

相關樣本/配對資料 的統計推論

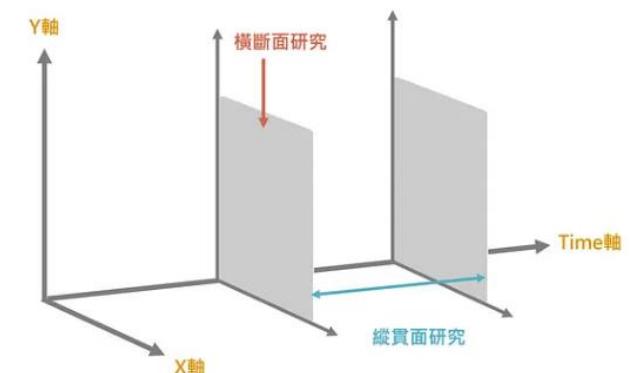
醫學研究部生統小組

陳俊朋

日期: 2024/6/12

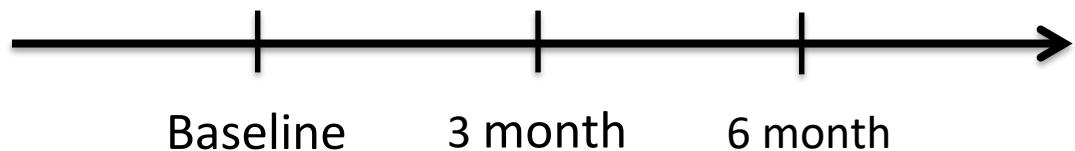
研究設計

Cross-sectional	Longitudinal
Single point in time	Period of time
Different samples	Same sample
Provides snapshot of society at a given point	Follows changes in participants over time
Compare many different variables at the same time	The difference between these studies is the timeline and variable
Cheaper	Expensive and require more resources

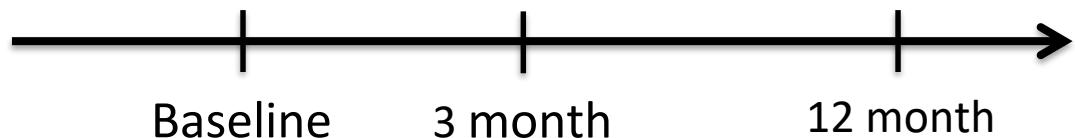


重複測量使用時機

- 相同受試者在不同時間點進行多次測量
 - 間隔時間相同



- 間隔時間不同



重複測量優缺點

- 優點
 - 需求樣本數較少
 - 不同時間點變化
 - 不同組別，隨著時間的變化
- 缺點
 - 資料完整性不佳
 - 個案須定期追蹤
 - 某些因素可能影響資料

重複測量統計

- 單組重複測量
 - 同一組內的個案，在多個時間點測量
- 多組重複測量
 - 將個案分組後，依照各組分類在多個時間點測量

重複測量統計方法

重複2次

- 類別變項
 - McNemar test (2*2 table)
 - McNemar-Bowker test (3*3 table)
- 連續變項
 - Paired t test
 - Wilcoxon signed-rank test
 - ANCOVA

重複3次以上

- 類別變項
 - Cochran's Q test
- 連續變項
 - Friedman test
 - Repeated measures ANOVA
 - Linear Mixed Model (LMM)
 - Generalized Estimating Equation (GEE)

重複測量建檔

- 寬資料

no	Group	Sex	MMSE_1	MMSE_2	MMSE_3
N01	0	0	30	28	30
N02	0	1	11	15	14
N03	0	1	26	28	26
N04	0	0	22	21	22
N05	0	0	27	26	25

- 長資料

no	Group	Sex	visit	MMSE
N01	0	0	1	30
N01	0	0	2	28
N01	0	0	3	30
N02	0	1	1	11
N02	0	1	2	15
N02	0	1	3	14
N03	0	1	1	26
N03	0	1	2	28
N03	0	1	3	26
N04	0	0	1	22
N04	0	0	2	21
N04	0	0	3	22
N05	0	0	1	27
N05	0	0	2	26
N05	0	0	3	25

McNemar test

- 僅適用2*2 table
- 單組同一人前後測/兩種不同診斷工具

International Urogynecology Journal
<https://doi.org/10.1007/s00192-021-04977-7>

ORIGINAL ARTICLE



Transvaginal repair of anterior vaginal wall prolapse with polyvinylidene fluoride (PVDF) mesh: an alternative for previously restricted materials?

Mohammad-Javad Eslami¹ • Mahtab Zargham¹ • Farshad Gholipour² • Mohammadreza Hajian³ • Katayoun Bakhtiari⁴ • Sakineh Hajebrahimi⁵ • Melina Eghbal⁶ • Ziba Farajzadegan⁷

Table 3 Urinary and vaginal symptoms at baseline and 12-month follow-up

	Preoperative	Postoperative (12 months)	P value
Storage symptoms (N=108)			
SUI: n (%)	74 (68.5%)	7 (6.5%)	<0.0001 *
UUI: n (%)	33 (30.5%)	9 (8.3%)	<0.0001 *
MUI: n (%)	29 (26.9%)	3 (2.8%)	<0.0001 *
Other storage symptoms	30 (27.8%)	8 (7.4%)	<0.0001 *
Voiding symptoms: n (%) (N=108)	24 (22.2%)	13 (12%)	0.019 *
ICIQ-UI (mean±SD) (N=98)	8.79±6.05	0.77±1.87	<0.0001 **
ICIQ-VS (mean±SD) (N=41)			
Vaginal symptoms	43.31±6.13	7.49±3.75	<0.0001 **
Sexual symptoms	43.17±6.15	7.82±4.89	<0.0001 **
Quality of life	8.27±0.98	1.6±1.73	<0.0001 **

* McNemar's test

** Wilcoxon signed-rank test

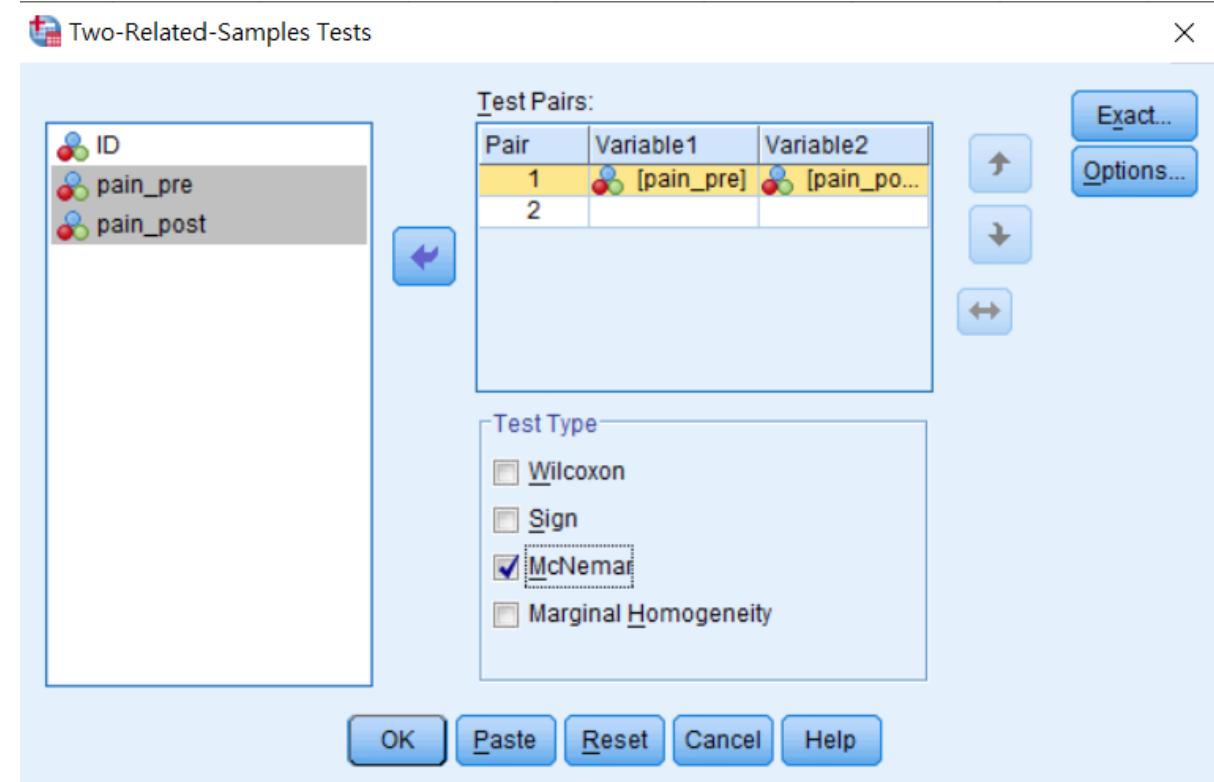
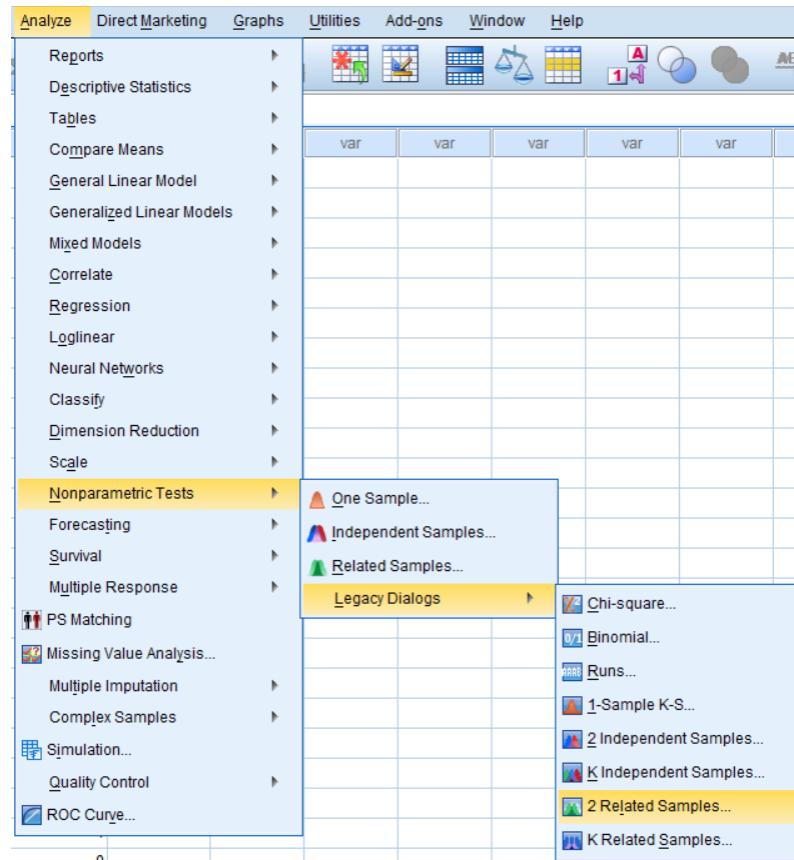
McNemar test dataset

