

Social relations models for binary, ranked and ordinal relations

567 Statistical analysis of social networks

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Adapting the normal SRM

Relational data are often valued, non-binary.

The standard SRM is appropriate for **normal** valued relational data.

However, relational data is often neither binary nor normal:

- Links can be **absent** or **continuous** within the same network.
 - meaning $Y_{i,j}$ is either 0 or some arbitrary real number;
 - communication networks (time spent communicating);
- Links can be **absent** or **ordinal** within the same network.
 - conflict networks (negative, positive or zero relation);
 - ranked nominations (friends are ranked, non-friends are not).

The SRM can be adapted via **ordinal probit models** to handle such data.

Ordinal data

An **ordinal variable** has a meaningful ordering to the possible outcomes. This is in contrast to a **categorical** (non-ordered) variable.

Ordinal variable

- continuous: (all real numbers)
- discrete: (counts, ranks, etc.)

Categorical variable

- non-orderable categories (religion, ethnicity, etc.)

Binary variable as ordinal

The simplest ordinal variable is a binary random variable $Y \in \{0, 1\}$. Let

$$\Pr(Y = 1) = \theta$$

$$\Pr(Y = 0) = 1 - \theta$$

This model for Y has the following **latent variable representation**:

$$\mu = \Phi^{-1}(\theta)$$

$$Z \sim N(\mu, 1)$$

$$Y = 1 \times (Z > 0)$$

Here, $\Phi^{-1}(\theta)$ is the θ -quantile of the standard normal distribution.

Probit representation

To confirm the representation, recall

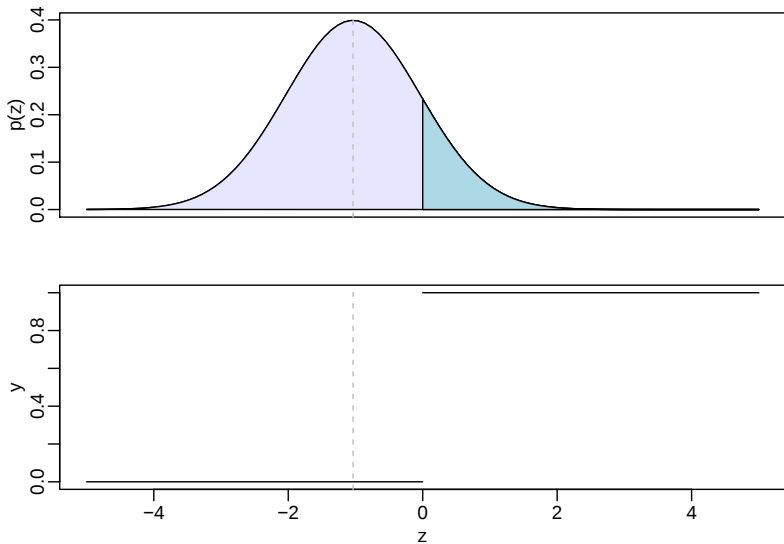
- If $Z \sim N(\mu, 1)$ then $Z - \mu \sim N(0, 1)$;
- If $Z \sim N(\mu, 1)$ then $Z = \mu + \epsilon$, where $\epsilon \sim N(0, 1)$;
- If $\epsilon \sim N(0, 1)$ then $-\epsilon \sim N(0, 1)$.

Now do the calculation:

$$\begin{aligned}\Pr(Y = 1) &= \Pr(Z > 0) \\ &= \Pr(-Z < 0) \\ &= \Pr([-Z + \mu] < \mu) \\ &= \Pr(\epsilon < \mu) = \Phi(\mu) = \theta\end{aligned}$$

