Supplementary Tutorial Duricki, Soleman and Moon (2016)

A click-by-click guide to analysing longitudinal data from animals where some data are missing.

Slides 1 to 31) How to enter and explore data graphically.

Slides 32 to 45) Analysis using RM ANCOVA.

Slides 46 to 74) Analysis using linear models with general covariance structures.

Slides 75 to 82) Restructuring data.

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Supplementary Tutorial, Slide 2: Loading data. Click File>Open>Data.

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Supplementary Tutorial, Slide 3: Navigate to "short_format" data file. You need to have downloaded this from the Nature Protocols website (Supplementary Data 1) or from www.lawrencemoon.co.uk/resources/linearmodels.asp

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Supplementary Tutorial, Slide 4: Data View

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Supplementary Tutorial, Slide 5 In Variable View, click on Blue icon ("…" in green circle) to discover the names of the Levels for the Factor "group". Missing values have been coded "999.00"

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Supplementary Tutorial, Slide 6: Variable View showing four Levels for "group" Factor

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Supplementary Tutorial, Slide 7 and Figure 2: Data View

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Supplementary Tutorial, Slide 8 and Figure 3: Data View of "long_format" data ("long_format.sav" data file from Supplementary Data 1) suitable for analysis using the Mixed Model>Linear procedure

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Supplementary Tutorial, Slide 9: Variable View for "long format" data

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Supplementary Tutorial, Slide 10: To plot individual rat performances over time, from "long format" data, click Graphs>Chart Builder

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7	1	Adult	Stroke	Young-NT3	.0	7	.03									
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14	2	Adult	Sham	Sham	.10	6	.16									
15	2	Adult	Sham	Sham	.10	7	.07									
16	2	Adult	Sham	Sham	.10	8	.10									
17	3	Adult	Sham	Sham	.06	1	.03									
18	3	Adult	Sham	Sham	.06	2	.06									
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Supplementary Tutorial, Slide 11: The Warning dialog reminds you to make sure you set up the variable types in Variable View properly (*i.e.*, select Nominal / Ordinal / Scale correctly for each variable)

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Supplementary Tutorial, Slide 12: Click on "Line" then drag the icon with three lines into the "Chart preview" window



Supplementary Tutorial, Slide 13:



Supplementary Tutorial, Slide 14 and Figure 4: Graphs showing individual rat performances over time arranged by group and colour coded according to Subject number (rat).



Supplementary Tutorial, Slide 15: How to generate a graph of mean group performance over time.



Supplementary Tutorial, Slide 16 and Figure 5: Graphs showing mean performances over time arranged by group. Note that the last data points for AAV-NT3 and AAV-GFP do not include data from rats with missing values



Supplementary Tutorial, Slide 17: Same data but plotted with missing values replaced by Last Value Carried Forward. Data points now include all available data from all animals.



Supplementary Tutorial, Slide 18: How to determine if group variances are similar *i.e.*, test if group variances are "heterogeneous"

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Supplementary Tutorial, Slide 19:



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Supplementary Tutorial, Slide 20:



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Supplementary Tutorial, Slide 21:



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Supplementary Tutorial, Slide 22:



Supplementary Tutorial, Slide 23



Supplementary Tutorial, Slide 24:



Supplementary Tutorial, Slide 25:



Supplementary Tutorial, Slide 26:



Supplementary Tutorial, Slide 27:

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20	-	21	Aged	Stroke	Aged-NT3	.13	.42	.40	.40	.37	.42	.40	.30	.20	
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Supplementary Tutorial, Slide 28:

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12	12	Adult	Stroke	Young-NT	mean_pos	top4	Sort the file	by grouping varia	ibles	.10	.06	.03	.03	
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21	22	Aged	Stroke	Aged-NT3	.10	.26	.10	.22	.11	.18	.10	.06	.13	
22	23	Aged	Stroke	Aged-NT3	.10	.20	.15	.13	.13	.07	.17	.10	.23	
23	24	Aged	Stroke	Aged-GFP	.03	.20	.33	.30	.29	.14	.10	.13	.39	
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Supplementary Tutorial, Slide 29:

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21	24	1	Aged	Stroke	Aged-	GFP	.03	.20	.33	.30	.29	.14	.10	.13	.39	
22	27	7	Aged	Stroke	Aged-	GFP	.06	.33	.13	.06	.17	.36	.06	.19	.45	
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Supplementary Tutorial, Slide 30:

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7	28	Aged	Stroke	Aged-I	💑 group		🖌 🔗 mean	_postop4		.22	.12	.10	.10	
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20	51	Aged	Stroke	Aged-NT3	.0	.23	.20	.23	.07	.13	.12	.13	.13	
21	24	Aged	Stroke	Aged-GFP	.03	.20	.33	.30	.29	.14	.10	.13	.39	
22	27	Aged	Stroke	Aged-GFP	.06	.33	.13	.06	.17	.36	.06	.19	.45	
23	32	Aged	Stroke	Aged-GFP	.03	.40	.33	.13	.30	.28	.23	.17	.48	
24	33	Aaed	Stroke	Aaed-GFP	.17	.33	.22	.26	.33	.40	.38	999.00	999.00	•
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Supplementary Tutorial, Slide 31: These correlations were generated without splitting the output by "group". There is a significant and positive correlation between most pairs of time points. The size of the correlation stays similar with increasing separation of time points. This suggests that a compound symmetric (CS) covariance structure may be appropriate.

