

# **Analysis of Longitudinal Data in Stata, Splus and SAS**

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# OUTLINE

- Longitudinal data
- Review
- Sample data set
- STATA (XTGEE, XTREG, GLLAMM6)
- SAS (Proc Mixed (Repetead, Random), Proc Glinmix, Proc Genmod)
- Splus (LME, YAGS)
- References

# Longitudinal Data

- **Longitudinal Studies:** studies in which the outcome variable is measured repeatedly over time. We do not necessarily require the same number of observations on each subject or that measurements be taken at the same times.

$y_{ij}$  = value of  $j^{th}$  observation on the  $i^{th}$  subject  
measures at time  $t_{ij}$ .

- **Repeated measures:** Older term used for a special set of longitudinal designs with measurements at a common set of occasions, usually in an experimental design.
- Models for the analysis of longitudinal data can be considered a special case of generalized linear models, with the peculiar feature that the residuals terms are correlated, as the observations at different time points in a longitudinal study are taken on the same subject. Any of the model being proposed must take this dependence into account.

# Potential Advantages of Longitudinal Studies

- Allow investigation of events that occur in time; essential to the study of normal growth and ageing.
- Essential to the study of temporal patterns of response to treatments.
- Permit more complete ascertainment of exposure histories in epidemiological studies.
- Reduce unexplained variability in the response by using subject as his or her own control.

# Normally Distributed Data - Marginal Models

With longitudinal data, we can consider models of the form

$$Y_{ij} = \beta_0 + \beta_1 X_{1ij} + \beta_2 X_{2ij} + \dots + \beta_Q X_{Qij} + \epsilon_{ij}$$

where the  $\epsilon_{ij}$  are correlated within individuals (i.e.  $Cov(\epsilon_{ij}, \epsilon_{ik}) \neq 0$ ) and the covariates  $(X_{1ij}, \dots, X_{Qij})$  include time,  $t_{ij}$  (or indicators of time trends), treatment/exposure indicators and their interactions.

Recall that the “compound symmetry” assumption is unrealistic for longitudinal studies, instead we need to consider alternative models for  $Cov(\epsilon_{ij}, \epsilon_{ik})$ .

# Models for the Covariance

:

Note that with  $p$  repeated measures, there are  $\frac{p(p+1)}{2}$  parameters in the covariance matrix.

In selecting a model for the covariance matrix, a balance must be struck:

- With too little structure (e.g., unstructured). there may be too many parameters to be estimated with a limited amount of data (information) available  $\implies$  weaker inferences concerning  $\beta$
- With too much structure (e.g., compound symmetry), there is more information available for estimating  $\beta$  but the potential risk of model misspecification  $\implies$  apparently stronger, but potentially biased, inferences concerning  $\beta$

## Other models

A number of additional models for the covariance that may be suitable for longitudinal data are

1. Autoregressive: The first-order autoregressive model, AR(1), has covariances of the form,  $Cov(Y_{ij}, Y_{ik}) = \sigma^2 \rho^{|j-k|}$ , i.e., homogeneous variances and correlations that decline over time.

		occasion			
		1	2	3	4
occasion	1	1	$\rho$	$\rho^2$	$\rho^3$
	2	$\rho$	1	$\rho$	$\rho^2$
	3	$\rho^2$	$\rho$	1	$\rho$
	4	$\rho^3$	$\rho^2$	$\rho$	1

Autoregressive models are appropriate for equally-spaced measurement.

2. Exponential correlation models can handle unequally-spaced measurements.

Suppose that measurements are made at times  $t_j$ , then the covariances are of the form,

$$Cov(Y_{ij}, Y_{ik}) = \sigma^2 \rho^{|t_j - t_k|}.$$

# STATA

**xtgee** fits generalized linear models of  $Y_{ij}$ , with covariates  $X_{ij}$ . Main components of a model:

1. **family** - assumed distribution of the response variables
2. **link** - link between response and its linear predictor
3. **corr** - structure of the working correlation