Probit representation

$$\theta = .15, \mu = \Phi^{-1}(.15) = -1.04$$



Probit regression

Now suppose we have

binary data $Y_1, \dots, Y_n$

we want to relate to

explanatory variables $\mathbf{x}_1, \dots, \mathbf{x}_n$

Latent variable model:

$$\epsilon_1, \dots, \epsilon_n \sim \text{i.i.d. } N(0, 1)$$

$$Z_i = \beta^T \mathbf{x}_i + \epsilon_i$$

$$Y_i = 1 \times (Z_i > 0)$$

Under this latent variable model, the Y_i 's are independent and

$$\begin{aligned} \Pr(Y_i = 1) &= \Pr(Z_i > 0) \\ &= \Pr(\beta^T \mathbf{x}_i + \epsilon_i > 0) \\ &= \Pr(-\epsilon_i < \beta^T \mathbf{x}_i) \\ &= \Pr(\epsilon_i < \beta^T \mathbf{x}_i) = \Phi(\beta^T \mathbf{x}_i) \end{aligned}$$

Probit regression

This latent variable model is exactly the same as **probit regression**:

$$\Pr(Y_1 = y_1, \dots, Y_n = y_n | \mathbf{x}_1, \dots, \mathbf{x}_n) = \prod_{i=1}^n \Phi(\beta^T \mathbf{x}_i)^{y_i} [1 - \Phi(\beta^T \mathbf{x}_i)]^{1-y_i}$$

Compare to **logistic regression**:

$$\Pr(Y_1 = y_1, \dots, Y_n = y_n | \mathbf{x}_1, \dots, \mathbf{x}_n) = \prod_{i=1}^n \left(\frac{e^{\beta^T \mathbf{x}_i}}{1 + e^{\beta^T \mathbf{x}_i}} \right)^{y_i} \left(\frac{1}{1 + e^{\beta^T \mathbf{x}_i}} \right)^{1-y_i}$$

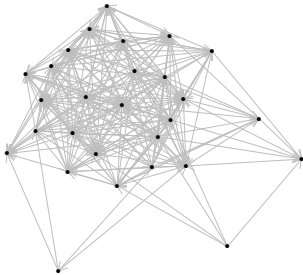
In fact, logistic regression has a latent variable representation also:

$$\epsilon_1, \dots, \epsilon_n \sim \text{i.i.d. } L(0, 1)$$

$$Z_i = \beta^T \mathbf{x}_i + \epsilon_i$$

$$Y_i = 1 \times (Z_i > 0)$$

Logistic versus probit regression



Sheep dominance data

- counts of dominance encounters between 28 male sheep;
- age of each sheep, in years.

Let's examine the effect of age difference on dominance.

Sheep dominance data

Let

- $\text{dom}[i, j]$ = indicator that i has dominated j at least once;
- $\text{aged}[i, j] = \text{age}_i - \text{age}_j$.

```
mean(aged[dom==1], na.rm=TRUE )
```

```
## [1] 2.184
```

```
mean(aged[dom==0], na.rm=TRUE )
```

```
## [1] -1.079051
```

```
summary( glm(dom~aged,family=binomial) )$coef
```

##	Estimate	Std. Error	z value	Pr(> z)
## (Intercept)	-0.8357694	0.08729001	-9.574628	1.022245e-21
## aged	0.2260270	0.02317097	9.754752	1.760303e-22

```
summary( glm(dom~aged,family=binomial(link=probit)) )$coef
```

##	Estimate	Std. Error	z value	Pr(> z)
## (Intercept)	-0.5138332	0.05111585	-10.05232	8.972511e-24
## aged	0.1385751	0.01337034	10.36437	3.601223e-25

Logit, probit and other binary regression models

Comparing logistic and probit regression output:

- $\hat{\beta}$'s are on different scales;
- inference (z-scores) are typically similar.

Both models are **binary regression models** of the form:

$$\Pr(Y_1 = y_1, \dots, Y_n = y_n | \mathbf{x}_1, \dots, \mathbf{x}_n) = \prod_{i=1}^n g(\beta^T \mathbf{x}_i)^{y_i} [1 - g(\beta^T \mathbf{x}_i)]^{1-y_i},$$

where g^{-1} is called the **inverse-link function**.

