

Bilinear random effects models for matrix data

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Duke STA 610

Matrix-valued data

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- $y_{i,j}$ = measurement specific to i th row unit, j th column unit.

Examples:

- Classical multivariate data: $y_{i,j}$ = j th variable for person i .
- Panel data: $y_{i,j}$ = outcome for unit i at time j (mv time series);
- Rating data: $y_{i,j}$ = rating of object i by person j .
- Dyadic data: $y_{i,j}$ = interaction between units i and j .

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Indices as grouping factors

In some situations, the row and column indices may be thought of as grouping factors.

- If the levels of an index set represent very different things (height, weight, income, education), then probably don't take this perspective.
- If they represent members of a common set of objects, then this perspective can be useful.
- a collection of survey participants, a collection of time points;
- a collection of movie raters, a collection of movies;
- a collection of people in a social network.

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Additive random effects models

Basic additive random effects model:

$$\begin{aligned}y_{i,j} &= \beta^\top x_{i,j} + a_i + b_j + \epsilon_{i,j} \\ a_1, \dots, a_m &\sim \text{i.i.d } N(0, \tau_a^2) \\ b_1, \dots, b_n &\sim \text{i.i.d } N(0, \tau_b^2).\end{aligned}$$

In a homework, we showed that this implies

$$\begin{aligned}\text{Cov}[y_{i,j}, y_{i',j'}] &= \tau_a^2 \text{ if } i = i' \\ \text{Cov}[y_{i,j}, y_{i',j'}] &= \tau_b^2 \text{ if } j = j' \\ \text{Cov}[y_{i,j}, y_{i',j'}] &= 0 \text{ if } i \neq i', j \neq j' .\end{aligned}$$

For example, for panel data this implies exchangeability among all observations within a common time point (they are all equally correlated).

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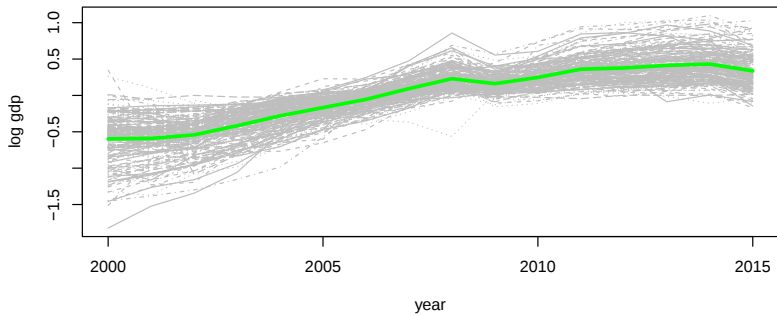
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Example: Per-capita GDP



Simple additive fit

```
xrow<-rep( rownames(Y) , times=ncol(Y))  
xcol<-rep( colnames(Y) , times=rep(nrow(Y),ncol(Y)) )  
y<-c(Y)
```

```
cbind( y,xrow,xcol)[1:10,]
```

##	y	xrow	xcol
## [1,]	"7.06969467420624"	"ALB"	"2000"
## [2,]	"7.47136990344077"	"DZA"	"2000"
## [3,]	"6.40732921228798"	"AGO"	"2000"
## [4,]	"9.21977166366161"	"ATG"	"2000"
## [5,]	"8.94497719738542"	"ARG"	"2000"
## [6,]	"6.43201494839607"	"ARM"	"2000"
## [7,]	"9.98345866321392"	"AUS"	"2000"
## [8,]	"10.1071329419972"	"AUT"	"2000"
## [9,]	"6.48478397663778"	"AZE"	"2000"
## [10,]	"9.96369634168077"	"BHS"	"2000"

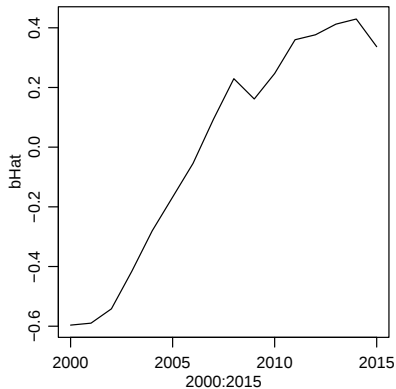
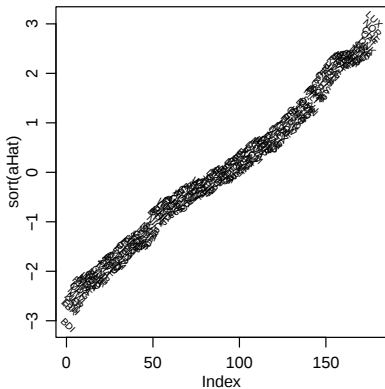
Simple additive fit

```
library(lme4)
fit<-lmer( y ~ 1 + (1|xrow) + (1|xcol) )
summary(fit)

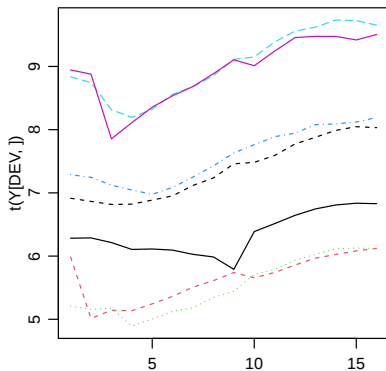
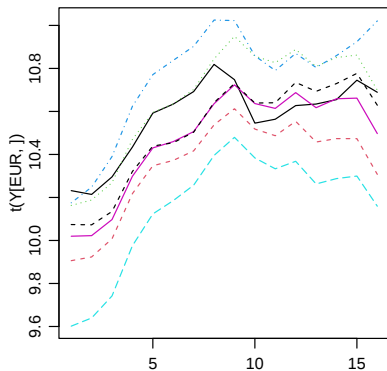
## Linear mixed model fit by REML ['lmerMod']
## Formula: y ~ 1 + (1 | xrow) + (1 | xcol)
##
## REML criterion at convergence: 660.1
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -5.6522 -0.5398  0.0014  0.5775  4.3365
##
## Random effects:
##   Groups    Name      Variance Std.Dev.
##   xrow      (Intercept) 2.32474  1.5247
##   xcol      (Intercept) 0.14474  0.3804
##   Residual                0.04703  0.2169
## Number of obs: 2848, groups:  xrow, 178; xcol, 16
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)   8.2542     0.1487   55.49
```