- 術前和術後有無疼痛比較

ID	pain_pre	pain_post
1	1	0
2	0	0
3	0	0
4	0	0
5	0	0
6	1	0
7	0	0
8	0	0
9	0	0
10	0	0
11	0	0
12	1	0
13	0	0
14	0	0
15	0	0

McNemar test analysis

分析>無母數檢定>歷史對話記錄>2個相關樣本



McNemar test output

McNemar Test

Crosstabs

pain_pre & pain_post

		pain_post		Total
		0	1	
pain_pre	0	21	0	
	1	5	1	
	Total	21	6	27

Test Statistics^a

	pain_pre & pain_post
N	27
Exact Sig. (2-tailed)	.063 ^b

a. McNemar Test

b. Binomial distribution used.

pain_post * pain_pre Crosstabulation

		pain_pre		Total
		0	1	
pain_post	0	Count	21	5
	0	% of Total	77.8%	18.5%
1	1	Count	0	1
	1	% of Total	0.0%	3.7%
Total		Count	21	6
		% of Total	77.8%	22.2%
				100.0%

Table.

	Pre-OP	Post-OP 1y	p value
Symptoms			
pain	6(22.2%)	1(3.7%)	0.063

McNemar test. * $p<0.05$, ** $p<0.01$

McNemar-Bowker test

- 適用3*3 table以上
- 單組同一人前後測/兩種不同診斷工具



Table 2. 2016 FSS by 2019 FSS Among Undergraduates (n = 338)

2016 FSS ^a	2019 FSS ^b				
	High	Marginal	Low	Very Low	Total
High	106 (31.4)	36 (10.7)	38 (11.2)	18 (5.3)	198 (58.6)
Marginal	21 (6.2)	16 (4.7)	16 (4.7)	11 (3.3)	64 (18.9)
Low	9 (2.7)	9 (2.7)	9 (2.7)	13 (3.8)	40 (11.8)
Very low	1 (0.3)	1 (0.3)	17 (5.0)	17 (5.0)	36 (10.6)
Total	137 (40.4)	62 (18.3)	80 (23.6)	59 (17.4)	338 (100)

FSS indicates food security status.

^aMeasured using the 10-item Food Security Survey Module; ^bMeasured using the 18-item Food Security Survey Module.

Note: Values displayed as n (% of total). Statistical test conducted is the McNemar-Bowker test ($\chi^2 = 47.878$, degrees of freedom = 6, P < 0.001) and paired samples Wilcoxon signed rank test (P < 0.001). Critical P ≤ 0.05.

McNemar-Bowker test dataset

- 新舊診斷工具一致性比較

ID	grade1	grade2
1	0	1
2	0	1
3	0	1
4	0	2
5	1	2
6	1	2
7	1	1
8	1	1
9	1	1
10	1	1
11	1	1
12	2	1
13	2	1
14	2	1
15	2	1

McNemar-Bowker test analysis

分析>叙述統計>交叉表

The screenshot illustrates the steps to perform a McNemar-Bowker test in SPSS:

- Analyze menu:** The "Analyze" menu is open, with "Descriptive Statistics" selected.
- Crosstabs dialog box:** The "Crosstabs" dialog box is open, showing the variable "grade2" assigned to Row(s) and "grade1" assigned to Column(s).
- Statistics sub-dialog box:** The "Statistics..." button is selected, showing the "Nominal" section. The "McNemar" checkbox is checked under "Nominal by Interval".
- Cell Display sub-dialog box:** The "Cells..." button is selected, showing the "Counts" section with "Observed" checked, and the "Residuals" section with "Total" checked.

McNemar-Bowker test output

grade2 * grade1 Crosstabulation

		grade1			Total	
		0	1	2		
grade2	0	Count	0	2	0	2
	0	% of Total	0.0%	7.4%	0.0%	7.4%
grade2	1	Count	7	7	8	22
	1	% of Total	25.9%	25.9%	29.6%	81.5%
grade2	2	Count	1	2	0	3
	2	% of Total	3.7%	7.4%	0.0%	11.1%
Total		Count	8	11	8	27
		% of Total	29.6%	40.7%	29.6%	100.0%

Table.

	Grade1			<i>p</i> value
	0	1	2	
Grade2				0.061
0	0(0.0%)	2(7.4%)	0(0.0%)	
1	7(25.9%)	7(25.9%)	8(29.6%)	
2	1(3.7%)	2(7.4%)	0(0.0%)	

McNemar test. **p*<0.05, ***p*<0.01

Chi-Square Tests

	Value	df	Asymp. Sig. (2-sided)
McNemar-Bowker Test	7.378	3	.061
N of Valid Cases	27		

Cochran's Q test

- 單組的相同個案，用同一工具測量三次/不同工具(類別型資料屬於二分類)

Open access

Original research

BMJ Open Implementation of a comprehensive surveillance system for recording suicides and attempted suicides in rural India

Lakshmi Vijayakumar,¹ Soumitra Pathare,² Nikhil Jain,² Renuka Nardodkar ,²
Deepa Pandit,² Sadhvi Krishnamoorthy,² Jasmine Kalha,² Laura Shields-Zeeman³

Table 4 Comparison of the ability of community surveillance in obtaining additional data on suicide compared with the hospital and police records

	Non-case	Case	Test statistics
Community surveillance	15	67	Cochran's
Hospital records	69	13	Q test 67.9 ($p<0.01$)
Police records	65	17	

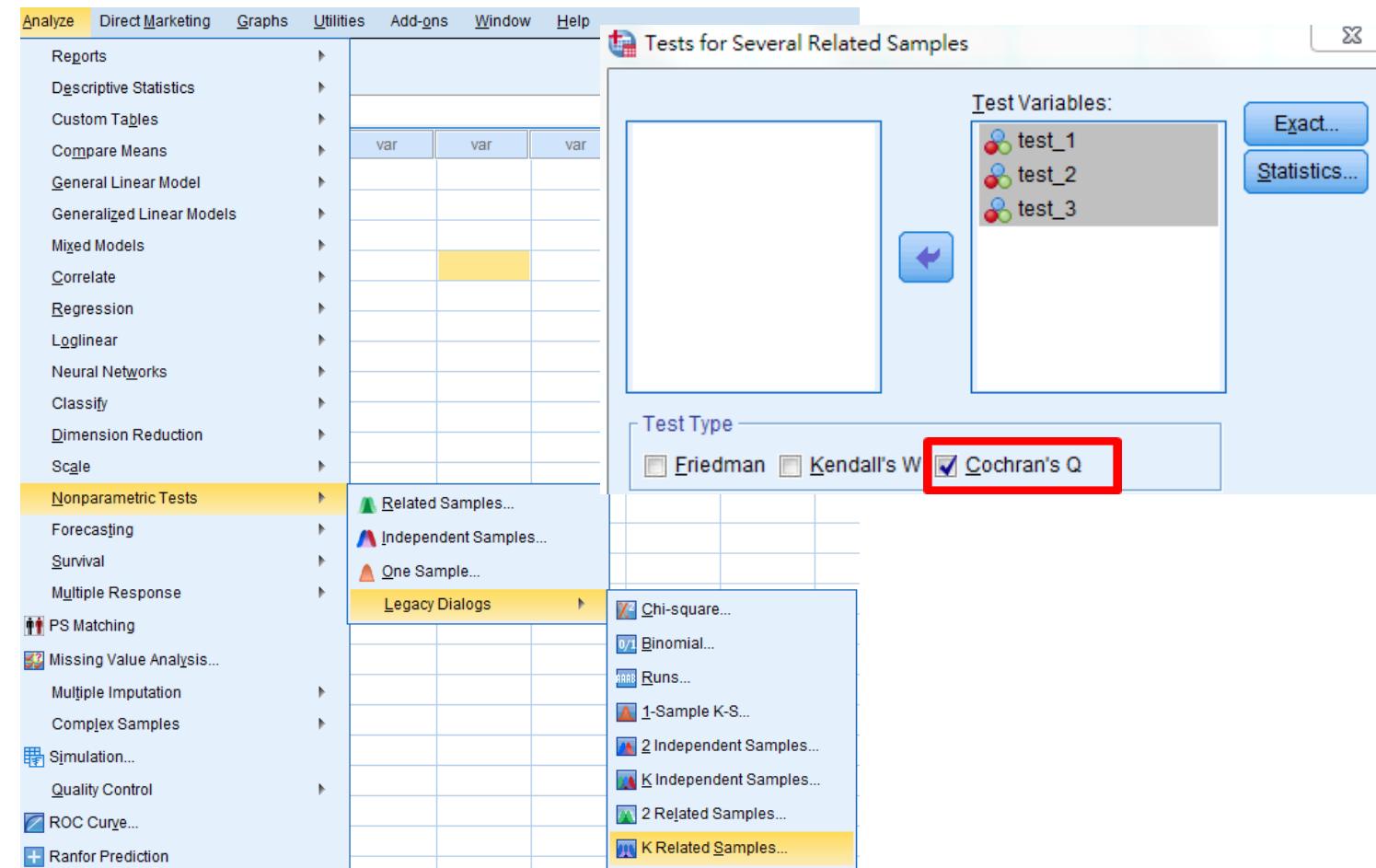
Cochran Q test dataset

- 同一組個案，測量三次時間點(類別型資料--二分類)
- 0: 無抽菸; 1: 有抽菸

no	test_1	test_2	test_3
1	0	1	1
2	0	1	1
3	1	1	1
4	1	0	1
5	0	0	0
6	1	1	1
7	1	1	1
8	1	0	1
9	1	1	1
10	0	0	1
11	1	1	1
12	1	1	1
13	1	0	1
14	1	0	0
15	0	0	1

Cochran Q test analysis

分析>無母數檢定>歷史對話記錄>K個相關樣本



Cochran Q test output

Cochran Test

Frequencies

	Value	
	0	1
test_1	5	10
test_2	7	8
test_3	2	13

Test Statistics

N	15
Cochran's Q	4.750 ^a
df	2
Asymp. Sig.	.093

a. 0 is treated as a success.