Link functions:

- **logistic regression:** $g^{-1}(\mu) = \exp(\mu) / [1 + \exp(\mu)]$;
- **probit regression:** $g^{-1}(\mu) = \Phi(\mu)$;
- **other binary regression:** $g^{-1}(\mu)$ is a strictly increasing function.

SRM for binary relational data

Recall the latent variable representation of probit regression:

$$\begin{aligned}Z_{i,j} &= \beta^T \mathbf{x}_{i,j} + \epsilon_{i,j} \\ Y_{i,j} &= 1 \times (Z_{i,j} > 0)\end{aligned}$$

We could estimate β with a probit regression analysis, but what about network dependence in the data?

Could there be

- across-row heterogeneity/within row correlation?
- across-column heterogeneity/within column correlation?
- within dyad correlation?

SRM for binary relational data

Recall the latent variable representation of probit regression:

$$\begin{aligned}Z_{i,j} &= \beta^T \mathbf{x}_{i,j} + \epsilon_{i,j} \\ Y_{i,j} &= 1 \times (Z_{i,j} > 0)\end{aligned}$$

Dependence on the Y -scale can be induced by dependence on the Z -scale:

$$\begin{aligned}\epsilon_{i,j} &= a_i + b_j + e_{i,j} \\ \{(a_1, b_1), \dots, (a_n, b_n)\} &\sim \text{i.i.d. } N(0, \Sigma_{ab}) \\ \{(e_{i,j}, e_{j,i}) : i \neq j\} &\sim \text{i.i.d. } N(0, \Sigma_e)\end{aligned}$$

$$\Sigma_{ab} = \begin{pmatrix} \sigma_a^2 & \sigma_{ab} \\ \sigma_{ab} & \sigma_b^2 \end{pmatrix} \quad \Sigma_e = \begin{pmatrix} 1 & \rho \\ \rho & 1 \end{pmatrix}$$

Note that the variance of $e_{i,j}$ is fixed at 1.

The scales of $e_{i,j}$ and β are not separately identifiable.

SRM for binary relational data

SRM probit model:

$$Y_{i,j} = 1 \times (Z_{i,j} > 0)$$

$$Z_{i,j} = \beta^T \mathbf{x}_{i,j} + \epsilon_{i,j}$$

$$\epsilon_{i,j} = a_i + b_j + e_{i,j},$$

and $\{a_i, b_j, e_{i,j}\}$ are random effects as described previously.

Parameter estimation:

- The likelihood can't be expressed in closed form;
- Bayesian parameter estimates can be obtained via MCMC.

The latter is provided in the package `amen`.

Binary SRM in amen

ame_bin

package:amen

R Documentation

AME fit for binary relational data

Description:

An MCMC routine providing a fit to an additive and multiplicative effects (AME) regression model for binary relational data

Usage:

```
ame_bin(Y, X, rvar = TRUE, cvar = TRUE, dcor = TRUE, R = 0, seed = 1, nscan = 50000, I
```

Arguments:

Y: an $n \times n$ square relational matrix

X: an $n \times n \times p$ array of covariates

rvar: logical: fit row random effects?

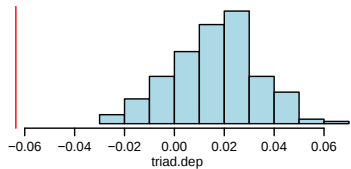
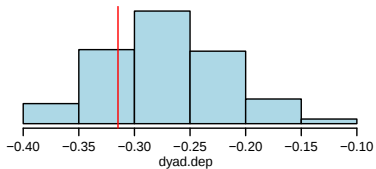
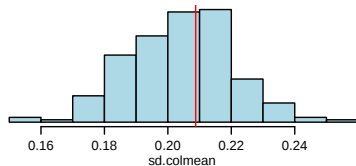
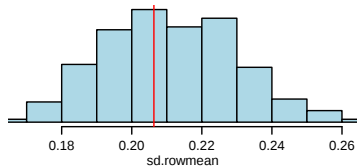
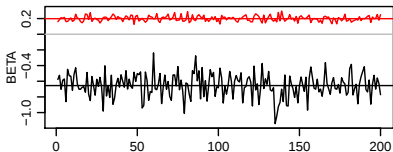
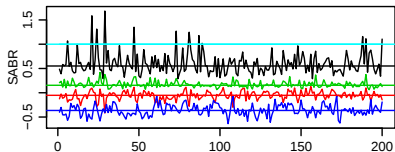
cvar: logical: fit column random effects?