Help

*Output2 [Document2] - PASW Statistics Viewer

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Title					Correlatio	ns				
group = Aged-N			mean_postop 1	mean postop	mean_poeton	mean nostop 4	mean_poston	mean nostop R	mean_postop	mean nostop
	mean postop1	Pearson Correlation	- 1	.543**	.393**	.482**	.443**	.345*	.521**	.522**
group = Aged-G		Siq. (2-tailed)		.000	.004	.000	.001	.012	.000	.000
Title		N	53	53		53	- 53	52	59	50
Correlatior	mean_postop2	Pearson Correlation	.543**	1	.545**	.413**	.531**	.433**	.636**	.499**
group = roung-		Sig. (2-tailed)	.000		.000	.002	.000	.001	.000	.000
Correlatior		N	53	53	53					
🖮 🤁 group = Sham	mean_postop3	Pearson Correlation	.393**	.545**	1	.436**	.483**	.371**	.394**	.444**
Title		Sig. (2-tailed)	.004	.000		.001	.000	.007	.005	001
		N	53	53	53	53	53	52	50	50
E Correlations	mean_postop4	Pearson Correlation	.482**	.413**	.436**	1	.521**	.502**	.446**	
→ Title		Sig. (2-tailed)	.000	.002	.001		.000	.000	.001	.002
Active Dataset		N	53	53	53	53	53	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~		
group = Aged-N	mean_postop5	Pearson Correlation	.443**	.531**	.483**	.521**	1	.488**	.407**	.499**
🖄 Title		Sig. (2-tailed)	.001	.000	.000	.000		.000	.003	.000
		N	53	53	53	53	53	52	50	
	mean_postop6	Pearson Correlation	.345*	.433**	.371**	.502**	.488**	1	.352 [*]	.285*
Correlation		Sig. (2-tailed)	.012	.001	.007	.000	.000		.012	.045
i group = Young-		N	52	52	52	52	52	52	50	50
Correlation	mean_postop7	Pearson Correlation	.521**	.636**	.394**	.446**	.407**	.352 [*]	1	452**
🖃 👎 group = Sham		Sig. (2-tailed)	.000	.000	.005	.001	.003	.012		.001
🔤 Title		N	50	50	50	50	50	50	50	50
Correlation	mean_postop8	Pearson Correlation	.522**	.499**	.444**	.427**	.499**	.285 [*]	.452**	1
Correlations		Sig. (2-tailed)	.000	.000	.001	.002	.000	.045	.001	
→ 🚔 Title		N	50	50	50	50	50	50	50	50
Notes	**. Correlation	is significant at the 0.01	level (2-tailed).							
Active Dataset	*. Correlation i	s significant at the 0.05	level (2-tailed).							
	1									
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Supplementary Tutorial, Slide 32: RM ANCOVA

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3	3		<u>C</u> orrela	te	•	848	Tepearea measures	•	.06	.10	.03	.03	.0	.03	.0	
4	4		<u>R</u> egres	sion	•		Variance Component:	S	.13	.13	.13	.03	.10	.03	.09	
5	5		Logline	ar		NТЗ	.0	.21	.09	.10	.14	.03	.10	.03	.09	
6	6		Classi <u>f</u> y	y 		NТЗ	.0	.20	.10	.13	.10	.03	.0	.13	.0	
7	7		Dimensi	ion Reduction		NT3	.0	.28	.25	.13	.07	.07	.06	.09	.13	
8	8		Scale			NT3	.03	.21	.24	.03	.13	.10	.10	.10	.03	
9	9		Nonpar	ametric lests		NT3	.07	.37	.09	.14	.07	.03	.12	.03	.0	
10	10		Foreca	süng		NТЗ	.14	.14	.16	.14	.18	.13	.10	.14	.03	
11	11		Surviva	Response		NТЗ	.0	.29	.06	.03	.14	.06	.03	.14	0.	
12	12			i Valua Analusia	r	NТЗ	.03	.07	.10	.14	.08	.10	.06	.03	.03	
13	13		Muttiple	y value Anal <u>y</u> sis		NT3	.13	.32	.13	.16	.21	.17	.03	.09	0.	
14	14		Comple	s Imputation		NT3	.0	.23	.13	.06	.13	.10	.03	.10	.03	
15	15		Quality	Control		ham	.03	.14	.10	.03	.07	.14	.07	.10	.03	
16	16			irua	r	NT3	.03	.19	.0	.20	.03	.13	.06	.03	.0	
17	17		7 3001		roung	NT3	.13	.20	.0	.06	.10	.0	.10	.10	.03	
18	18		Adult	Stroke	Young	-NT3	.06	.44	.16	.14	.14	.07	.03	.12	.16	
19	19		Aged	Stroke	Aged	-NT3	.04	.26	.17	.19	.25	.20	.50	.13	.10	
20	21		Aged	Stroke	Aged	-NT3	.13	.42	.46	.46	.37	.42	.46	.30	.26	
21	22		Aged	Stroke	Aged	-NT3	.10	.26	.10	.22	.11	.18	.10	.06	.13	
22	23		Aged	Stroke	Aged	-NT3	.10	.20	.15	.13	.13	.07	.17	.10	.23	
23	24		Aged	Stroke	Aged-	GFP	.03	.20	.33	.30	.29	.14	.10	.13	.39	
24	25		Aaed	Stroke	Aaed	-NT3	.03	.20	.38	.03	.13	.0	.10	.21	.03	T
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Supplementary Tutorial, Slide 33