Table.

	Test1 (n=15)	Test2 (n=15)	Test3 (n=15)	p value
Smoking				0.093
No	5 (33.3%)	7 (46.7%)	2 (13.3%)	
Yes	10 (66.7%)	8 (53.3%)	13 (86.7%)	

Cochran Q test.

Paired t test

- 針對常態分布的連續數值或是大樣本的資料做平均值的比較
- 用於前後測比較。



ORIGINAL RESEARCH
published: 11 February 2022
doi: 10.3389/fmed.2021.698728



Subtypes of Premorbid Metabolic Syndrome and Associated Clinical Outcomes in Older Adults

Chu-Sheng Lin^{1,2,3,4}, Wei-Ju Lee^{4,5,6}, Shih-Yi Lin², Hui-Ping Lin⁷, Ran-Chou Chen^{5,7}, Chi-Hung Lin^{4,5,8} and Liang-Kung Chen^{4,5,9,10*}

TABLE 1 | Demographic data of study participants in 2014 and 2016.

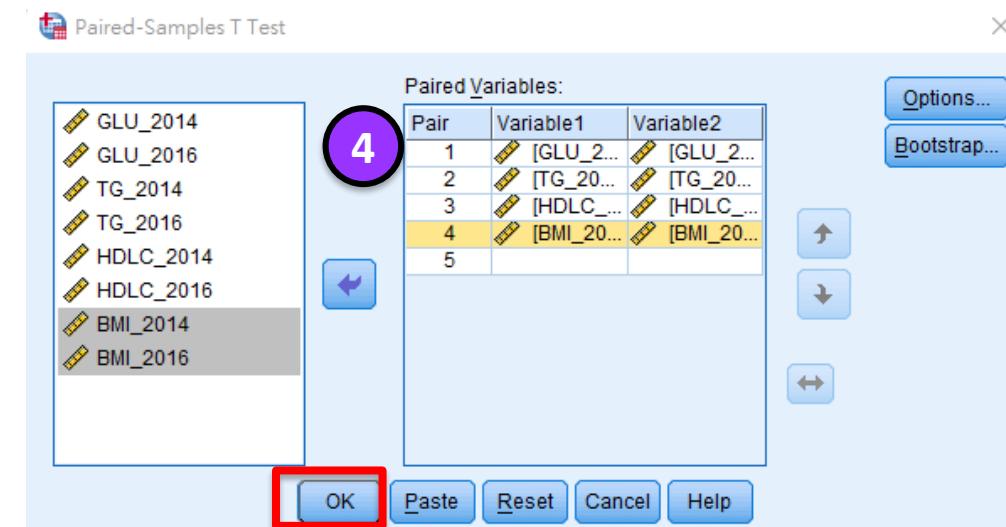
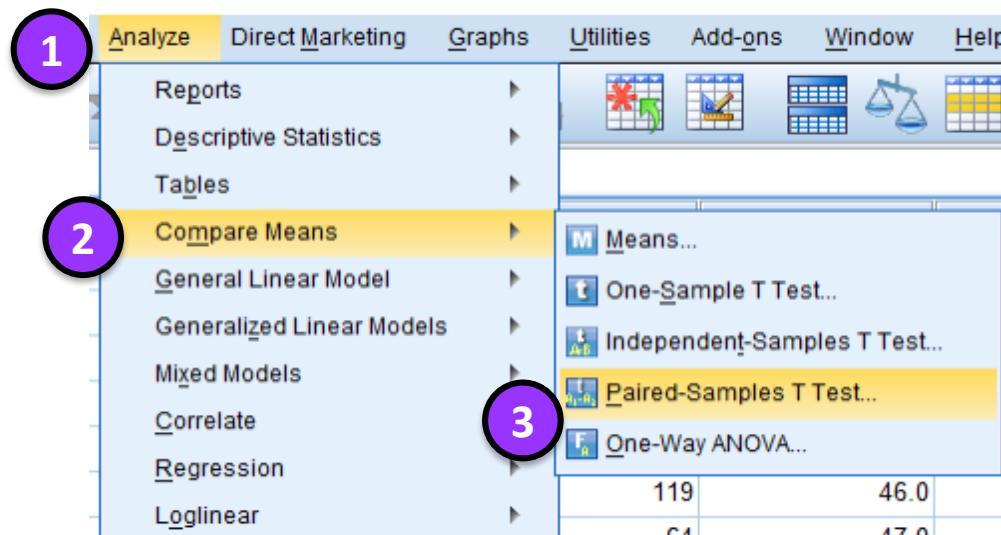
	2014 (n = 4,537)		2016 (n = 4,537)		p-value
	n	%	n	%	
Age	71.75	±5.93	73.75	±5.93	–
Gender					1.000
Male	2,207	(48.6%)	2,207	(48.6%)	
Female	2,330	(51.4%)	2,330	(51.4%)	
Smoking	309	(6.8%)	297	(6.5%)	0.119
Drinking	547	(12.1%)	581	(12.8%)	0.862
Exercise	3,237	(71.3%)	3,462	(76.3%)	0.930
BMI	24.49	±3.47	24.41	±3.42	0.004**
Height (cm)	157.67	±8.15	157.59	±8.12	0.063
Weight (kg)	60.99	±10.38	60.78	±10.47	<0.001**
Waist (cm)	83.20	±9.72	84.59	±9.66	<0.001**
SBP (mmHg)	133.91	±18.19	134.63	±18.52	0.007**
DBP (mmHg)	78.14	±10.91	77.21	±11.26	<0.001**
Pulse pressure (mmHg)	55.77	±13.82	57.42	±13.74	<0.001**
Fasting plasma glucose (mg/dl)	104.02	±21.92	104.89	±25.17	0.009**
Triglycerides (mg/dl)	117.03	±66.71	114.85	±67.83	0.024*
HDL cholesterol (mg/dl)	55.61	±15.66	55.60	±15.78	0.922

Paired t test dataset

caseno	GLU_2014	GLU_2016	TG_2014	TG_2016	HDLC_2014	HDLC_2016	BMI_2014	BMI_2016
1	91	91	245	180	34.0	42.0	27.68	27.27
2	103	224	89	104	45.0	35.0	19.15	20.16
3	92	96	96	56	54.0	47.0	20.94	23.57
4	97	96	123	92	57.0	57.0	25.54	25.57
5	88	307	125	106	42.0	87.0	18.97	17.10
6	94	94	101	119	46.0	47.0	21.50	21.79
7	92	91	49	64	47.0	50.0	25.67	26.58
8	151	89	83	69	39.0	47.0	24.61	26.37
9	97	97	120	59	42.0	44.0	23.95	24.31
10	149	156	142	128	50.0	50.0	25.54	23.84
11	113	124	337	104	58.0	80.0	23.28	23.74
12	99	92	114	76	67.0	60.0	20.20	20.57

Paired t test analysis

分析>比較平均數法>成對樣本T檢定



Paired T-test SPSS output

Paired Samples Statistics

	Mean	N	Std. Deviation	Std. Error Mean
Pair 1	GLU_2014	104.02	4537	21.923
	GLU_2016	104.89	4537	.325
Pair 2	TG_2014	117.03	4537	25.166
	TG_2016	114.85	4537	.990
Pair 3	HDLC_2014	55.615	4537	66.712
	HDLC_2016	55.598	4537	1.007
Pair 4	BMI_2014	24.4877	4537	15.6628
	BMI_2016	24.4141	4537	.2325
				.2343
				.05148
				.05076

在Glucose的部分，
2014和2016有統計差異
2016平均高於2014 ($p=0.009$)

Paired Samples Test

	Paired Differences					t	df	Sig. (2-tailed)			
	Mean	Std. Deviation	Std. Error Mean	95% Confidence Interval of the Difference							
				Lower	Upper						
Pair 1	GLU_2014 - GLU_2016	-.869	22.503	.334	-1.524	-.214	-2.600	4536	.009		
Pair 2	TG_2014 - TG_2016	2.186	65.013	.965	.294	4.078	2.265	4536	.024		
Pair 3	HDLC_2014 - HDLC_2016	.0164	11.3191	.1680	-.3131	.3458	.098	4536	.922		
Pair 4	BMI_2014 - BMI_2016	.07361	1.73773	.02580	.02303	.12418	2.853	4536	.004		

Wilcoxon signed-rank test

- 針對非常態分布的連續數值或小樣本，檢定兩組資料的中位數
- 用於前後測比較

Clinical Rheumatology
<https://doi.org/10.1007/s10067-023-06735-0>

ORIGINAL ARTICLE



Potential alleviation of bone mineral density loss with Janus kinase inhibitors in rheumatoid arthritis