dcor: logical: fit a dyadic correlation?

SRM probit in amen

```
XD<-outer(X,X,"-")
```

```
fit_ame_bin<-ame(YB,XD,model="bin")
```



Posterior analysis

The `summary` command provides a canned summary of the fitted model:

```
summary(fit_ame_bin)

##
## beta:
##           pmean   psd z-stat p-val
## intercept -0.654 0.149 -4.374    0
## .dyad      0.200 0.034  5.835    0
##
## Sigma_ab pmean:
##           a      b
## a  0.603 -0.057
## b -0.057  0.172
##
## rho pmean:
## -0.359
```

Posterior analysis

Confidence intervals can be obtained via the quantile command:

```
apply( fit_ame_bin$BETA , 2 , quantile, prob=c(.025,.5, .975))
```

```
##      intercept      .dyad
## 2.5% -0.9423416 0.1398794
## 50%  -0.6554213 0.1985873
## 97.5% -0.4028070 0.2637526
```

```
apply( fit_ame_bin$SABR , 2 , quantile, prob=c(.025,.5, .975))
```

```
##           va           cab           vb           rho ve
## 2.5%  0.3111828 -0.26359446 0.07500074 -0.5818362  1
## 50%   0.5511227 -0.05253038 0.15423887 -0.3646930  1
## 97.5% 1.2468605  0.09875416 0.33123579 -0.1495458  1
```

Posterior inference

Summary of results:

Strong evidence of an age effect:

- Older sheep dominate younger ones.

Row and column heterogeneity is not substantial:

- Compare $\hat{\sigma}_a^2 = 0.55$ and $\hat{\sigma}_b^2 = 0.15$ to the error variance $\sigma_e^2 = 1$.

There is evidence that dominance tends to go one-way in a dyad:

- The 95% CI for ρ is $(-0.58, -0.15)$.

Goodness of fit

The default `amen` plot provides four goodness of fit plots:

Outdegree distribution: Comparing outdegree distribution to simulated;

Indegree distribution: Comparing indegree distribution to simulated;

Reciprocity: Comparing fraction reciprocated ties to simulated fraction;

Transitivity: Comparing number of triangles to simulated number.

These statistics are computed with the commands

```
t_degree
```

```
t_recip
```

```
t_trans
```

Model comparison

Let's use these statistics to evaluate the fit of three models:

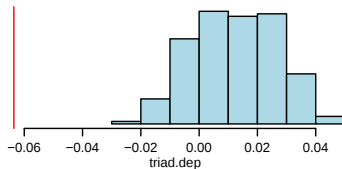
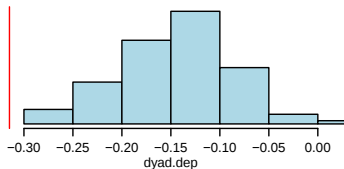
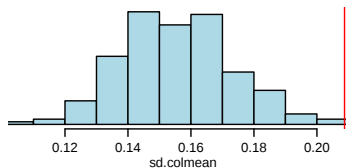
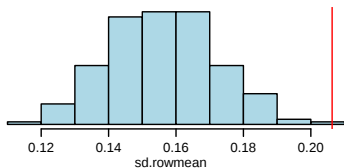
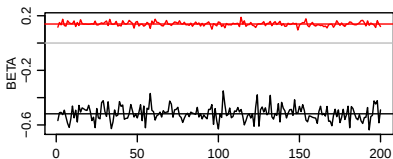
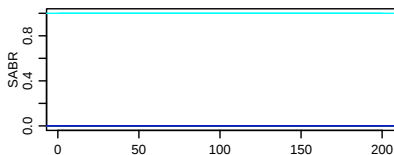
fit_bin_000: the probit model, fix $\sigma_a^2 = \sigma_b^2 = \rho = 0$.