# Stata-xtgee

```
*****
*
* Sample program for NASUG 2001
* Data set: depress.dat from Hasbekt & Everitt
* Rino Bellocco
*****
infile subj group pre dep1 dep2 dep3 dep4 dep5 dep6
using c:\rino\nasug\depress.dat, clear
(61 observations read)
```

```
subj  group  pre  dep1  dep2  dep3  dep4  dep5  dep6
  1      0    18   17   18   15   17   14   15
  2      0    27   26   23   18   17   12   10
```

Observations are correlated!

```
      | pre  dep1  dep2  dep3  dep4  dep5  dep6
-----+-----
pre   | 1.0000
dep1  | 0.2027  1.0000
dep2  | 0.2292  0.1937  1.0000
dep3  | 0.1683  0.0700  0.5645  1.0000
dep4  | 0.0561  0.0594  0.5125  0.9015  1.0000
dep5  | 0.1160  0.0654  0.5256  0.9160  0.9606  1.0000
dep6  | 0.1037  0.0184  0.5045  0.9035  0.9499  0.9743  1.0000
```

# Stata-xtgee

First step is to reshape the data so that we can use models.

```
reshape long dep, i(subj) j(visit) (note: j = 1 2 3 4 5 6)
```

subj	visit	group	pre	dep
1	1	0	18	17
1	2	0	18	18
1	3	0	18	15
1	4	0	18	17
1	5	0	18	14
1	6	0	18	15
2	1	0	27	26
2	2	0	27	23
2	3	0	27	18
2	4	0	27	17
2	5	0	27	12
2	6	0	27	10

# Stata-xtgee

First, I run a model with independence structure

```
xtgee dep group pre visit, i(subj) t(visit) corr(indep) link(iden) fam(normal) nmp
```

```
GEE population-averaged model
Group variable:          subj      Number of obs      =      295
Link:                    identity   Number of groups   =      61
Family:                  Gaussian   Obs per group: min =      1
Correlation:            independent max                 =      6
Scale parameter:        25.80052   Wald chi2(3)      =     144.15
                          Prob > chi2 =      0.0000
Pearson chi2(291):      7507.95   Deviance          =     7507.95
Dispersion (Pearson):  25.80052   Dispersion        =     25.80052
```

```
-----+-----
      dep |      Coef.   Std. Err.      z    P>|z|     [95% Conf. Interval]
-----+-----
      group | -4.290664   .6072954    -7.07   0.000   -5.480941   -3.100387
        pre |  .4769071   .0798565     5.97   0.000    .3203913    .633423
      visit | -1.307841   .169842    -7.70   0.000   -1.640725   -.9749569
       _cons |  8.233577   1.803945     4.56   0.000    4.697909   11.76924
-----+-----
```

# Stata-xtgee

Then I fit a GLM with an exchangeable structure

```
. xtgee dep group pre visit, i(subj) t(visit) corr(exc) link(iden) fam(normal)
```

```
Iteration 1: tolerance = .04984936
```

```
Iteration 2: tolerance = .0004433
```

```
Iteration 3: tolerance = 4.602e-06
```

```
Iteration 4: tolerance = 4.782e-08
```

```
GEE population-averaged model
Group variable:          subj      Number of obs      =      295
Link:                   identity    Number of groups   =      61
Family:                 Gaussian    Obs per group: min =      1
Correlation:           exchangeable          avg =      4.8
Scale parameter:       25.56569          max =      6
Wald chi2(3)          =      135.08
Prob > chi2           =      0.0000
```

```
-----+-----
```

dep	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
group	-4.024676	1.081131	-3.72	0.000	-6.143654	-1.905698
pre	.4599018	.1441533	3.19	0.001	.1773666	.742437
visit	-1.226764	.1175009	-10.44	0.000	-1.457062	-.9964666
_cons	8.432806	3.120987	2.70	0.007	2.315783	14.54983

```
-----+-----
```

# Stata-xtgee

Then I fit a model with unstructured correlation

```
xtgee dep group pre visit, i(subj) t(visit) corr(uns) link(iden) fam(normal)
```

```
GEE population-averaged model
Group and time vars:      subj visit
Link:                     identity
Family:                   Gaussian
Correlation:              unstructured
Scale parameter:         25.87029

Number of obs      =      295
Number of groups   =       61
Obs per group: min =       1
                  avg =     4.8
                  max =       6
Wald chi2(3)      =     94.13
Prob > chi2       =     0.0000
```

