A of dimension $n \times m$

$a[k, i]$ = respondent k answer to the i -th case.

All the cases are measured on the same numerical scale. We also allow the value NA (not measured).

The usual *scalar product* of the vectors x and y is determined by the rule $x \bullet y = \sum_i x_i \cdot y_i$. The *length* of the vector x is $|x| = \sqrt{x \bullet x}$.

Weighted scalar product $x \odot y = \sum_i w_i \cdot x_i \cdot y_i$ can be introduced for a given vector of nonnegative weights w . The *weighted length* of the vector x is $\|x\| = \sqrt{x \odot x}$.

Let A be the matrix of a two-mode network and $W = A \bullet \text{diag}(\sqrt{w}) = [\sqrt{w_i} \cdot a_{i,j}]$. Then $A^T \odot A = W^T \bullet W$ holds.



Salton's cosine

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Salton's index $S(i, j)$ or cosine similarity of cases i and j is defined as the cosine between the vectors / variables $A[:, i]$ and $A[:, j]$

$$S(i, j) = \cos(A[:, i], A[:, j]) = \frac{\sum_k a[k, i] \cdot a[k, j]}{\sqrt{\sum_k a[k, i]^2 \cdot \sum_k a[k, j]^2}} = \frac{A[:, i] \bullet A[:, j]}{|A[:, i]| \cdot |A[:, j]|}$$

In general, $S(i, j) \in [-1, 1]$ and $S(i, i) = 1$. $S(x, a \cdot x) = 1$ for $a > 0$.

The Salton's index can be transformed into dissimilarity (for clustering) in several ways:

$$d(i, j) = \frac{1 - S(i, j)}{2}$$

$$\delta(i, j) = \frac{\arccos(S(i, j))}{\pi}$$



Computing the Salton's similarity matrix

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From matrix A we calculate (projection to cases) a square *matching matrix* of cases (more precisely, answers to questions)

$$C = A^T \bullet A$$

of dimensions $m \times m$. Its entry

$$c[i, j] = \sum_k a^T[i, k] \cdot a[k, j] = \sum_k a[k, i] \cdot a[k, j]$$

is equal to the scalar product in the numerator of Salton's similarity.

More, $c[i, i] = \sum_k a[k, i] \cdot a[k, i] = \sum_k a[k, i]^2$ is the same as the term appearing in the denominator. Thus, Salton's similarity can be expressed in terms of the matching matrix

$$S(i, j) = \frac{c[i, j]}{\sqrt{c[i, i] \cdot c[j, j]}}$$



Salton's similarity for (p, n, a, r, z) -vectors

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Let's look at a special case where only three values are possible: p , n , and $a = r = z$. The data is recoded as follows: $p \rightarrow 1$, $n \rightarrow -1$, $z \rightarrow 0$. Let $A_s = A(1, -1, 0, 0, 0)$ contains the recoded data. Then from the form for $c[i, j]$ it follows

$$\begin{aligned} c[i, j] &= \#(1, 1 | -1, -1) - \#(1, -1 | -1, 1) = \\ &= \# \text{consistent answers} - \# \text{opposite answers} \end{aligned}$$

and

$$c[i, i] = \# \text{answers to question } i$$

Thus, $S(i, j)$ is equal to the consistency of the answers divided (normalized) by the geometric mean of the numbers of answers to both questions.

We can relatively easily show that $S(i, j) = 1 \Leftrightarrow a[., i] = a[., j]$ and $S(i, j) = -1 \Leftrightarrow a[., i] = -a[., j]$.