fit_bin_001: dyadic correlation ρ estimated, $\sigma_a^2 = \sigma_b^2 = 0$.

fit_bin_111: full SRM covariance model.

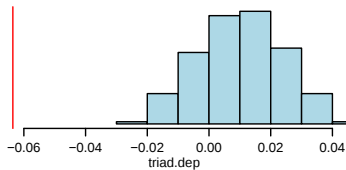
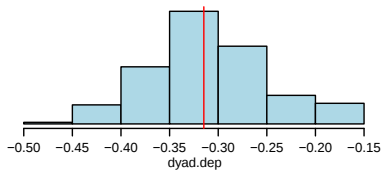
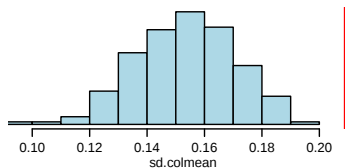
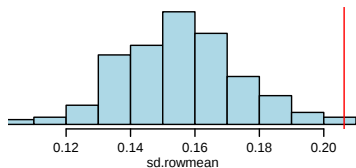
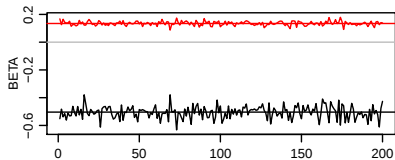
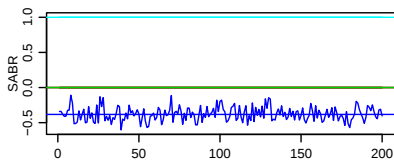
Model comparison

```
fit_bin_000<-ame(YB,XD,model="bin",rvar=FALSE,cvar=FALSE,dcor=FALSE)
```



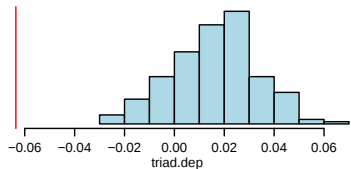
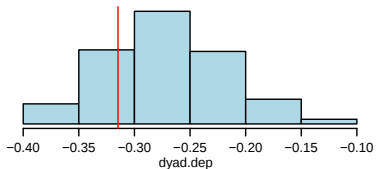
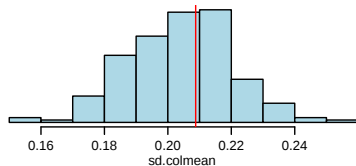
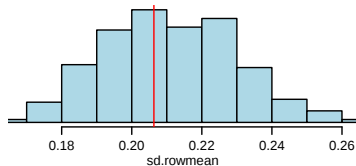
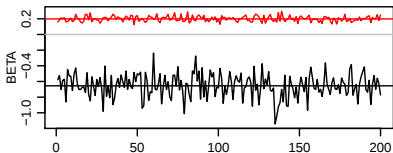
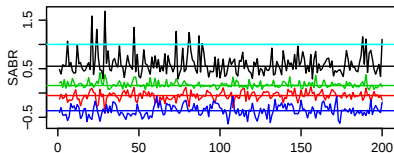
Model comparison

```
fit_bin_001<-ame(YB,XD,model="bin",rvar=FALSE,cvar=FALSE,dcor=TRUE)
```



Model comparison

```
fit_bin_111<-ame(YB,XD,model="bin")
```



Model comparison

The SRM model `fit_bin_111` looks best based on these statistics, except possibly for the transitivity statistic (more on that soon).