Yun-Wen Chen^{1,2} · Hsin-Hua Chen^{1,2,3,4,5,6,7} · Wen-Nan Huang^{1,3,4,8} · Jun-Peng Chen^{9,10} · Yi-Hsing Chen^{1,4}.
 Yi-Ming Chen^{1,3,4,10,11}

Table 2 Comparisons of the changes in BMD and T-score from the pretreatment to posttreatment assessment based on the RA treatment

JAKi		Before		After		p-value
BMD, L spine	1.02	(0.92,	1.20)	1.03	(0.88,	1.21)
BMD, left femoral neck	0.84	(0.76,	0.90)	0.84	(0.76,	0.92)
BMD, right femoral neck	0.87	(0.77,	0.92)	0.87	(0.75,	0.94)
T-score, L spine	-1.5	(-2.5,	-0.6)	-1.6	(-2.5,	-1.0)
T-score, left femoral neck	-1.5	(-2.0,	-1.0)	-1.7	(-2.6,	-1.1)
T-score, right femoral neck	-1.4	(-1.9,	-0.8)	-1.8	(-2.7,	-1.0)

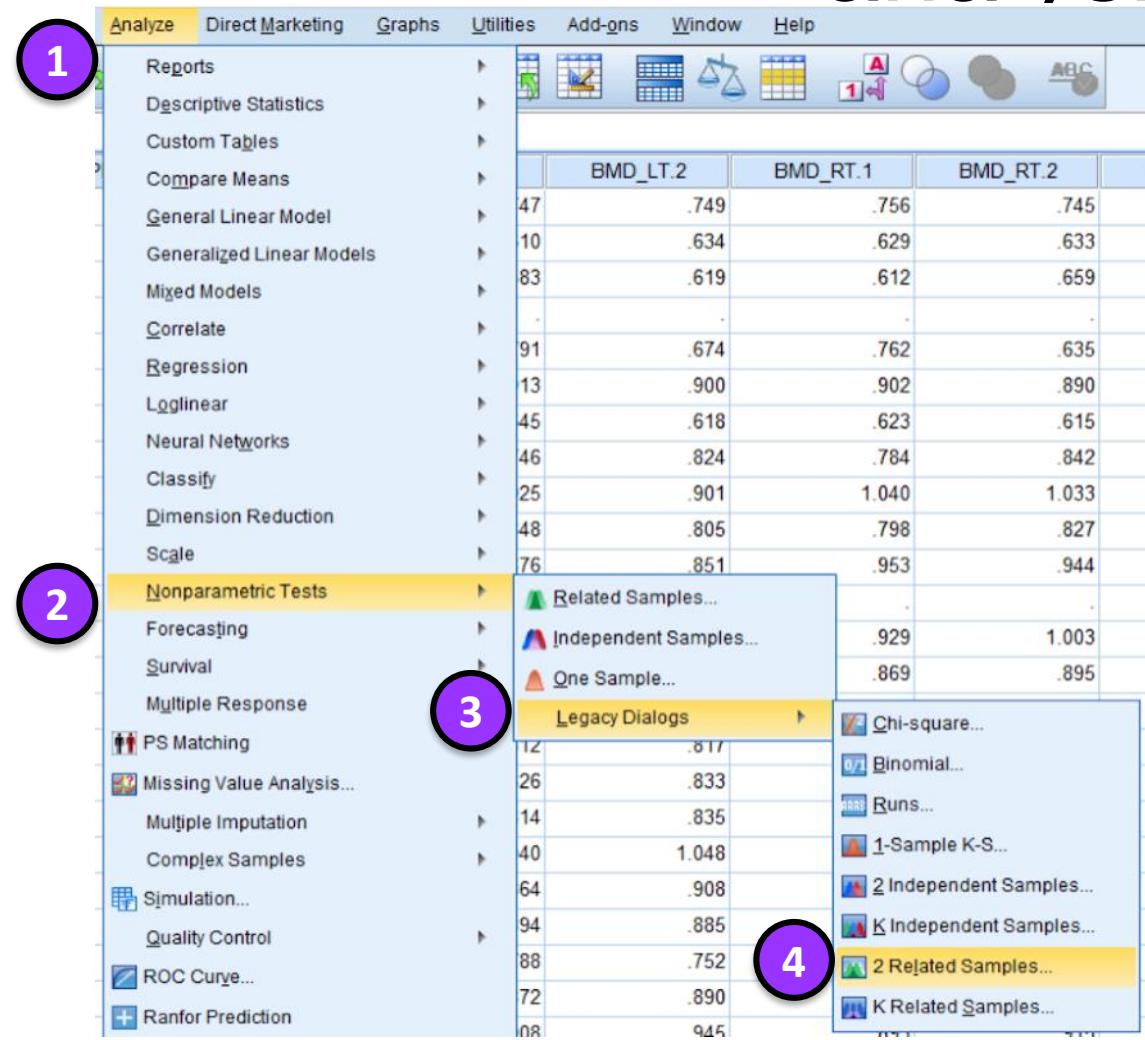
Wilcoxon signed-ranks test. * $p < 0.05$, ** $p < 0.01$

Wilcoxon signed-rank test dataset

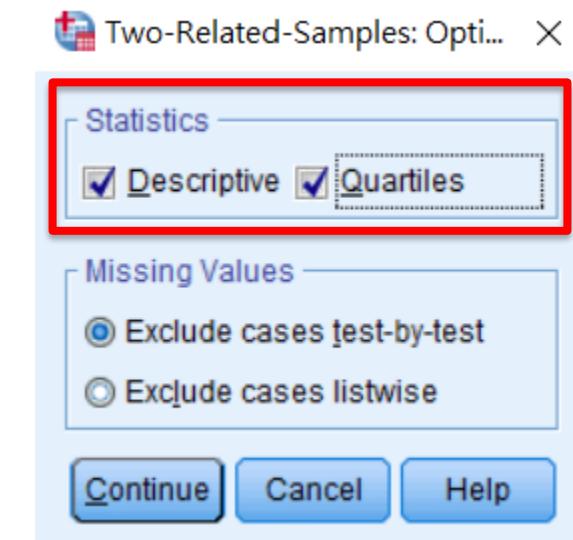
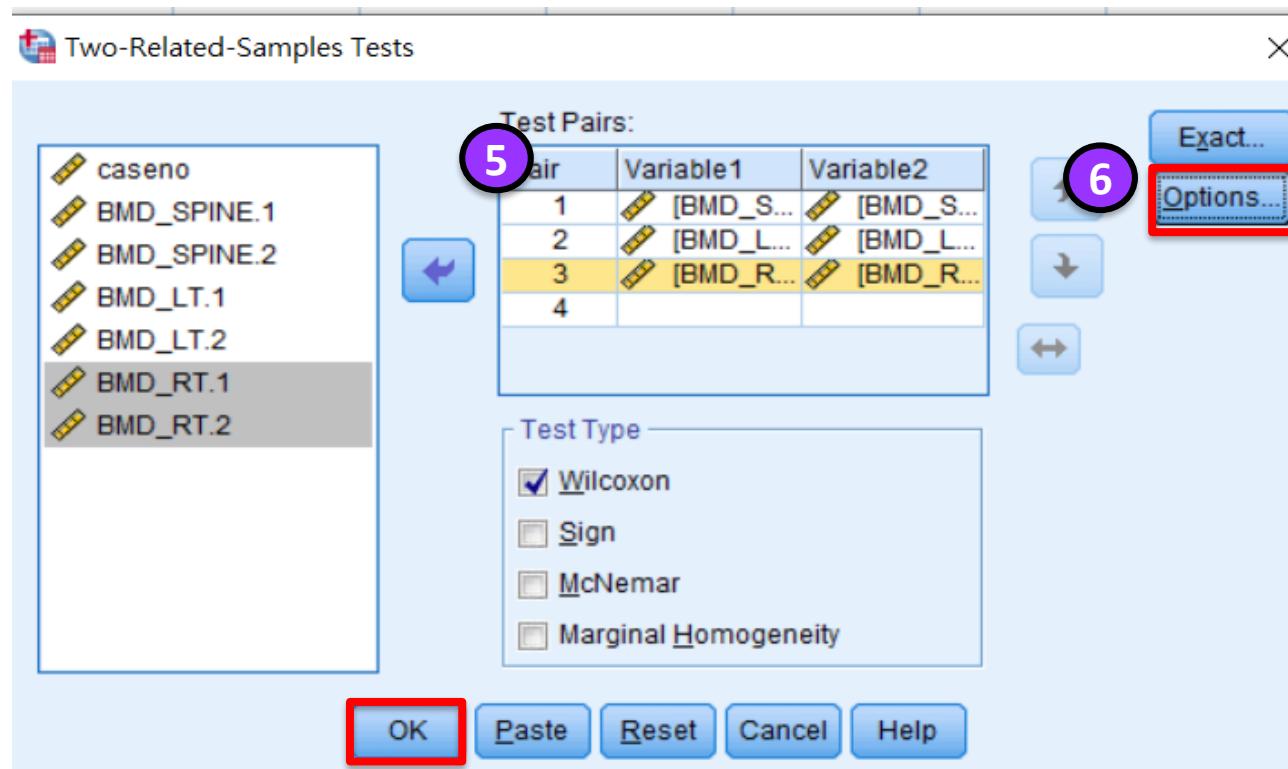
caseno	BMD_SPINE.1	BMD_SPINE.2	BMD_LT.1	BMD_LT.2	BMD_RT.1	BMD_RT.2
1	.805	.788	.747	.749	.756	.745
2	.866	.900	.610	.634	.629	.633
3	.804	.868	.583	.619	.612	.659
4	.714	.839
5	.973	.813	.791	.674	.762	.635
6	.	.	.913	.900	.902	.890
7	.823	.798	.645	.618	.623	.615
8	1.151	1.232	.746	.824	.784	.842
9	1.086	.	.925	.901	1.040	1.033
10	.989	.866	.848	.805	.798	.827
11	1.169	1.078	.876	.851	.953	.944
12	.899	.881
13	1.232	1.199	.888	.982	.929	1.003
14	1.020	.974	.881	.918	.869	.895
15	1.081	1.030	.772	.783	.796	.782