US Senate 106/1 dendrogram

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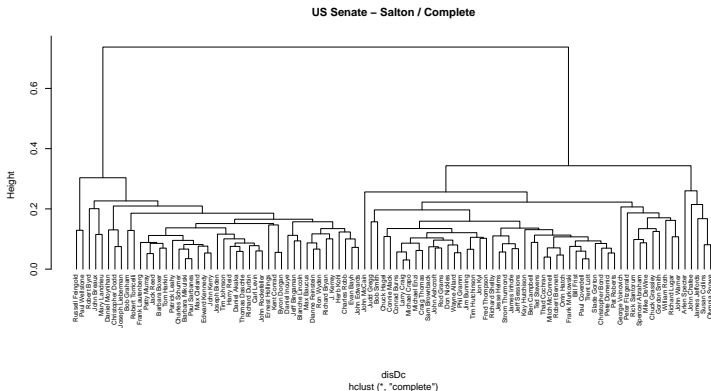
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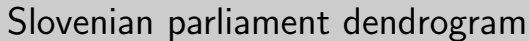
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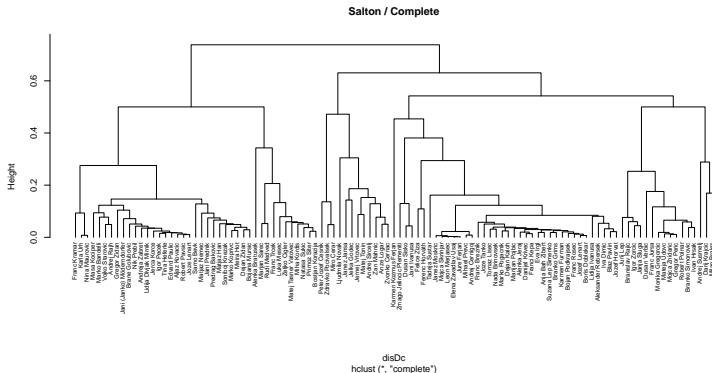
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Slovenian parliament 2 most similar neighbors network

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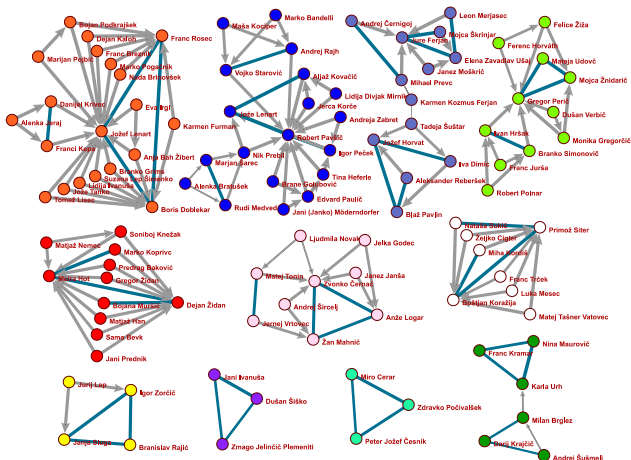
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3/12/22, 8:12 PM

9. Bidirected Arcs to Edges (MAX) of N8 (107)



file:///C:/Users/vlado/DL/data/parliament/parliament.svg



1/1



Jaccard's similarity

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Jaccard's similarity can be generalized to the scale (p, n, z) as follows:
By recoding, we create the matrices $A_p = A(1, 0, 0, 0, 0)$ and $A_n = A(0, 1, 0, 0, 0)$ and calculate the matching matrix

$$C = A_p^T \bullet A_p + A_n^T \bullet A_n$$

with entries

$$c[i, j] = \#(1, 1 | -1, -1) = \# \text{consistent answers}$$

and $c[i, i] = \# \text{answers to question } i$.

Therefore Jaccard's similarity ($J = \frac{|X \cap Y|}{|X \cup Y|}$)

$$J(i, j) = \frac{c[i, j]}{c[i, i] + c[j, j] - c[i, j]}$$

and the Jaccard dissimilarity $d_J(i, j) = 1 - J(i, j)$



Jaccard's similarity

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There are some generalizations of the Jaccard similarity $J(x, y)$ to (nonnegative) real vectors. Perhaps the most famous is (Ružička)

$$J_1(x, y) = \frac{\sum_i \min(x_i, y_i)}{\sum_i \max(x_i, y_i)}$$

We can easily check that for binary vectors x and y hold $\sum_i \min(x_i, y_i) = |X \cap Y|$ and $\sum_i \max(x_i, y_i) = |X \cup Y|$. So, for binary vectors we $J_1(x, y) = J(x, y)$.