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3	3	Adult	Sham	Sham	.06			10	.03	.03	.0	.03	.0		
4	4	Adult	Stroke	Young-NT3	.0	<u>Wi</u> thin-Subje	ct Factor Name:	13	.13	.03	.10	.03	.09		
5	5	Adult	Stroke	Young-NT3	.0		wave	10	.14	.03	.10	.03	.09		
6	6	Adult	Stroke	Young-NT3	.0	Number of L	evels: 8	13	.10	.03	.0	.13	.0		
7	7	Adult	Stroke	Young-NT3	.0	0. et et	wave(8)	13	.07	.07	.06	.09	.13		
8	8	Adult	Stroke	Young-NT3	.03	Auu		03	.13	.10	.10	.10	.03		
9	9	Adult	Stroke	Young-NT3	.07	Change		14	.07	.03	.12	.03	.0		
10	10	Adult	Stroke	Young-NT3	.14	Remove		14	.18	.13	.10	.14	.03		
11	11	Adult	Stroke	Young-NT3	.0	Magauna Mar		p3	.14	.06	.03	.14	.0		
12	12	Adult	Stroke	Young-NT3	.03	weasure <u>N</u> ai	me.	14	.08	.10	.06	.03	.03		
13	13	Adult	Stroke	Young-NT3	.13	l r		16	.21	.17	.03	.09	.0		
14	14	Adult	Stroke	Young-NT3	.0	Add		D6	.13	.10	.03	.10	.03		
15	15	Adult	Sham	Sham	.03	Change		03	.07	.14	.07	.10	.03		
16	16	Adult	Stroke	Young-NT3	.03	Remove		20	.03	.13	.06	.03	.0		
17	17	Adult	Stroke	Young-NT3	.13	Itemove		D6	.10	.0	.10	.10	.03		
18	18	Adult	Stroke	Young-NT3	.06			14	.14	.07	.03	.12	.16		
19	19	Aged	Stroke	Aged-NT3	.04	Define	Reset Cancel	Help 19	.25	.20	.50	.13	.10		
20	21	Aged	Stroke	Aged-NT3	.13	.42	.46	.46	.37	.42	.46	.30	.26		
21	22	Aged	Stroke	Aged-NT3	.10	.26	.10	.22	.11	.18	.10	.06	.13		
22	23	Aged	Stroke	Aged-NT3	.10	.20	.15	.13	.13	.07	.17	.10	.23		
23	24	Aged	Stroke	Aged-GFP	.03	.20	.33	.30	.29	.14	.10	.13	.39		
24	25	Aaed	Stroke	Aaed-NT3	.03	.20	.38	.03	.13	.0	.10	.21	.03		-
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Supplementary Tutorial, Slide 34: Note that the Repeated Measure only includes the eight post-treatment time points and does <u>not</u> include the pre-operative baseline measurement time point (as this reduces the power of the test).