Wilcoxon signed-rank SPSS analysis-1

分析>無母數檢定>歷史對話記錄
>2個相關樣本



Wilcoxon signed-rank SPSS analysis-2



Wilcoxon signed-rank SPSS output

Descriptive Statistics

	N	Mean	Std. Deviation	Minimum	Maximum	Percentiles		
						25th	50th (Median)	75th
BMD_SPINE.1	28	1.05589	.185310	.714	1.478	.93050	1.01900	1.19175
BMD_LT.1	28	.82861	.131799	.536	1.097	.76075	.83700	.90450
BMD_RT.1	28	.84746	.141434	.579	1.187	.76650	.87050	.92300
BMD_SPINE.2	28	1.04739	.196029	.725	1.463	.87125	1.02250	1.20650
BMD_LT.2	28	.83682	.135854	.553	1.110	.75550	.84100	.91550
BMD_RT.2	28	.84875	.146636	.560	1.194	.75375	.87250	.93675

Test Statistics^a

	BMD_SPINE. 2 - BMD_SPINE. 1	BMD_LT.2 - BMD_LT.1	BMD_RT.2 - BMD_RT.1
Z	-.529 ^b	-1.378 ^b	-.216 ^b
Asymp. Sig. (2-tailed)	.597	.168	.829

在BMD spine的部分，使用JAK
Before和After無統計差異
After的中位數高於Before
(p=0.597)

a. Wilcoxon Signed Ranks Test

b. Based on negative ranks.

Friedman test

- 相同個案，測量三次時間點(數值型資料)
- 整體顯著可再做事後檢定(Dunn-Bonferroni)



The effect of foot reflexology massage on burn-specific pain anxiety and sleep quality and quantity of patients hospitalized in the burn intensive care unit (ICU)

Reza Alinia-najjar^a, Masoumeh Bagheri-Nesami^{b,c,*}, Seyed Afshin Shorofi^{b,d}, Seyed Nouraddin Mousavinasab^e, Kiarash Saatchi^f

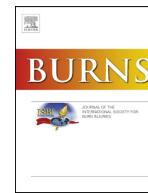
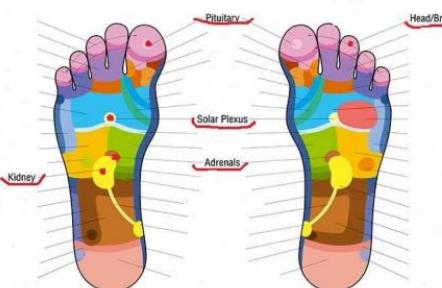
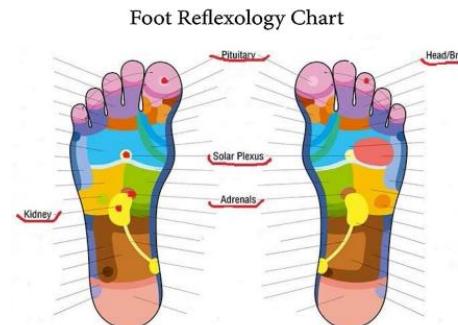


Table 3 – Comparison of median of duration of last night's sleep on the third, fourth, fifth and sixth days of hospitalization in two groups.

		Duration of last night's sleep	Third day (before the intervention)	Fourth day (before the intervention)	Fifth day (before the intervention)	Sixth day (before the intervention)	Friedman statistic and p-value
Intervention	Median	4.5 (3–7.5) ^a	6 (3–7.3)	6 (3–7.5)	8 (5.5–9.75)	X ² =60.184	
Control	Median	4.5 (3–7.5)	4.30 (3–7.3)	4.5 (3–7.5)	4.5 (3–7)	p<0.001	
Mann–Whitney U test and p-values		Z=0.000	Z=2.28	Z=3.24	Z=6.12	X ² =12.183	
		p=1.000	p=0.023	p<0.001	p<0.001	p=0.007	

^a (Min–Max).



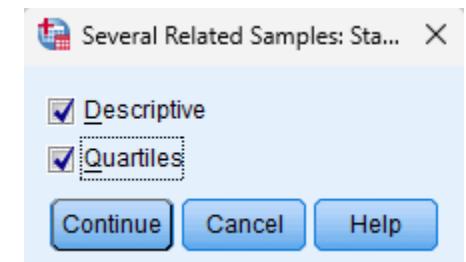
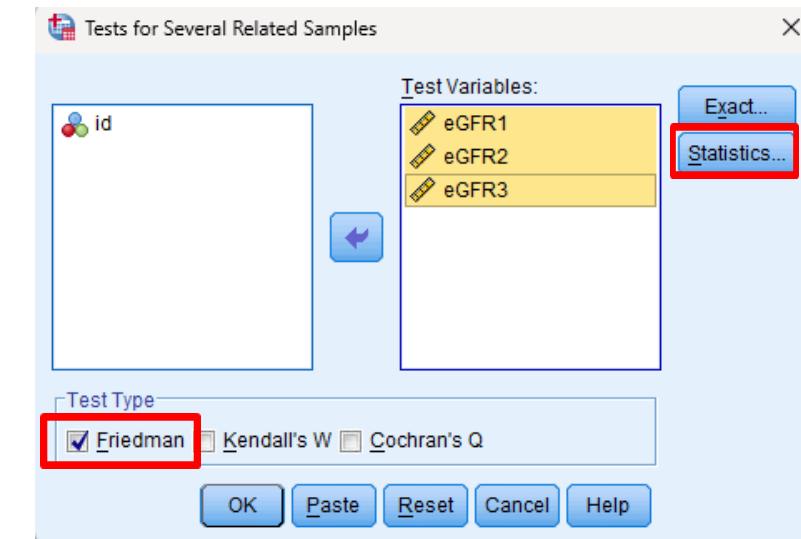
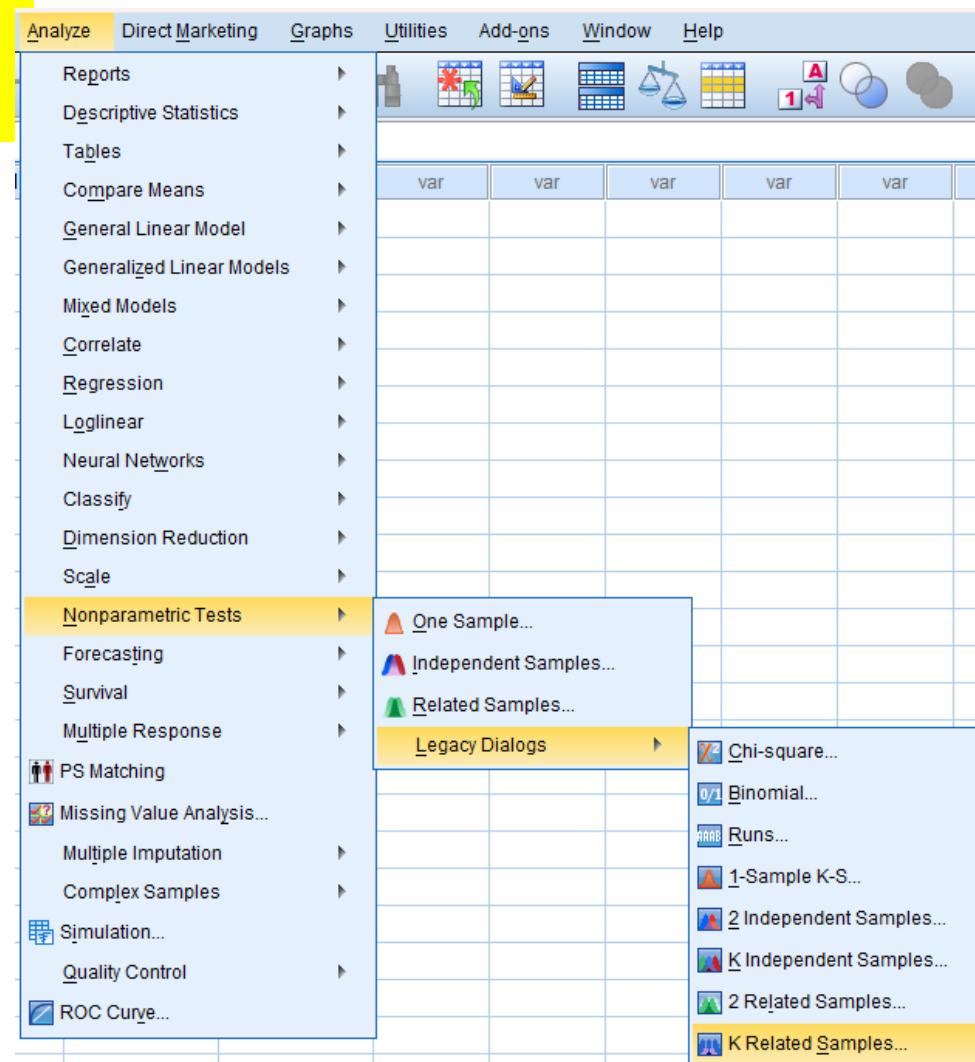
Friedman test dataset

- 同一組個案，測量三次時間點(數值型資料)
- 整體顯著可再做事後檢定(Dunn-Bonferroni)

id	eGFR1	eGFR2	eGFR3
1	60.6	80.2	13.6
2	85.9	14.4	48.1
3	56.7	60.6	78.9
4	47.3	32.7	89.1
5	42.7	59.7	97.9
6	100.4	71.1	60.7
7	42.8	80.7	38.9
8	80.2	75.2	32.1
9	14.4	73.4	48.1
10	60.6	21.2	80.3
11	38.9	50.6	30.7
12	32.1	62.6	59.1
13	48.1	56.8	15.1
14	80.3	58.2	36.5
15	30.7	55.4	74.5