---

dep	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
group	-4.134413	.9986306	-4.14	0.000	-6.091693	-2.177133
pre	.3399185	.1326684	2.56	0.010	.0798932	.5999437
visit	-1.228327	.1492831	-8.23	0.000	-1.520916	-.9357372
_cons	11.13045	2.892903	3.85	0.000	5.460464	16.80044

---

# Stata-xtgee

## And finally a model with AR1 structure

```
xtgee dep group pre visit, i(subj) t(visit) corr(ar1) link(iden) fam(normal)
note:  some groups have fewer than 2 observations
      not possible to estimate correlations for those groups
      8 groups omitted from estimation
```

```
Iteration 1: tolerance = .10070858
Iteration 2: tolerance = .00136623
Iteration 3: tolerance = .00002736
Iteration 4: tolerance = 5.508e-07
```

```
GEE population-averaged model
Group and time vars:      subj visit      Number of obs      =      287
Link:                     identity        Number of groups   =      53
Family:                   Gaussian        Obs per group: min =      2
Correlation:              AR(1)          avg                =      5.4
Scale parameter:          25.82413        max                =      6
                          Wald chi2(3)   =      64.55
                          Prob > chi2    =      0.0000
```

---

dep	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
group	-4.218194	1.053504	-4.00	0.000	-6.283023	-2.153364
pre	.4268002	.1376156	3.10	0.002	.1570785	.6965219
visit	-1.181975	.1907298	-6.20	0.000	-1.555799	-.8081517
_cons	9.037864	3.036076	2.98	0.003	3.087264	14.98846

---

# SAS-GLM

Here, I show what I think is the equivalent procedure in SAS (codes are reported at the end). Independence:

The REG Procedure  
Model: MODEL1  
Dependent Variable: dep

## Analysis of Variance

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	3	3719.12937	1239.70979	48.05	<.0001
Error	291	7507.95172	25.80052		
Corrected Total	294	11227			

Root MSE	5.07942	R-Square	0.3313
Dependent Mean	11.32915	Adj R-Sq	0.3244
Coeff Var	44.83496		

## Parameter Estimates

Variable	DF	Parameter Estimate	Standard Error	t Value	Pr >  t
Intercept	1	8.23358	1.80395	4.56	<.0001
group	1	-4.29066	0.60730	-7.07	<.0001
pre	1	0.47691	0.07986	5.97	<.0001
visit	1	-1.30784	0.16984	-7.70	<.0001

# SAS-GLM

## Unrestricted Covariance structure

Effect	group	Estimate	Standard Error	DF	t Value	Pr >  t
Intercept		6.2422	2.8737	58	2.17	0.0339
group	0	4.1207	0.9739	58	4.23	<.0001
group	1	0	.	.	.	.
pre		0.3641	0.1292	58	2.82	0.0066
visit		-1.1091	0.1426	58	-7.78	<.0001

## Compound structure

Effect	group	Estimate	Standard Error	DF	t Value	Pr >  t
Intercept		4.4124	3.1901	58	1.38	0.1719
group	0	4.0216	1.0887	58	3.69	0.0005
group	1	0	.	.	.	.
pre		0.4598	0.1452	58	3.17	0.0025
visit		-1.2259	0.1167	233	-10.50	<.0001

## AR1 structure

Effect	group	Estimate	Error	DF	t Value	Pr >  t
Intercept		5.0946	2.9691	58	1.72	0.0915
group	0	4.0317	1.0015	58	4.03	0.0002
group	1	0	.	.	.	.
pre		0.4296	0.1331	58	3.23	0.0021
visit		-1.2221	0.1844	233	-6.63	<.0001