Ordered probit models for ordinal data

The original sheep data consists of counts:

```
YV[1:8,1:8]
```

```
##      V1 V2 V3 V4 V5 V6 V7 V8
## [1,] NA  0  0  0  0  0  0  1
## [2,]  0 NA  0  0  5  2  1  0
## [3,]  0  0 NA  0  7  4  0  0
## [4,]  0  0  8 NA  0  0  0  0
## [5,]  0  0  0  0 NA  1  0  0
## [6,]  0  0  0  7  0 NA  0  0
## [7,]  0  0  1  0  0  0 NA  0
## [8,]  0  1  1  4  5  3  0 NA
```

Ideally, this information should be taken into consideration.

- In principle, throwing away information is not efficient.
- Effects of some covariates might distinguish high values of the relation.

Ordered probit models for ordinal data

Latent variable model:

$$\begin{aligned}Z_{i,j} &= \beta^T x_{i,j} + \epsilon_{i,j} \\ \epsilon_{i,j} &= a_i + b_j + e_{i,j}\end{aligned}$$

Binary probit:

$$Y_{i,j} = \begin{cases} 0 & \text{if } Z_{i,j} < 0 \\ 1 & \text{if } Z_{i,j} > 0 \end{cases}$$

What if $Y_{i,j} \in \{y_1, \dots, y_K\} = \mathcal{Y}$?

Here, $\mathcal{Y}$ is an ordered, countable set of possible values of $Y_{i,j}$.

Ordered probit models for ordinal data

Latent variable model:

$$\begin{aligned}Z_{i,j} &= \beta^T \mathbf{x}_{i,j} + \epsilon_{i,j} \\ \epsilon_{i,j} &= a_i + b_j + e_{i,j}\end{aligned}$$

Ordered probit:

$$Y_{i,j} = \begin{cases} y_1 & \text{if } Z_{i,j} \in (-\infty, c_1) \\ y_2 & \text{if } Z_{i,j} \in (c_1, c_2) \\ \vdots & \\ y_{K-1} & \text{if } Z_{i,j} \in (c_{K-2}, c_{K-1}) \\ y_K & \text{if } Z_{i,j} \in (c_{K-1}, \infty) \end{cases}$$

Ordered probit models for ordinal data

```
table(c(YB))
```

```
##
```

```
##    0    1
```

```
## 506 250
```

```
table(c(YV))
```

```
##
```

```
##    0    1    2    3    4    5    6    7    8    9   10   11   12
```

```
## 506 100  59  34  20  13   5   7   5   1   3   1   2
```

```
round( table(c(YV))/sum(table(c(YV))) ,3 )
```

```
##
```

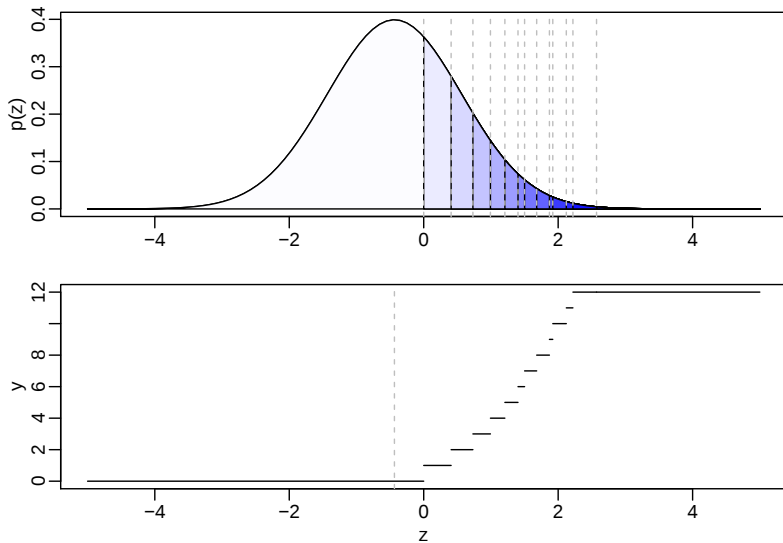
```
##      0      1      2      3      4      5      6      7      8      9     10     11
```

```
## 0.669 0.132 0.078 0.045 0.026 0.017 0.007 0.009 0.007 0.001 0.004 0.001
```

```
##      12
```

```
## 0.003
```