Friedman test analysis

分析>無母數檢定>歷史對話記錄>K個相關樣本



Friedman test output

Descriptive Statistics

	N	Mean	Std. Deviation	Minimum	Maximum	Percentiles		
						25th	50th (Median)	75th
eGFR1	57	50.565	20.7201	14.4	100.4	32.590	48.090	67.365
eGFR2	57	59.949	17.1628	14.4	99.8	52.150	60.570	71.830
eGFR3	57	49.082	21.6933	13.6	97.9	32.070	42.910	69.035

Friedman Test

Ranks

	Mean Rank
eGFR1	1.84
eGFR2	2.33
eGFR3	1.82

Test Statistics^a

N	57
Chi-Square	9.509
df	2
Asymp. Sig.	.009

a. Friedman Test

Table. (N=57)

	Baseline	3 month	6 month	P value
eGFR	48.1(32.6-67.4)	60.6(52.2-71.8)	42.9(32.1-69.0)	0.009

Friedman test. Median (IQR)

Friedman test (Post-hoc)

Name	Type	Width	Decimals	Label	Values	Missing	Columns	Align	Measure
id	Numeric	2	0		None	None	10	Right	Nominal
eGFR1	Numeric	2	1		None	None	10	Right	Scale
eGFR2	Numeric	2	1		None	None	10	Right	Scale
eGFR3	Numeric	2	1		None	None	10	Right	Scale

分析>無母數檢定>相關樣本

The screenshot shows the SPSS menu bar at the top. Below it, the 'Analyze' menu is open, and the 'Nonparametric Tests' option is selected. Under 'Nonparametric Tests', the 'Related Samples...' option is highlighted. To the right, the 'Nonparametric Tests: Two or More Related Samples' dialog box is displayed. The 'Fields' tab is selected. In the 'Fields:' list, 'id' is listed under 'Sort: None'. In the 'Test Fields:' list, 'eGFR1', 'eGFR2', and 'eGFR3' are listed. A red box highlights the 'Fields' tab and the 'Test Fields:' list.

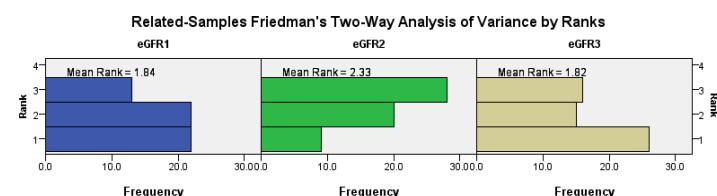
Friedman test (Post-hoc output)

1

Hypothesis Test Summary

Null Hypothesis	Test	Sig.	Decision
1 The distributions of eGFR1, eGFR2 and eGFR3 are the same.	Related-Samples Friedman's Two-Way Analysis of Variance by Ranks	.009	Reject the null hypothesis.

Asymptotic significances are displayed. The significance level is .05.



Total N	57
Test Statistic	9.509
Degrees of Freedom	2
Asymptotic Sig. (2-sided test)	.009

2

Related Samples Test View
Continuous Field Information
Pairwise Comparisons

Test: Friedman ▾ Field(s): eGFR1, eGFR2, eGFR3(Test 1) ▾ View: Related Samples Test View ▾

Each node shows the sample average rank.

Sample1-Sample2	Test Statistic	Std. Error	Std. Test Statistic	Sig.	Adj.Sig.
eGFR3-eGFR1	.018	.187	.094	.925	1.000
eGFR3-eGFR2	.509	.187	2.716	.007	.020
eGFR1-eGFR2	-.491	.187	-2.622	.009	.026

3

Each row tests the null hypothesis that the Sample 1 and Sample 2 distributions are the same.
Asymptotic significances (2-sided tests) are displayed. The significance level is .05.

Repeated measures ANOVA

- Outcome為數值型資料且需符合常態分佈
- 寬資料，有一次資料missing則排除此個案

Neurodegeneration

ORIGINAL RESEARCH

Effects of vitamin D supplementation on cognitive function and blood A β -related biomarkers in older adults with Alzheimer's disease: a randomised, double-blind, placebo-controlled trial

Jingya Jia,¹ Jing Hu,¹ Xiaoxu Huo,¹ Rujuan Miao,¹ Yanping Zhang,² Fei Ma  ¹

Table 2 The level of biochemical parameters at baseline, 6 months and 12 months between the two groups

Items	groups	Cases (n)	Treatment time*			Repeated measures†		
			Baseline	6 months	12 months	Interaction effect, P	Time effect, P	Group effect, P
A β 40 (pg/mL)	Intervention	105	27.78±20.80	26.88±20.81	28.62±20.79	0.468	0.027	0.480
	Control	105	33.46±24.06	32.43±24.00	33.22±23.20			
A β 42 (pg/mL)	Intervention	105	34.39±21.71	32.33±21.70	30.50±21.69	<0.001	<0.001	0.987
	Control	105	33.28±21.26	33.76±21.24	33.19±20.51			
P τ 1 (pg/mL)	Intervention	105	48.78±44.39	48.89±44.39	48.89±44.39	0.951	0.727	0.272
	Control	105	44.51±37.92	44.63±37.92	44.74±37.65			
P τ 2 (pg/mL)	Intervention	105	10.43±19.02	10.57±19.02	10.69±19.03	0.142	<0.001	0.133
	Control	105	11.01±17.44	11.17±17.44	11.55±17.53			
P τ 1mRNA	Intervention	105	28.85±2.59	30.19±2.61	29.77±2.59	0.040	<0.001	0.070
	Control	105	28.89±2.58	30.41±2.58	29.86±2.60			
P τ 2mRNA	Intervention	105	4.00±0.43	4.34±0.53	4.17±0.53	0.121	<0.001	0.409
	Control	105	4.18±0.39	4.60±0.61	4.34±0.38			
APP	Intervention	105	44.75±39.77	41.69±39.73	39.96±39.59	<0.001	<0.001	<0.001
	Control	105	44.71±33.20	42.56±33.20	45.56±33.24			
APPmRNA	Intervention	105	32.59±1.53	30.28±1.50	27.96±1.54	<0.001	<0.001	<0.001
	Control	105	32.41±1.41	31.44±1.47	31.55±1.38			
BACE1 (μ g/mL)	Intervention	105	300.83±11.96	298.14±11.97	296.23±12.01	<0.001	<0.001	<0.001
	Control	105	299.94±12.05	297.76±12.06	300.62±11.84			
BACE1mRNA	Intervention	105	27.59±1.29	25.29±1.55	23.48±1.91	<0.001	<0.001	<0.001
	Control	105	27.24±1.32	26.09±1.31	27.82±1.27			
25-D (ng/mL)	Intervention	105	18.82±2.91	20.90±3.12	22.77±3.41	<0.001	<0.001	<0.001
	Control	105	19.44±2.81	19.18±2.81	19.08±2.84			
1,25-D (ng/mL)	Intervention	105	30.37±2.60	31.79±2.57	33.61±2.77	<0.001	<0.001	<0.001
	Control	105	30.22±2.67	29.01±2.67	30.71±2.64			

*Presented as mean±SD.

†P value for group (intervention vs control) derived from analysis of covariance adjusted for respective baseline value and for age, gender and education. A β , amyloid beta; APP, A β protein precursor; BACE1, β -secretase 1; 25-D, 25-hydroxy vitamin D; 1,25-D, 1,25-dihydroxy vitamin D; PS, presenilin.