# SAS-GLM

```
libname rino 'c:\rino\nasug';
data rino;
infile 'c:\rino\nasug\depress.dat';
input subj group pre dep1 dep2 dep3 dep4 dep5 dep6;

if dep1=-9 then dep1=.;
if dep2=-9 then dep2=.;
if dep3=-9 then dep3=.;
if dep4=-9 then dep4=.;
if dep5=-9 then dep5=.;
if dep6=-9 then dep6=.;
run;
proc means;
var dep1 dep2 dep3 dep4 dep5 dep6 group pre;
run;

data rino1;
set rino;

visit=1; dep=dep1;t=1;output;
visit=2; dep=dep2;t=2;output;
visit=3; dep=dep3;t=3;output;
visit=4; dep=dep4;t=4;output;
visit=5; dep=dep5;t=5;output;
visit=6; dep=dep6;t=6;output;
run;
proc means;
var dep time pre group;
run;

/* proc print data=rino1;
run;
*/
proc reg data=rino1;
model dep=group pre visit ;
run;

proc mixed data=rino1 noclprint method=ml ;
class subj group t;
```

```
model dep = group pre visit /s;
repeated t /type=un subject=subj r;
title 'unrest.cov. structure, linear trend, ML';
run;
```

```
proc mixed data=rino1 noclprint method=ml;
class subj group t;
model dep = group pre visit /s;
repeated t /type=cs subject=subj r;
title 'compound structure, linear trend, ML';
run;
```

```
proc mixed data=rino1 noclprint method=ml;
class subj group t;
model dep = group pre visit /s;
repeated t /type=ar(1) subject=subj r;
title 'ar1 structure, linear trend, ML';
run;
```

```
proc mixed data=rino1 noclprint method=ml;
class subj group t;
model dep = group pre visit /s;
random intercept /type =un sub=subj s;
title 'random intercept, linear trend, ML';
run;
```

```
proc mixed data=rino1 noclprint method=ml;
class subj group t;
model dep = group pre visit /s;
random intercept visit /type =un sub=subj s;
title 'random intercept, linear trend, ML';
run;
```

## **Stata SAS- comparison**

Similar results are observed, however not the same estimates are produced. Testing and comparison of models with different covariance structures will be reported in a future paper (most likely an STB bulletin).

# Normally Distributed Data

## Random Effect Models

This approach assumes that the correlation arises among repeated measures as the regression coefficients vary across individuals.

That is, each subject is assumed to have an (unobserved) underlying level of response which persists across the  $p$  measurements.

This subject effect is treated as random and the model becomes

$$Y_{ij} = \beta_0 + \beta_1 X_{1ij} + \beta_2 X_{2ij} + \dots + \beta_{p-1} X_{p-1,ij} + b_i + e_{ij}$$

or

$$Y_{ij} = (\beta_0 + b_i) + \beta_1 X_{1ij} + \beta_2 X_{2ij} + \dots + \beta_{p-1} X_{p-1,ij} + e_{ij}$$

(also known as “random intercepts model”).

In the model

$$Y_{ij} = \beta_0 + \beta_1 X_{1ij} + \beta_2 X_{2ij} + \dots + \beta_{p-1} X_{p-1,ij} + b_i + e_{ij}$$

the response for the  $i^{\text{th}}$  subject is assumed to differ from the population mean, by a subject effect,  $b_i$ , and a within-subject measurement error,  $e_{ij}$ .

Alternatively, we have decomposed

$$\epsilon_{ij} = b_i + e_{ij}.$$

Furthermore, it is assumed that

$$b_i \stackrel{d}{=} N(0, \sigma_b^2); \quad e_{ij} \stackrel{d}{=} N(0, \sigma_e^2)$$

and that  $b_i$  and  $e_{ij}$  are mutually independent.

The introduction of a random subject effect induces correlation among the repeated measures.

It can be shown that the following correlation structure results:

$$\text{Var}(Y_{ij}) = \sigma_b^2 + \sigma_e^2$$

$$\text{Cov}(Y_{ij}, Y_{ik}) = \sigma_b^2$$

$$\implies \text{Corr}(Y_{ij}, Y_{lj}) = \frac{\sigma_b^2}{\sigma_b^2 + \sigma_e^2}$$