Estimated effects

```
aHat<-ranef(fit)[[1]][[1]] ; names(aHat)<-rownames(ranef(fit)[[1]] )  
bHat<-ranef(fit)[[2]][[1]] ; names(bHat)<-rownames(ranef(fit)[[2]] )
```



Lack of additivity



Multiplicative models

Consider an “interaction:”

$$y_{i,j} = \mu + a_{i,1} + b_{j,1} + a_{i,2}b_{j,2} + \epsilon_{i,j}$$

- For countries with $a_{i,2} \approx 0$, the time trend is $\{b_{1,1}, \dots, b_{1,T}\}$.
- For countries with $a_{i,2} \approx 1$, the time trend is $\{b_{1,1} + b_{2,1}, \dots, b_{1,T} + b_{2,T}\}$.

Such a model allows for time trajectories that vary across units.

To accommodate more trajectories, we increase the number of interactions:

$$y_{i,j} = \mu + a_i + b_j + \mathbf{u}_i^\top \mathbf{v}_j + \epsilon_{i,j}$$
$$\mathbf{u}_i^\top \mathbf{v}_j = u_{i,1}v_{j,1} + \dots + u_{i,r}v_{j,r}.$$

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AMMI models

Such models are called “AMMI models”:

- additive main effects;
- multiplicative interactions.

The basic structure is used widely in modern multivariate analysis:

- panel data;
- consumer ratings of products;
- recommender systems;
- network analysis;
- genomics and bioinformatics.

The models are nonlinear, but are the next easiest thing.

Bilinear estimation via linear methods

Consider iterative estimation:

Update column effects: Given row effects $(a_i, \mathbf{u}_i)$,

$$\begin{aligned} y_{i,j} &= \mu + (a_i, \mathbf{u}_i)^\top (b_j, \mathbf{v}_j) + \epsilon_{i,j} \\ &\equiv \mu + \mathbf{x}_i^\top \beta_j + \epsilon_{i,j} \end{aligned}$$

which is *linear* in the column effects $(b_j, \mathbf{v}_j)$.

Similarly,

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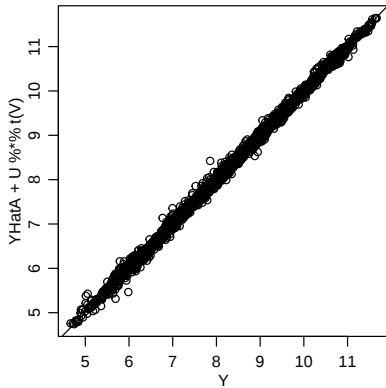
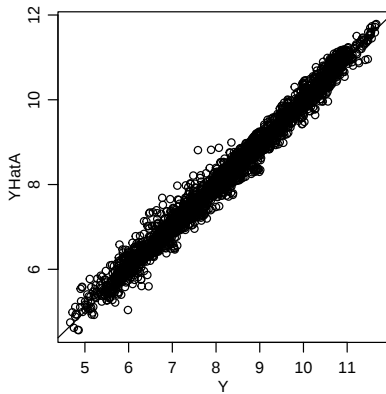
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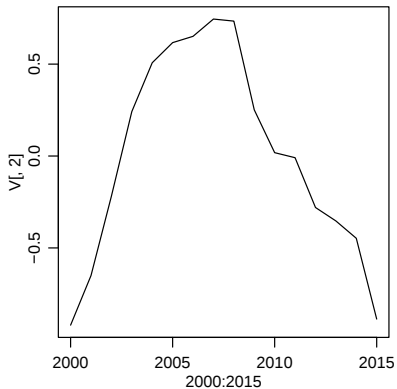
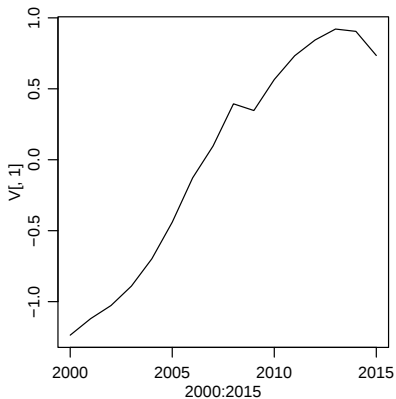
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Two-dimensional GDP model



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