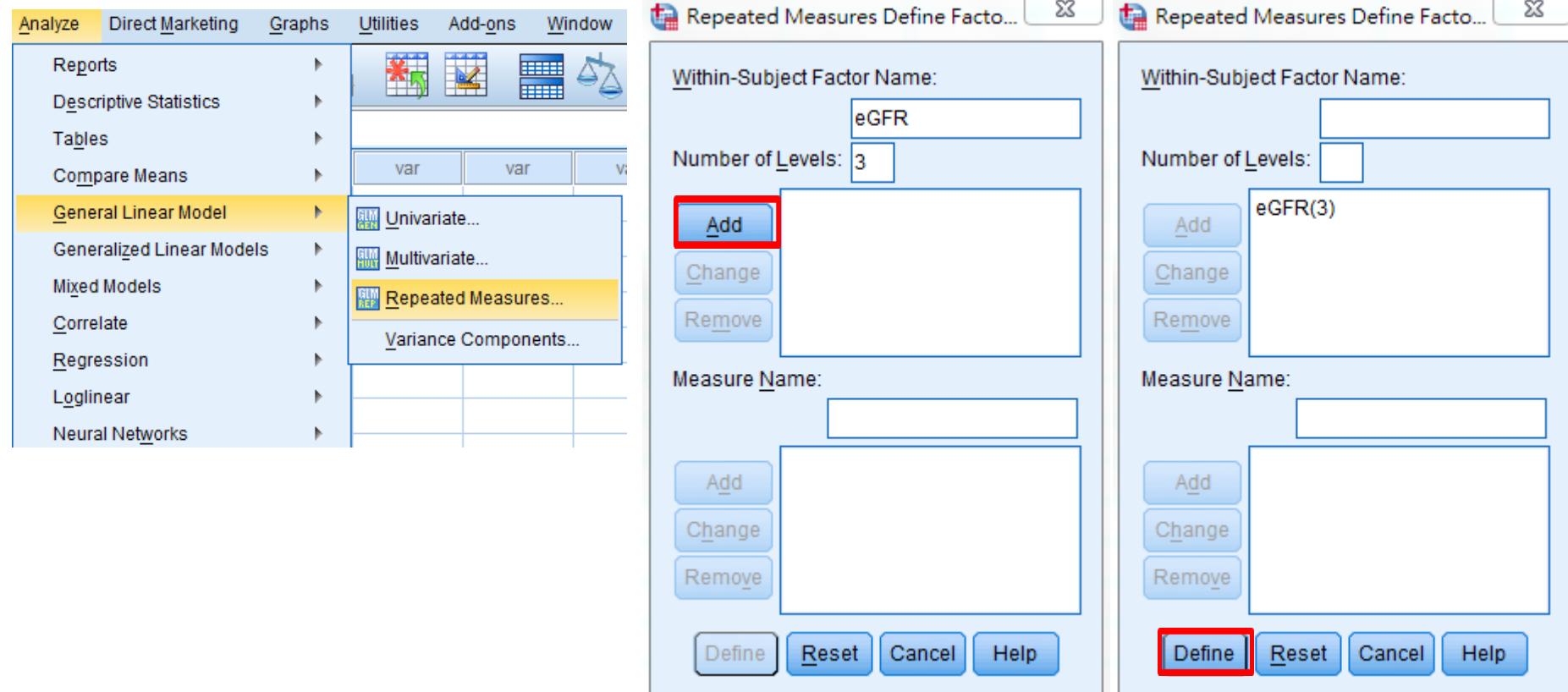
Repeated measures ANOVA dataset

- 兩組測量三個時間點(數值型資料)
- 整體顯著可再做事後檢定

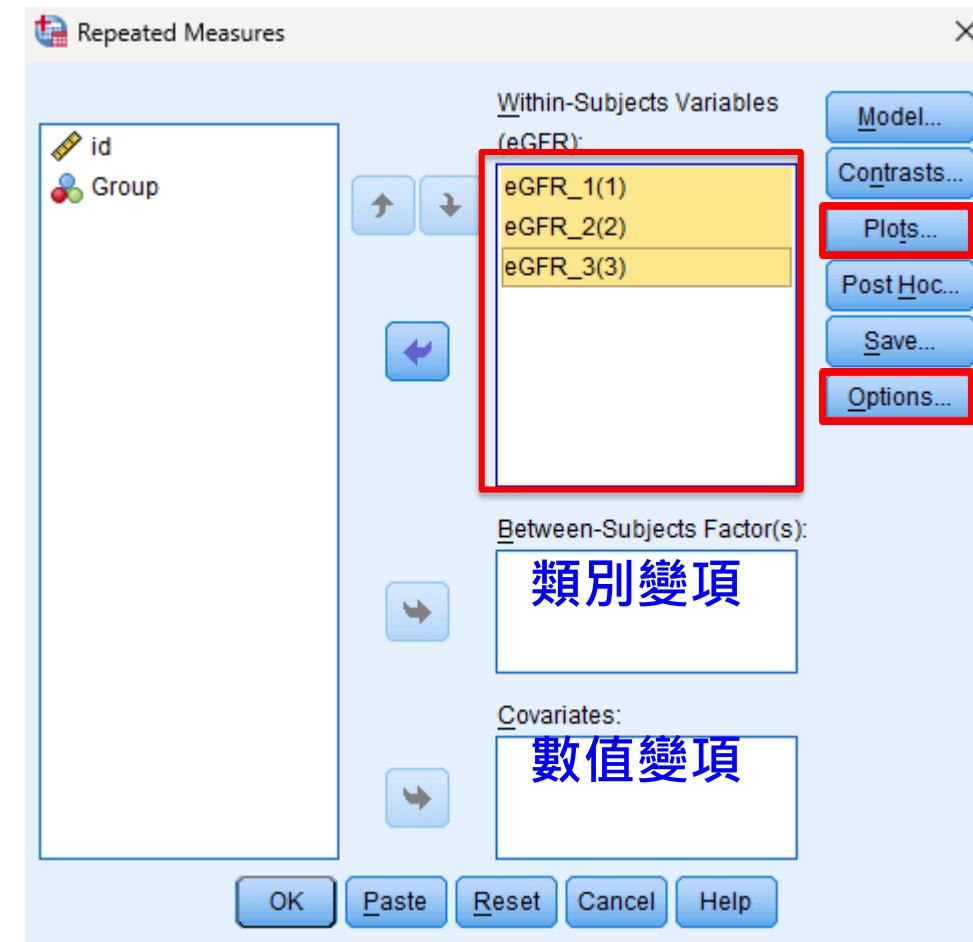
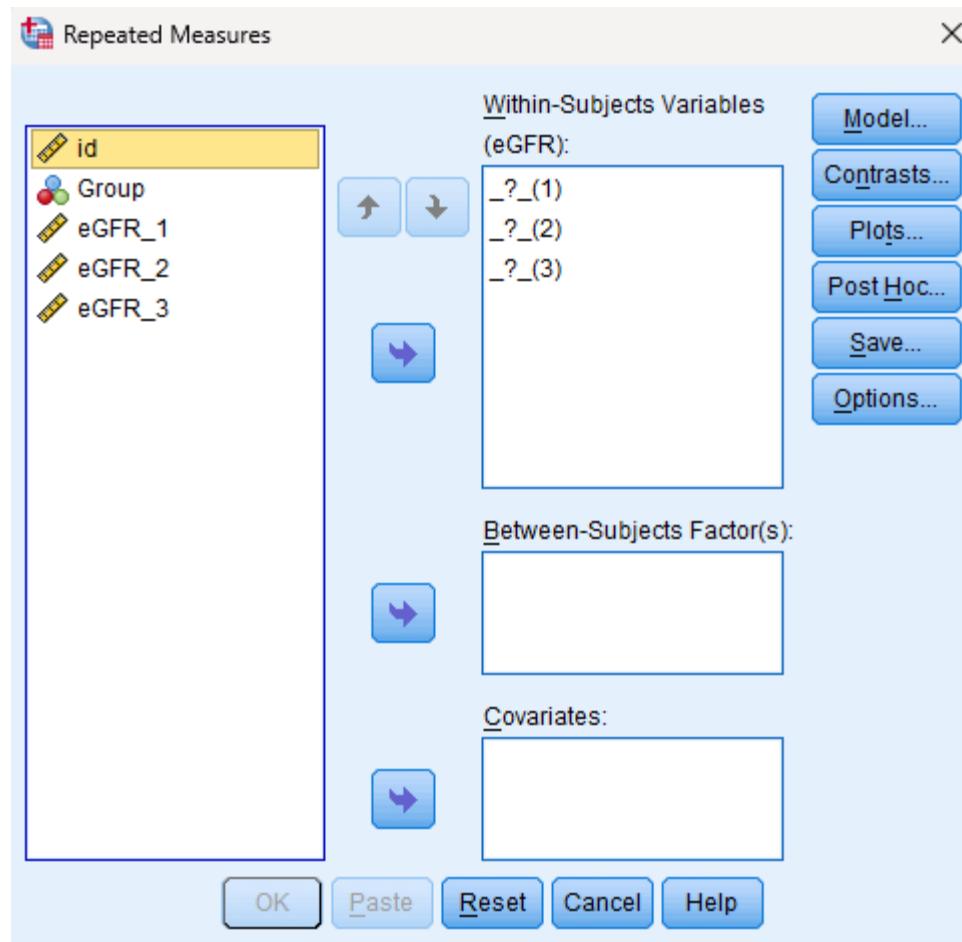
id	Group	eGFR_1	eGFR_2	eGFR_3
1	1	51.58	53.66	53.36
2	1	3.25	3.59	3.29
3	1	4.63	3.46	3.16
4	1	6.35	9.14	8.84
5	1	6.94	15.11	14.81
6	1	11.74	35.74	35.44
7	1	13.61	14.83	14.53
8	1	14.03	51.58	51.28
9	1	14.38	22.64	22.34
10	1	14.70	12.23	11.93
11	1	16.97	25.45	25.15
12	1	19.99	38.94	38.64
13	1	21.23	31.18	30.88
14	1	21.99	32.92	32.62
15	0	26.60	30.72	30.42