= correlation of observations on the same individual

**Stata** can fit this model using the **XTREG** procedure.

# XTREG/Stata

```
. xtreg dep group pre visit, i(subj) mle
```

```
Random-effects ML regression      Number of obs      =      295
Group variable (i) : subj         Number of groups   =      61

Random effects u_i ~ Gaussian    Obs per group: min =      1
                                   avg =      4.8
                                   max =      6

Log likelihood = -832.36607       LR chi2(3)         =      111.62
                                   Prob > chi2         =      0.0000
```

dep	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
group	-4.021599	1.08894	-3.69	0.000	-6.155882	-1.887316
pre	.4597672	.1451952	3.17	0.002	.1751898	.7443446
visit	-1.225857	.1168668	-10.49	0.000	-1.454912	-.9968024
_cons	8.434001	3.142894	2.68	0.007	2.274042	14.59396
/sigma_u	3.805795	.4160801	9.15	0.000	2.990293	4.621297
/sigma_e	3.346938	.15434	21.69	0.000	3.044438	3.649439
rho	.5638883	.0600327			.4451442	.6771015

```
Likelihood ratio test of sigma_u=0: chibar2(01)= 127.28
Prob>=chibar2 = 0.000
```

# SAS

random intercept, linear trend, ML

## Covariance Parameter Estimates

Cov Parm	Subject	Estimate
UN(1,1)	subj	14.4836
Residual		11.2021

## Fit Statistics

-2 Log Likelihood	1664.7
AIC (smaller is better)	1676.7
AICC (smaller is better)	1677.0
BIC (smaller is better)	1689.4

## Solution for Fixed Effects

Effect	group	Standard		DF	t Value	Pr >  t
		Estimate	Error			
Intercept		4.4124	3.1901	58	1.38	0.1719
group	0	4.0216	1.0887	233	3.69	0.0003
group	1	0	.	.	.	.
pre		0.4598	0.1452	233	3.17	0.0017
visit		-1.2259	0.1167	233	-10.50	<.0001

## Type 3 Tests of Fixed Effects

Effect	Num	Den	F Value	Pr > F
	DF	DF		
group	1	233	13.64	0.0003
pre	1	233	10.03	0.0017
visit	1	233	110.35	<.0001



# Splus

```
> summary(rem0)
Linear mixed-effects model fit by REML
Data: rino
      AIC      BIC    logLik
1678.536 1700.576 -833.2679

Random effects:
Formula: visit ~ 1 | subj
      (Intercept) Residual
StdDev:    3.923239 3.353891

Fixed effects: dep ~ visit + pre + group
              Value Std.Error  DF   t-value p-value
(Intercept)  8.435886  3.224813 233    2.61593  0.0095
visit       -1.224393  0.117018 233   -10.46327 <.0001
pre         0.459552  0.149022  58    3.08379  0.0031
group      -4.016623  1.117115  58   -3.59553  0.0007

Correlation:
      (Intr)  visit    pre
visit -0.107
pre   -0.960  0.005
group -0.130 -0.040 -0.066

Standardized Within-Group Residuals:
      Min      Q1      Med      Q3      Max
-3.840718 -0.5559042 -0.03438542 0.4645086 3.912141

Number of Observations: 295 Number of Groups: 61
```

## Random Intercepts and Slopes Models

A natural extension of the random intercepts model. The introduction of random intercepts and slopes induces a covariance matrix that depends on time ( $t_{ij}$ ).

Consider the following model with intercepts and slopes that vary randomly among subjects

$$Y_{ij} = \beta_0 + \beta_1 t_{ij} + b_{i0} + b_{i1} t_{ij} + e_{ij}$$

Assume that  $b_{i0}$  and  $b_{i1}$  have mean zero and let  $Var(e_{ij}) = \sigma_e^2$ ,  $Var(b_{i0}) = \sigma_{00}^2$ ,  $Var(b_{i1}) = \sigma_{11}^2$ , and  $Cov(b_{i0}, b_{i1}) = \sigma_{01}$ .

Then, it can be shown that

$$Var(Y_{ij}) = \sigma_{00}^2 + 2t_{ij}\sigma_{01} + \sigma_{11}^2 t_{ij}^2 + \sigma_e^2$$

and

$$Cov(Y_{ij}, Y_{ik}) = \sigma_{00}^2 + (t_{ij} + t_{ik})\sigma_{01} + \sigma_{11}^2 t_{ij} t_{ik}$$

That is, the covariance matrix is a function of time. Stata has limited resources for modeling longitudinal data (GLLAMM6 is a routine provided

by Rabe-Hesketh which allows to fits this model, but it is not part of regular Stata and as, Sophia has told me, GLLAMM6 is intended for non-normal data where no exact method exists; instead we can use PROC MIXED in SAS and LME in Splus.