Unfortunately, this measure does not work well for vectors containing negative values.

Another generalization of Jaccard's similarity is

$$J_2(x, y) = \frac{x \bullet y}{(x - y)^2 + x \bullet y}, \text{ for } x \neq 0 \text{ and } y \neq 0$$

and $J_2(0, 0) = 1$.

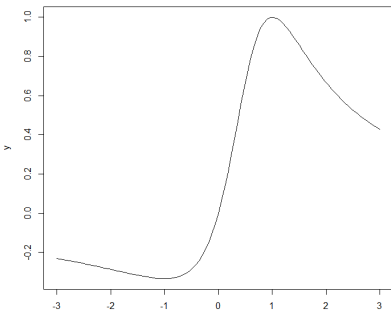
Again, for binary vectors x and y we have $x \bullet y = |X \cap Y|$ and $(x - y)^2 + x \bullet y = |X \cup Y|$.

Jaccard's similarity

The following properties hold for general real vectors:

- $J_2(x, y) = J_2(y, x)$
- $J_2(x, y) \leq 1$
- $J_2(x, y) = 1 \Leftrightarrow x = y$
- $J_2(a \cdot x, a \cdot y) = J_2(x, y)$ for $a \neq 0$.

For nonnegative vectors x and y , $J_2(x, y) \geq 0$. But in the case of $y = -x$ we get $J_2(x, -x) = -\frac{1}{3}$. If we look a little more generally, we get $J_2(x, a \cdot x) = \frac{a}{(1-a)^2 + a}$.





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$-\frac{1}{3}$ seems to be the smallest possible value of measure J_2 .

That turns out to be true. The proof is short: the inequality $0 \leq (x + y)^2$ is transformed into $-3 \cdot x \bullet y \leq (x - y)^2 + x \bullet y$ which means $J_2(x, y) \geq -\frac{1}{3}$; the equality holds iff $x = -y$.

Measure J_2 is mapped to the interval $[1, 0]$ by transformation

$$J_3(x, y) = \frac{3 \cdot J_2(x, y) + 1}{4}.$$

If we express (simplify) J_3 with scalar products, we get

$$J_3(x, y) = \frac{\left(\frac{x+y}{2}\right)^2}{(x - y)^2 + x \bullet y}$$



Generalized binary similarities

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Most of similarities between binary vectors are defined/expressed using matching counters $a = \#(1, 1)$, $b = \#(1, 0)$, $c = \#(0, 1)$ and $d = \#(0, 0)$. This approach can be extended also to signed data. To obtain the counters used in a selected similarity we can use the approach we used for extending the Jaccard's similarity.

For example, to get counters $pn = \#(p, n)$ we create matrices $A_p = A(1, 0, 0, 0, 0)$ and $A_n = A(0, 1, 0, 0, 0)$ and compute their product

$$C_{pn} = A_p^T \bullet A_n$$

with entries

$$c_{pn}[i, j] = \#(p, n) = \#\text{consistent answers}$$

and $c_{pn}[i, i] = 0$. Note that, since $C_{np} = C_{pn}^T$

$$c_{pn}[i, j] + c_{pn}[j, i] = \#(p, n | n, p) = \#\text{opposite answers}$$



Manhattan-Jaccard's dissimilarity

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I also tried the dissimilarity that uses the Jaccard's normalization on the Manhattan distance

$$MJ(x, y) = \frac{\sum_i w_i \cdot |x_i - y_i|}{|X \cup Y|}$$

where $|X \cup Y| = \sum_i \max(|x_i|, |y_i|)$.



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- 1 Generalization of some other (dis)similarities for binary vectors to (p,n,a,r,z) -vectors [5].
- 2 US senate, Bundestag, Parlameter. Additional data: parties, short names of cases, keywords/tags, types of cases: **poslanci**, **besede**, **ki so zaznamovale sejo**
- 3 Schoch [7], complex numbers.
- 4 semirings [3].

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