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4	4	Adult	Stroke	You 🏼 🍕	rat		(wave):		Contrasts	.03	.10	.03	.09		
5	5	Adult	Stroke	You 🍧	Adult_vs_Aged	+	mean_p	ostopi(i) ostop2(2)	Ploto	.03	.10	.03	.09		
6	6	Adult	Stroke	You	n ijai y		mean_p	ostop3(3)	FIO <u>I</u> S	.03	.0	.13	.0		
7	7	Adult	Stroke	You			mean_po	ostop4(4)	Post Hoc	.07	.06	.09	.13		
8	8	Adult	Stroke	You		•	mean_po	ostop5(5) ostop6(6)	Save	.10	.10	.10	.03		
9	9	Adult	Stroke	You			mean_p	ostop7(7)	Options	.03	.12	.03	.0		
10	10	Adult	Stroke	You			mean_po	ostop8(8)		.13	.10	.14	.03		
11	11	Adult	Stroke	You			D			.06	.03	.14	.0		
12	12	Adult	Stroke	You				-Subjects Factori	(8):	.10	.06	.03	.03		
13	13	Adult	Stroke	You				4P		.17	.03	.09	.0		
14	14	Adult	Stroke	You						.10	.03	.10	.03		
15	15	Adult	Sham				Covariate	· ·		.14	.07	.10	.03		
16	16	Adult	Stroke	You		_	mea	n preop		.13	.06	.03	.0		
17	17	Adult	Stroke	You			•			.0	.10	.10	.03		
18	18	Adult	Stroke	You						.07	.03	.12	.16		
19	19	Aged	Stroke	Ag	ſ	OK Paste	Reset Ca	ancel Help		.20	.50	.13	.10		
20	21	Aged	Stroke	Ag						.42	.46	.30	.26		
21	22	Aged	Stroke	Aged-N1	r3 .10	.26	.10	.22	.11	.18	.10	.06	.13		
22	23	Aged	Stroke	Aged-N1	r3 .10	.20	.15	i .13	.13	.07	.17	.10	.23		
23	24	Aged	Stroke	Aged-GF	P .03	.20	.33	.30	.29	.14	.10	.13	.39		
24	25	Aaed	Stroke	Aaed-N	13 .03	.20	.38	.03	.13	.0	.10	.21	.03		-
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Supplementary Tutorial, Slide 35:

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2	2	Adult	Sham	🔛 Re	peated Me	asures				.03	.16	.07	.10			
3	3	Adult	Sham			🔢 Repeated Measu	ires: Save			.03	.0	.03	.0			
4	4	Adult	Stroke	Your 💊 r	rat	Predicted Values				.03	.10	.03	.09			
5	5	Adult	Stroke	Your 💑	Adult_vs_A	Unstandardized	Unstandardi	zedi	Contrasts	.03	.10	.03	.09			
6	6	Adult	Stroke	Your 🏼 💑 i	njury	Standard error	Standardize	d.	Plots	.03	.0	.13	.0			
7	7	Adult	Stroke	Your					Post <u>H</u> oc	.07	.06	.09	.13			
8	8	Adult	Stroke	Your		Diagnostics	Deleted		<u>S</u> ave	.10	.10	.10	.03			
9	9	Adult	Stroke	Your		Cook's distance			Options	.03	.12	.03	.0			
10	10	Adult	Stroke	Your		Leverage values				.13	.10	.14	.03			
11	11	Adult	Stroke	Your		Coefficient Statistics				.06	.03	.14	.0			
12	12	Adult	Stroke	Your		Create coefficien	t statistics		1	.10	.06	.03	.03			
13	13	Adult	Stroke	Your		Create a new	/ dataset			.17	.03	.09	.0			
14	14	Adult	Stroke	Your		Dataset nam	e:			.10	.03	.10	.03			
15	15	Adult	Sham			🔘 Write a new	data file			.14	.07	.10	.03			
16	16	Adult	Stroke	Your		File			1	.13	.06	.03	.0			
17	17	Adult	Stroke	Your						.0	.10	.10	.03			
18	18	Adult	Stroke	Your		Continue	Cancel Help			.07	.03	.12	.16			
19	19	Aged	Stroke	Age				_		.20	.50	.13	.10			
20	21	Aged	Stroke	Age						.42	.46	.30	.26			
21	22	Aged	Stroke	Aged-NT3		.10 .26	.10	.22	.11	.18	.10	.06	.13			
22	23	Aged	Stroke	Aged-NT3		.10 .20	.15	.13	.13	.07	.17	.10	.23			
23	24	Aged	Stroke	Aged-GFP		.03 .20	.33	.30	.29	.14	.10	.13	.39			
24	25	Aged	Stroke	Aged-NT3		.03 .20	.38	.03	.13	.0	.10	.21	.03	~		
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2	2	Adult	Sham			Estimated Marginal Means					.03	.16	.07	.10		
3	3	Adult	Sham			Factor(s) and Factor Interactions:		Display <u>M</u> e	ans for:		.03	.0	.03	.0		
4	4	Adult	Stroke	You		(OVERALL)		(OVERALI group	_)	в	.03	.10	.03	.09		
5	5	Adult	Stroke	You		wave	*	wave			.03	.10	.03	.09		
6	6	Adult	Stroke	You		group*wave		group*wa	ve		.03	.0	.13	.0		
7	7	Adult	Stroke	You							.07	.06	.09	.13		
8	8	Adult	Stroke	You				🗹 Compa	re main effects		.10	.10	.10	.03		
9	9	Adult	Stroke	You				Confidence	e interval adjustment	t	.03	.12	.03	.0		
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11	11	Adult	Stroke	You							.06	.03	.14	.0		
12	12	Adult	Stroke	You		Display	_				.10	.06	.03	.03		
13	13	Adult	Stroke	You		Descriptive statistics	T	ransformatio	n matrix		.17	.03	.09	.0		
14	14	Adult	Stroke	You		Estimates of effect size	✓ ⊢	lomogeneity t	ests		.10	.03	.10	.03		
15	15	Adult	Sham			Observed power		pread vs. lev	el plot		.14	.07	.10	.03		
16	16	Adult	Stroke	You		Parameter estimates		esidual plot			.13	.06	.03	.0		
17	17	Adult	Stroke	You		Sop matrices Sop matrices		ack of fit	able function		0.	.10	.10	.03		
18	18	Adult	Stroke	You	Ч			eneral estim			.07	.03	.12	.16		
19	19	Aged	Stroke	Ag		Significance level: .05 Confide	nce interv	vals are 95.0	%		.20	.50	.13	.10		
20	21	Aged	Stroke	Ag	_	Continue	Concol	Holo	1		.42	.46	.30	.26		
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Supplementary Tutorial, Slide 38 Ensure there is ONLY one full stop (period), at the end. Now click Run>All