# STATA

```
gen cons=1 eq cons: cons eq slope: visit  
gllamm6 dep group pre  
visit, i(subj) nrf(2) eqs(cons slope) trace
```

```
gllamm model
```

```
log likelihood = -820.90341
```

---

dep	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
group	-3.459758	.9574966	-3.61	0.000	-5.336417	-1.583099
pre	.5769432	.0954126	6.05	0.000	.3899379	.7639484
visit	-1.240965	.1552877	-7.99	0.000	-1.545324	-.9366072
_cons	5.499468	2.249447	2.44	0.014	1.090632	9.908304

---

```
Variance at level 1
```

```
8.1725165 (.86878708)
```

```
Variances and covariances of random effects
```

```
***level 2 (subj)
```

```
var(1): 23.758474 (5.8717413)
```

```
cov(1,2): -2.2504823 (.98450321) cor(1,2): -.53217727
```

```
var(2): .75269674 (.18593369)
```

---

# SAS

random intercept + slope, linear trend, ML

## Iteration History

Iteration	Evaluations	-2 Log Like	Criterion
0	1	1792.01280464	
1	2	1642.82321420	0.00000252
2	1	1642.82181110	0.00000000

Convergence criteria met.

## Covariance Parameter Estimates

Cov Parm	Subject	Estimate
UN(1,1)	subj	22.3135
UN(2,1)	subj	-2.4981
UN(2,2)	subj	0.8352
Residual		8.3660

## Fit Statistics

-2 Log Likelihood	1642.8
AIC (smaller is better)	1658.8

random intercept + slope, linear trend, ML

12:30 Saturday, Mar

The Mixed Procedure

## Fit Statistics

AICC (smaller is better)            1659.3  
 BIC (smaller is better)            1675.7

Null Model Likelihood Ratio Test

DF	Chi-Square	Pr > ChiSq
3	149.19	<.0001

Solution for Fixed Effects

Effect	group	Estimate	Standard Error	DF	t Value	Pr >  t
Intercept		4.2101	3.2138	58	1.31	0.1954
group	0	4.0397	1.0922	181	3.70	0.0003
group	1	0	.	.	.	.
pre		0.4682	0.1456	181	3.22	0.0015
visit		-1.2097	0.1651	52	-7.33	<.0001

# Splus

```
> summary(rem1)
Linear mixed-effects model fit by REML
Data: rino
      AIC      BIC    logLik
1659.905 1689.292 -821.9527

Random effects:
Formula: ~ visit | subj
Structure: General positive-definite
           StdDev  Corr
(Intercept) 4.8414891 (Intercept)
visit 0.9303804 -0.572
Residual 2.8915377

Fixed effects: dep ~ visit + pre + group
           Value Std.Error DF  t-value p-value
(Intercept) 8.243741 3.247253 233 2.538682 0.0118
visit -1.206358 0.167118 233 -7.218614 <.0001
pre 0.468243 0.149474 58 3.132615 0.0027
group -4.034921 1.121173 58 -3.598840 0.0007

Correlation:
(Intr) visit pre
visit -0.139
pre -0.956 0.005
group -0.126 -0.047 -0.067

Standardized Within-Group Residuals:
      Min      Q1      Med      Q3      Max
-3.315408 -0.5357005 -0.09072777 0.4617966 3.058502

Number of Observations: 295 Number of Groups: 61
```

## Non Normal Data

In this case, we cannot always specify a likelihood with an arbitrary structure. We can define random effect models by introducing a random intercept and slope into the linear predictor (generalized linear mixed models). These models can be difficult to estimate (GLLMM6).

In the GEE approach, we can specify any covariance structure and link function without specifying the joint distribution of the the repeated observations.

REM and GEE lead to different interpretations of between subject effects. In the first case, a between subject effect stands for the difference between subjects conditional on the same random effect, while the parameters of GEE represent the average difference between subject.



## References

- Laird, Ware paper on REM, (Biometrics 1982)
- Zeger, Liang, Albert, on GEE (Biometrics, 1988)
- Horton, Lipsitz, on GEE software, (The American Statistician, 1998)