Link function for ordered probit



Fitting ordered probit models in amen

ame_ord

package:amen

R Documentation

AME fit for ordinal relational data

Description:

An MCMC routine providing a fit to an additive and multiplicative effects (AME) regression model for ordinal relational data

Usage:

```
ame_ord(Y, X, rvar = TRUE, cvar = TRUE, dcor = TRUE, R = 0, seed = 1, nscan = 50000, burn = 500, odens = 25, plot = TRUE, print = TRUE)
```

Arguments:

Y: an $n \times n$ square relational matrix

X: an $n \times n \times p$ array of covariates

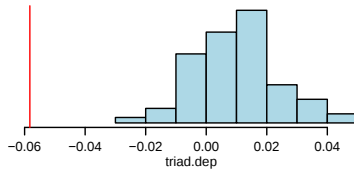
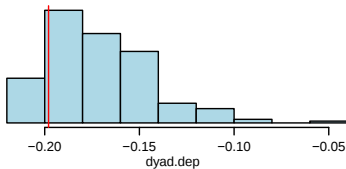
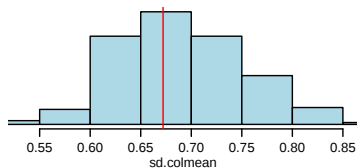
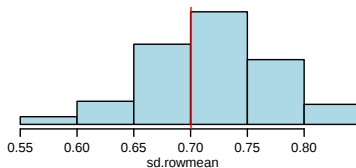
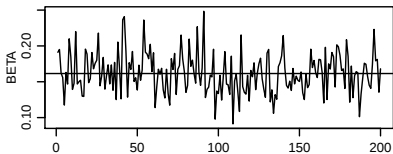
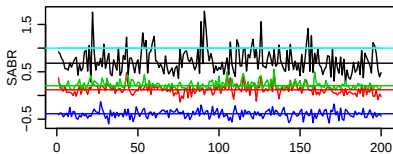
rvar: logical: fit row random effects?

cvar: logical: fit column random effects?

dcor: logical: fit a dyadic correlation?

SRM probit in amen

```
fit_ame_ord<-ame(YV,XD,model="ord")
```



Posterior analysis

Summary:

```
summary(fit_ame_ord)

##
## beta:
##      pmean   psd z-stat p-val
## .dyad 0.162 0.028  5.72    0
##
## Sigma_ab pmean:
##      a      b
## a 0.731 0.125
## b 0.125 0.224
##
## rho pmean:
## -0.388
```

Posterior analysis

Confidence intervals

```
apply( fit_ame_ord$BETA , 2 , quantile, prob=c(.025,.5, .975))
```

```
##           .dyad
## 2.5%  0.1138031
## 50%   0.1615003
## 97.5% 0.2231967
```

```
apply( fit_ame_ord$SABR , 2 , quantile, prob=c(.025,.5, .975))
```

```
##           va           cab           vb           rho ve
## 2.5%  0.3728079 -0.06367146  0.1060518 -0.5601718  1
## 50%   0.6811580  0.12026448  0.2057822 -0.3863772  1
## 97.5% 1.4126096  0.37453941  0.4239469 -0.2343322  1
```

These results are very similar to those obtained from the binary probit analysis using the dichotomized data.