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Supplementary Tutorial, Slide 45: Scroll down to group * wave and then inspect pairwise comparisons

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Supplementary Tutorial, Slide 46. Make sure you downloaded and unzipped the "Supplementary Data 1" file from Nature Protocols or from

www.lawrencemoon.co.uk/resources/mixedmodels.asp

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Supplementary Tutorial, Slide 47 and Figure 3

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Supplementary Tutorial, Slide 48 and Figure 3

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Supplementary Tutorial, Slide 49 and Figure 4

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Supplementary Tutorial, Slide 50 and Figure 4

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Supplementary Tutorial, Slide 51 and Figure 5

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Supplementary Tutorial, Slide 52 and Figure 5

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Supplementary Tutorial, Slide 54 and Figure 6

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Supplementary Tutorial, Slide 55: Analysis using linear model and a Compound Symmetric covariance structure and estimated using REML



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Supplementary Tutorial, Slide 57: Comparing -2LL from two different models

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Supplementary Tutorial, Slide 58: Comparing -2LL from two different models

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Supplementary Tutorial, Slide 59: Comparing -2LL from two different models

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Supplementary Tutorial, Slide 60 and Figure 7: Analysis using linear model and CS covariance structure estimated using REML



Supplementary Tutorial, Slide 61 and Figure 7

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Supplementary Tutorial, Slide 66: The vast majority of circles fall on the diagonal line, with only a few outliers (green circle). This Q-Q plot indicates the assumption of normality is reasonable.



Supplementary Tutorial, Slide 67: A scatterplot of the residuals against their predicted values shows no obvious correlation within groups. However, the variability of the residuals is smaller for the sham and young groups than for the two aged groups.



Supplementary Tutorial, Slide 68: Making the model more parsimonious

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Supplementary Tutorial, Slide 69: Analysis using CS covariance structure with estimation performed using REML

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Supplementary Tutorial, Slide 72:



Supplementary Tutorial, Slide 73:

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🍎 M	lodel Dimensi		7	Aged-NT3	Aged-GFP	048	.035	293.544	.171	118	.021				
i	nformation Crit				Young-NT3	.073*	.034	280.688	.033	.006	.140				
F	ixed Effects				Sham	.089	.061	277.155	.150	032	.210				
	🚡 Type III Tes			Aged-GFP	Aged-NT3	.048	.035	293.544	.171	021	.118				
🖨 🖷 📴 C	ovariance Par				Young-NT3	.122*	.037	286.931	.001	.048	.195				
	🔄 Title				Sham	.137*	.063	281.456	.031	.013	.262				
	stimated Marc			Young-NT3	Aged-NT3	073*	.034	280.688	.033	140	006				
	🖻 Title 🔋				Aged-GFP	122*	.037	286.931	.001	195	048				
	🚡 1. Grand M				Sham	.016	.063	275.252	.804	108	.139				
	/ 2. group			Sham	Aged-NT3	089	.061	277.155	.150	210	.032				
	Estim				Aged-GFP	137*	.063	281.456	.031	262	013				
	📲 Pairwi	1			Young-NT3	016	.063	275.252	.804	139	.108				
	🛄 Univar		8	Aged-NT3	Aged-GFP	153*	.035	293.544	.000	222	083				
	3. wave				Young-NT3	.111*	.034	280.688	.001	.044	.178				
	Estim				Sham	.114	.061	277.155	.064	007	.235				
	🛁 👰 Pairwi			Aged-GFP	Aged-NT3	.153*	.035	293.544	.000	.083	.222				
	Univar				Young-NT3	.264*	.037	286.931	.000	.190	.337				
	4. group *				Sham	.267*	.063	281.456	.000	.142	.391				
	Title			Young-NT3	Aged-NT3	111*	.034	280.688	.001	178	044				
	- 🔓 Estim				Aged-GFP	264*	.037	286.931	.000	337	190				
	la Pairwi				Sham	.003	.063	275.252	.963	120	.126				
.	6. group *			Sham	Aged-NT3	114	.061	277.155	.064	235	.007				
	Title				Aged-GFP	267	.063	281.456	.000	391	142				
	Estim				Young-NT3	003	.063	275.252	.963	126	.120				
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Supplementary Tutorial, Slide 75: Restructuring data. First save the data file by File>Save As because the Restructuring overwrites your existing file. Next click Data>Restructure.

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		8	D <u>e</u> fine Dates				group	mean_preop	mean_posto	p mean_postop	mean_postop	mean_postop	mean_postop	mean_postop	mean_postop	mean_postop	var
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2			Validation		•	m	Sham	10		3 .10 3 .17	06	.07	.03	.00	.03	10	
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4		\mathbf{Q}	dentify Unusual C	Cases		<e< th=""><th>Young-NT3</th><th>.0</th><th>.2</th><th>5 .13</th><th>.13</th><th>.13</th><th>.03</th><th>.10</th><th>.03</th><th>.09</th><th></th></e<>	Young-NT3	.0	.2	5 .13	.13	.13	.03	.10	.03	.09	
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6		3Ú	Sort Varia <u>b</u> les			<e< th=""><th>Young-NT3</th><th>.0</th><th>.20</th><th>.10</th><th>.13</th><th>.10</th><th>.03</th><th>.0</th><th>.13</th><th>0.</th><th></th></e<>	Young-NT3	.0	.20	.10	.13	.10	.03	.0	.13	0.	
7		4	Tra <u>n</u> spose		-	ke	Young-NT3	.0	.28	3	.13	.07	.07	.06	.09	.13	
8		*	<u>R</u> estructure		-	ke	Young-NT3	.03	.2	.24	.03	.13	.10	.10	.10	.03	
9			Merge Files		•	ke	Young-NT3	.07	.37	7 .09	.14	.07	.03	.12	.03	0.	
10		a	Aggregate		-	ke	Young-NT3	.14	.14	.16	.14	.18	.13	.10	.14	.03	
11			Ort <u>h</u> ogonal Desigr	n	•	ke	Young-NT3	.0	.29	3 .06	.03	.14	.06	.03	.14	.0	
12		₩.	Copy <u>D</u> ataset		·	<e< th=""><th>Young-NT3</th><th>.03</th><th>.07</th><th>7 .10</th><th>.14</th><th>.08</th><th>.10</th><th>.06</th><th>.03</th><th>.03</th><th></th></e<>	Young-NT3	.03	.07	7 .10	.14	.08	.10	.06	.03	.03	
13			Split <u>F</u> ile			ke	Young-NT3	.13	.32	2	.16	.21	.17	.03	.09	.0	
14		Ħ	Select Cases		-	ke	Young-NT3	.0	.23	3	.06	.13	.10	.03	.10	.03	
15		ata -	– Weight Cases			m	Sham	.03	.14	4 .10	.03	.07	.14	.07	.10	.03	
16						ke	Young-NT3	.03	.19	9	.20	.03	.13	.06	.03	0.	
17			17	Adult	Strok	ke	Young-NT3	.13	.20	0. (.06	.10	.0	.10	.10	.03	
18			18	Adult	Strok	ke	Young-NT3	.06	.4	4 .16	.14	.14	.07	.03	.12	.16	
19			19	Aged	Strok	<e< th=""><th>Aged-NT3</th><th>.04</th><th>.26</th><th>6 .17</th><th>.19</th><th>.25</th><th>.20</th><th>.50</th><th>.13</th><th>.10</th><th></th></e<>	Aged-NT3	.04	.26	6 .17	.19	.25	.20	.50	.13	.10	
20			21	Aged	Strok	<e< th=""><th>Aged-NT3</th><th>.13</th><th>.42</th><th>2</th><th>.46</th><th>.37</th><th>.42</th><th>.46</th><th>.30</th><th>.26</th><th></th></e<>	Aged-NT3	.13	.42	2	.46	.37	.42	.46	.30	.26	
21			22	Aged	Strok	<e< th=""><th>Aged-NT3</th><th>.10</th><th>.26</th><th>6 .10</th><th>.22</th><th>.11</th><th>.18</th><th>.10</th><th>.06</th><th>.13</th><th></th></e<>	Aged-NT3	.10	.26	6 .10	.22	.11	.18	.10	.06	.13	
22			23	Aged	Strok	<e< th=""><th>Aged-NT3</th><th>.10</th><th>.20</th><th>.15</th><th>.13</th><th>.13</th><th>.07</th><th>.17</th><th>.10</th><th>.23</th><th></th></e<>	Aged-NT3	.10	.20	.15	.13	.13	.07	.17	.10	.23	
23			24	Aged	Strok	<e< th=""><th>Aged-GFP</th><th>.03</th><th>.20</th><th>.33</th><th>.30</th><th>.29</th><th>.14</th><th>.10</th><th>.13</th><th>.39</th><th></th></e<>	Aged-GFP	.03	.20	.33	.30	.29	.14	.10	.13	.39	
24	4		25	Aaed	Strok	(e	Aaed-NT3	.03	.20	.38	.03	.13	.0	.10	.21	.03	
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Supplementary Tutorial, Slide 78: Restructuring data