The role of covariance

Regression modeling:

$$Y_{i,j} = \beta^T \mathbf{x}_{i,j} + \epsilon_{ij}$$
$$\mathbf{Y} = \langle \mathbf{X}, \beta \rangle + \mathbf{E}$$

OLS estimation:

$$\hat{\beta} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$$

Precision of OLS estimators:

Let $\mathbf{C} = \text{Cov}[\mathbf{E}]$

$$\begin{aligned}\text{Cov}[\hat{\beta}] &= (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{C} \mathbf{X} (\mathbf{X}^T \mathbf{X})^{-1} \\ &= (\mathbf{X}^T \mathbf{X})^{-1} \sigma^2 \text{ if } \mathbf{C} = \sigma^2 \mathbf{I}\end{aligned}$$

For networks and relational data, typically $\mathbf{C} \neq \sigma^2 \mathbf{I}$.

Accurate standard errors can't be obtained unless we know/estimate $\text{Cov}[\mathbf{E}]$.

Justifying the SRM

The social relations covariance model can be by a symmetry principle.

Exchangeability:

1. randomly sample n individuals from a population;
2. observe $\epsilon_{i,j}$ = directed relation between i th and j th person.

Consider a probability model $\Pr(\mathbf{E})$ for the possible outcomes of $\mathbf{E} = \{\epsilon_{i,j}\}$

$$\Pr \left(\begin{pmatrix} NA & -0.94 & 0.15 \\ -0.7 & NA & 0.63 \\ 0.63 & -0.42 & NA \end{pmatrix} \right) = ?$$

$$\Pr \left(\begin{pmatrix} NA & 0.63 & -0.42 \\ 0.15 & NA & -0.94 \\ 0.63 & -0.7 & NA \end{pmatrix} \right) = ?$$

Note the second matrix is the same as the first with (1,2,3) relabelled as (2,3,1).

Justifying the SRM

Let π be some permutation of $\{1, \dots, n\}$.

$$\mathbf{E} = \{\epsilon_{i,j} : i \neq j\}$$
$$\mathbf{E}_\pi = \{\epsilon_{\pi_i, \pi_j} : i \neq j\}$$

Exchangeability: A probability distribution $\Pr(\mathbf{E})$ is **exchangeable** if

$$\Pr(\mathbf{E}) = \Pr(\mathbf{E}_\pi)$$

for all $\mathbf{E}$ and permutations π .

Exchangeability can be justified by

- random sampling of nodes from a population;
- symmetry in the uncertainty in the $\epsilon_{i,j}$'s.

Justifying the SRM

Interesting result:

Suppose

1. $\epsilon_{i,j}$'s are normal and mean zero;
2. our probability for $\mathbf{E} = \{\epsilon_{i,j}\}$ is exchangeable.

Then

$$\begin{aligned}\epsilon_{i,j} &= a_i + b_j + e_{i,j} \\ \{(a_1, b_1), \dots, (a_n, b_n)\} &\sim \text{i.i.d. } N(0, \Sigma_{ab}) \\ \{(e_{i,j}, e_{j,i}) : i \neq j\} &\sim \text{i.i.d. } N(0, \Sigma_e)\end{aligned}$$

for some Σ_{ab} and Σ_e (Li and Loken, 2002).

Interpretation:

normality + exchangeability $\Rightarrow$ SRM

Justifying the SRM

There is a stronger result than this: If $\mathbf{E}$ is exchangeable, then

$$\epsilon_{i,j} = a_i + b_j + e_{i,j}$$

$$\text{Cov}[(a_i, b_i)] = \Sigma_{ab}$$

$$\text{Cov}[(e_{i,j}, e_{j,i})] = \Sigma_e$$

for some Σ_{ab}, Σ_e .

Interpretation:

exchangeability $\Rightarrow$ SRM covariance structure

Implication:

$$\text{Cov}[\hat{\beta}] = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{C} \mathbf{X} (\mathbf{X}^T \mathbf{X})^{-1},$$

where $\mathbf{C} = \text{Cov}[(\epsilon_{i,j}, \epsilon_{j,i})]$.

Under exchangeability, the SRM provides appropriate SEs and CIs for β .