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	Name		V Restructure Data Wi	zard - Step 3	of 7	leasure	Role	
1	rat	Numeric 8				minal	🔪 Input	·
2	Adult_vs_Aged	Numeric 8	Variables to Ca	ases: Sele	ect Variables	minal	🔪 Input	
3	injury	Numeric 8	For each variable group vo	ou have in the cur	rent data the restructured file will have one target variable.	minal	🔪 Input	
4	group	Numeric 8	In this step, choose how to	identify case arc	hups in the restructured data, and choose which variables belong with	minal	🔪 Input	
5	mean_preop	Numeric 8	each target variable.			ale	🔪 Input	
6	mean_postop1	Numeric 8	Optionally, you can also ch	noose variables to	copy to the new file as Fixed Variables.	ale	🔪 Input	
7	mean_postop2	Numeric 8				ale	🔪 Input	
8	mean_postop3	Numeric 8	Variables in the <u>C</u> urrent F	ile:		ale	🔪 Input	
9	mean_postop4	Numeric 8	💑 rat		Case Group Identification	ale	🔪 Input	
10	mean_postop5	Numeric 8	Adult_vs_Aged		Use selected variable 💎	ale	🦒 Input	
11	mean_postop6	Numeric 8	aroun		Variable: 💦 rat	ale	🔪 Input	
12	mean_postop7	Numeric 8	💉 group			ale	🔪 Input	
13	mean_postop8	Numeric 8	💉 mean_postop1		Variables to be Transposed	ale	🔪 Input	
14			mean_postop2		Target Variable: outcome			
15			/ mean_postop3					
16			mean_postop4					
17			💞 mean_postop6		✓ mean_postop2			
18			💉 mean_postop7		💉 🖋 mean_postop4			
19			// mean_postop8					
20			_		Fixed Variable(s):			_
21					Aduit_vs_Aged			
22					group			
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Supplementary Tutorial, Slide 79: In this case, you only have One index variable. See next slide for an explanation of the indexing variable.

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	Name	Туре	V 🔝 Restruct	ure Data Wizard - Ste	ep 4 of 7		×	leasure	Role		
1	rat	Numeric 8						minal	ゝ Input		
2	Adult_vs_Aged	Numeric 8	Varia	bles to Cases:	Create Index Varia	bles		minal	ゝ Input		
3	injury	Numeric 8	In the curr	ent data, values for a varia	ible group appear in a single case	in multiple variables. For example	e, a	minal	ゝ Input		
4	group	Numeric 8	single cas	e contains the values for v	v1, w2, and w3.			minal	ゝ Input		
5	mean_preop	Numeric 8	In the new	/ data, values for a variable	e group will appear in multiple cas	es in a single variable. For exampl	le,	ale	ゝ Input		
6	mean_postop1	Numeric 8	nere will t	oe inree cases, one each t	or wit, wz, and ws.	was practed from the original app	- For	ale	ゝ Input		
7	mean_postop2	Numeric 8	example, a	s a new variable triat identi an index named "w" would	have the values 1, 2, and 3.	was created from the original cas	e. rur	ale	ゝ Input		
8	mean_postop3	Numeric 8						ale	ゝ Input		
9	mean_postop4	Numeric 8						ale	ゝ Input		
10	mean_postop5	Numeric 8			How many index variables d	o you want to create?		ale	ゝ Input		
11	mean_postop6	Numeric 8		1 1 2 0.11 1 1 3 0.05	One One			ale	ゝ Input		
12	mean_postop7	Numeric 8		2 1 1 0.08	Use this when a variabl	e group records the effects of a si	ingle	ale	ゝ Input		
13	mean_postop8	Numeric 8		2 1 3 0.06	factor, treatment or con	dition.		ale	ゝ Input		
14				1 1 1 1 0.07	-						
15				1 1 1 2 0.11	◯ More than one Hov	v many? 2					
16				1 1 2 1 0.08	Use this when a variabl	e group records the effects of mor	re than				
17				1 1 2 3 0.06	one factor, treatment or	condition.				-	
18				1 0.08 2 0.07	\square					_	
19				2 1 0.11 2 0.11 3 1 0.07 2 0.05	O Non <u>e</u>					_	
20				1 0.06 2 0.08 5 1 0.09 2 0.04	Use this if index informa	tion is stored in one of the sets of				_	
21	1			5 1 0.02 2 0.06	variables to be transpos	eu.				_	
22			_							-	
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Supplementary Tutorial, Slide 80: Type "wave" into the "Name" for your first indexing variable. This generates a new variable called "wave" with eight levels corresponding to your Repeated Measure time points.

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	Name	Type		Restructure Data Wizard - Step 5 of 7	X leasure	Role]
1	rat	Numeric	8		minal	🔪 Input	_
2	Adult vs Aged	Numeric	8	Variables to Cases: Create One Index Variable	minal	🔪 Input	
3	injury	Numeric	8	You have chosen to create one index variable. The variable's values can be sequential numbers or the names	minal	🔪 Input	-
4	group	Numeric	8	of variables in a group.	minal	🔪 Input	
5	mean_preop	Numeric	8	In the table you can specify the name and label for the index variable.	ale	🔪 Input	
6	mean_postop1	Numeric	8		ale	🔪 Input	
7	mean_postop2	Numeric	8	⊂What kind of index values?	ale	ゝ Input	
8	mean_postop3	Numeric	8	Somertial rumbers	ale	ゝ Input	
9	mean_postop4	Numeric	8	Index Values: 1, 2, 3, 4, 5, 6, 7, 8	ale	ゝ Input	
10	mean_postop5	Numeric	8		ale	ゝ Input	
11	mean_postop6	Numeric	8	Index Values: ween notion? ween notion? ween notion? ween notion?	ale	🔪 Input	
12	mean_postop7	Numeric	8	mear_bostop1, mear_bostop2, mear_bostop3, mear_bostop4, mear_bostop4, mear_bostop4	ale	ゝ Input	
13	mean_postop8	Numeric	8	Edit the Index Variable Name and Label:	ale	ゝ Input	
14				Name Label Levels Index Values			
15				1 wave 8 1, 2, 3, 4, 5, 6, 7, 8			
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Supplementary Tutorial, Slide 81: Click Next, Next, Finish, OK.

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	Name	Туре	V	🗱 Restructure Data Wizard - Finish 🛛 🔊	leasure	Role		
1	rat	Numeric	8		minal	ゝ Input		-
2	Adult_vs_Aged	Numeric	8	Finish	minal	ゝ Input		
3	injury	Numeric	8		minal	ゝ Input		
4	group	Numeric	8		minal	🖒 Input		
5	mean_preop	Numeric	8		ale	🖒 Input		
6	mean_postop1	Numeric	8		ale	🖒 Input		
7	mean_postop2	Numeric	8	What do you want to do?	ale	🖒 Input		
8	mean_postop3	Numeric	8		ale	ゝ Input		
9	mean_postop4	Numeric	8		ale	ゝ Input		
10	mean_postop5	Numeric	8	Use this when you want to replace the current file immediately.	ale	ゝ Input		
11	mean_postop6	Numeric	8		ale	ゝ Input		
12	mean_postop7	Numeric	8		ale	ゝ Input		
13	mean_postop8	Numeric	8		ale	ゝ Input		
14				Case and the second sec				
15				Use this when you want to save or modify the syntax before you				
16				restructure the data.				
17								
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Supplementary Tutorial, Slide 82: Annoyingly, SPSS does not carry forward the "missing value" designation. To fix this, click "Variable View" and the blue "…" icon in "Missing" for "outcome". Click on "Discrete missing values" and enter "999.00" in one box.

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		· 🥣	i 🚣 🗉	a a a					ABC		
;	Name	Туре	Width	Decimals	Label	Values	Missing	Columns	Align	Measure	Role
1	rat	Numeric	8	0		None	None	8	≣ Right	💑 Nominal	🔪 Input
2	Adult_vs_Aged	Numeric	8	2		{1.00, Adult}	. None	8	 ■ Right	💑 Nominal	🔪 Input
3	injury	Numeric	8	0		{1, Sham}	None	8	≡ Right	💑 Nominal	🔪 Input
4	group	Numeric	8	2		{1.00, p}	None	8	≣ Right	💑 Nominal	🔪 Input
5	wave	Numeric	4	0		None	None	8	≣ Right	💑 Nominal	🔪 Input
6	outcome	Numeric	8	2		None	None	8	≣ Right	🛷 Scale	🔪 Input
7											
8					🔛 Missi	ng Values					
9											
10					© <u>N</u> on	nissing values					
11						rete missing value	s				
12					999	.00					
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Supplementary Tutorial, Slide 83: Users of SPSS version 19 or later can use the GENLINMIXED procedure to fit linear models with general error covariance structures and different variance components for different groups of subjects.

Entire books have been written about the GENLINMIXED procedure, so a full explanation is beyond the scope of this article.

However, the following Syntax may be used as guidance to begin fitting a model which would allow males and females to have different error variance components for a Scale (i.e., quantitative, continuous) dependent variable called "weight", an independent variable called "iv1" and a repeated measure called "time".

GENLINMIXED

/DATA_STRUCTURE SUBJECTS=subject REPEATED_MEASURES=time GROUPING=male COVARIANCE_TYPE=IDENTITY /FIELDS TARGET=weight TRIALS=NONE OFFSET=NONE /TARGET_OPTIONS DISTRIBUTION=NORMAL LINK=IDENTITY /FIXED EFFECTS=time male iv1 USE_INTERCEPT=TRUE ...

Specify whatever BUILD_OPTIONS and EMMEANS_OPTIONS you want. The output format needs to be changed from "Diagram" to "Table".

We thank an anonymous reviewer for providing this suggestion.