Repeated measures ANOVA analysis-1

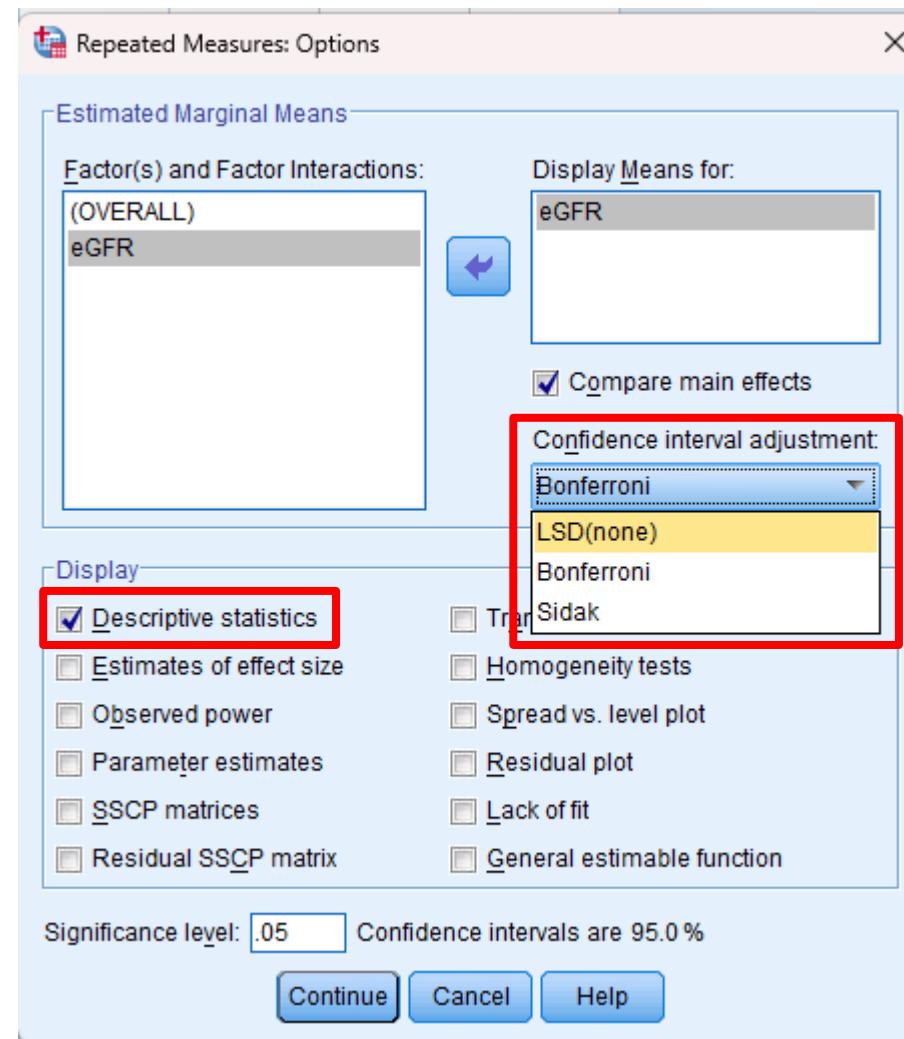
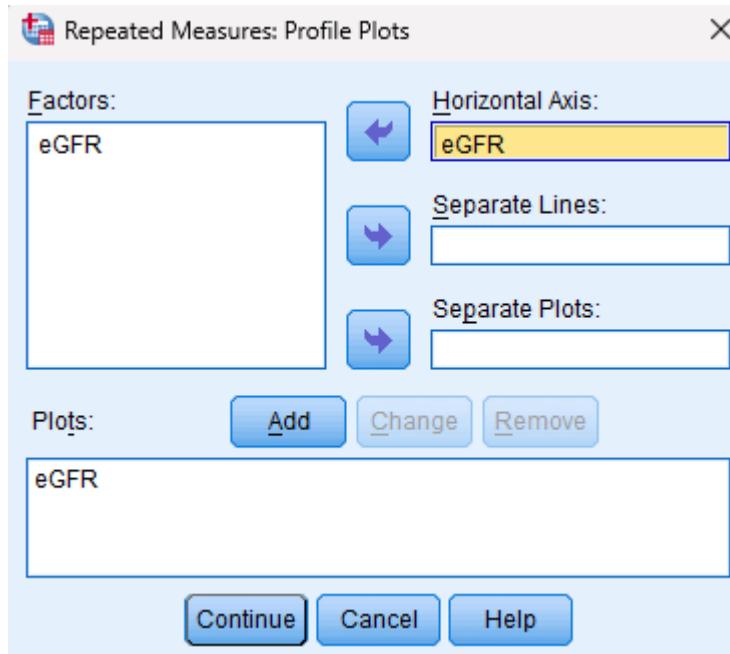
分析>一般線性模式>重複測量



Repeated measures ANOVA analysis-2



Repeated measures ANOVA analysis-3



Repeated measures ANOVA output

Descriptive Statistics

	Mean	Std. Deviation	N
eGFR_1	59.3108	23.02112	1127
eGFR_2	57.5311	23.78461	1127
eGFR_3	57.2311	23.78461	1127

Mauchly's Test of Sphericity^a

Measure: MEASURE_1

Within Subjects Effect	Mauchly's W	Approx. Chi-Square	df	Sig.	Epsilon ^b		
					Greenhouse-Geisser	Huynh-Feldt	Lower-bound
eGFR	.000	36007.760	2	.000	.500	.500	.500

Tests the null hypothesis that the error covariance matrix of the orthonormalized transformed dependent variables is proportional to an identity matrix.

a. Design: Intercept

Within Subjects Design: eGFR

P<0.05 代表資料未符合球形假設

b. May be used to adjust the degrees of freedom for the averaged tests of significance. Corrected tests are displayed in the Tests of Within-Subjects Effects table.

Tests of Within-Subjects Effects

Measure: MEASURE_1 符合球

Source	形假設	Type III Sum of Squares	df	Mean Square	F	Sig.
eGFR	Sphericity Assumed	2848.636	2	1424.318	20.420	.000
	Greenhouse-Geisser	2848.636	1.000	2848.636	20.420	.000
	Huynh-Feldt	2848.636	1.000	2848.636	20.420	.000
	Lower-bound	2848.636	1.000	2848.636	20.420	.000
Error(eGFR)	Sphericity Assumed	157081.715	2252	69.752		
	Greenhouse-Geisser	157081.715	1126.000	139.504		
	Huynh-Feldt	157081.715	1126.000	139.504		
	Lower-bound	157081.715	1126.000	139.504		

不符合球
形假設

Pairwise Comparisons

Measure: MEASURE_1

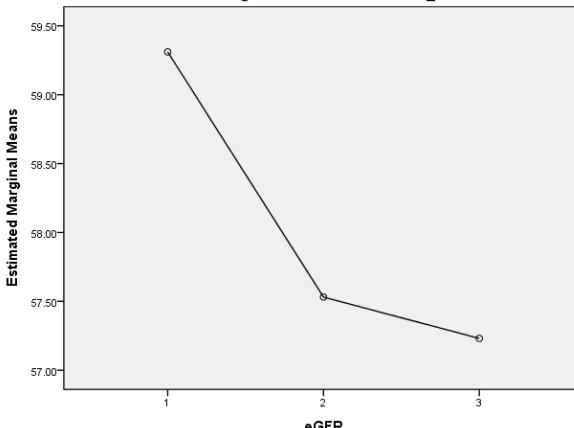
(I) eGFR	(J) eGFR	Mean Difference (I-J)	Std. Error	Sig. ^b	95% Confidence Interval for Difference ^b	
					Lower Bound	Upper Bound
1	2	1.780 [*]	.431	.000	.747	2.813
	3	2.080 [*]	.431	.000	1.047	3.113
2	1	-1.780 [*]	.431	.000	-2.813	-.747
	3	.300 [*]	.000	.000	.300	.300
3	1	-2.080 [*]	.431	.000	-3.113	-1.047
	2	-.300 [*]	.000	.000	-.300	-.300

Based on estimated marginal means

*. The mean difference is significant at the .05 level.

b. Adjustment for multiple comparisons: Bonferroni.

Estimated Marginal Means of MEASURE_1



Repeated measures ANOVA analysis

The image shows two overlapping dialog boxes from SPSS:

Repeated Measures dialog box (left):

- Within-Subjects Variables (eGFR):** eGFR_1(1), eGFR_2(2), eGFR_3(3)
- Between-Subjects Factor(s):** Group
- Covariates:** None
- Buttons:** Model..., Contrasts..., Plots..., Post Hoc..., Save..., Options..., OK, Paste, Reset, Cancel, Help.

Repeated Measures: Profile Plots dialog box (right):

- Factors:** Group, eGFR
- Horizontal Axis:** eGFR
- Separate Lines:** Group
- Separate Plots:** None
- Plots:** eGFR*Group
- Buttons:** Add (highlighted with a red box), Change, Remove, Continue, Cancel, Help.

Repeated measures ANOVA output

Descriptive Statistics

Group	Mean	Std. Deviation	N
eGFR_1	0	66.1851	19.40228
	1	58.3516	23.32970
	Total	59.3108	23.02112
eGFR_2	0	63.7180	19.50203
	1	56.6678	24.20522
	Total	57.5311	23.78461
eGFR_3	0	63.4180	19.50203
	1	56.3678	24.20522
	Total	57.2311	23.78461

Mauchly's Test of Sphericity^a

Measure: MEASURE_1

Within Subjects Effect	Mauchly's W	Approx. Chi-Square	df	Sig.	Epsilon ^b		
					Greenhouse-Geisser	Huynh-Feldt	Lower-bound
eGFR	.000	36298.754	2	.000	.500	.500	.500

Tests the null hypothesis that the error covariance matrix of the orthonormalized transformed dependent variables is proportional to an identity matrix.

a. Design: Intercept + Group
Within Subjects Design: eGFR

b. May be used to adjust the degrees of freedom for the averaged tests of significance. Corrected tests are displayed in the Tests of Within-Subjects Effects table.

Tests of Within-Subjects Effects

Measure: MEASURE_1

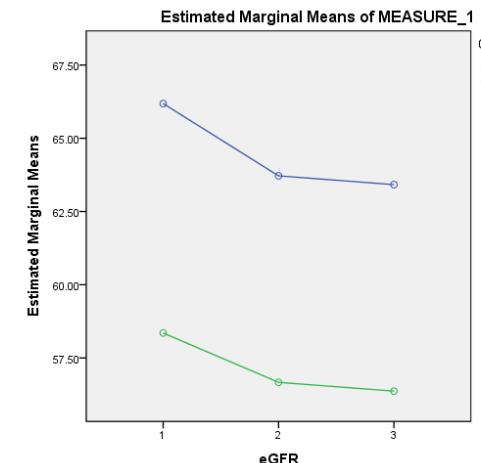
Source	Type III Sum of Squares	df	Mean Square	F	Sig.
eGFR	Sphericity Assumed	1621.180	810.590	11.614	.000
	Greenhouse-Geisser	1621.180	1621.180	11.614	.001
	Huynh-Feldt	1621.180	1619.739	11.614	.001
	Lower-bound	1621.180	1621.180	11.614	.001
eGFR * Group	Sphericity Assumed	49.520	24.760	.355	.701
	Greenhouse-Geisser	49.520	49.520	.355	.552
	Huynh-Feldt	49.520	49.476	.355	.552
	Lower-bound	49.520	49.520	.355	.552
Error(eGFR)	Sphericity Assumed	157032.196	69.792		
	Greenhouse-Geisser	157032.196	1125.000	139.584	
	Huynh-Feldt	157032.196	1126.001	139.460	
	Lower-bound	157032.196	1125.000	139.584	

Tests of Between-Subjects Effects

Measure: MEASURE_1

Transformed Variable: Average

Source	Type III Sum of Squares	df	Mean Square	F	Sig.
Intercept	5369346.712	1	5369346.712	3565.368	.000
Group	19420.818	1	19420.818	12.896	.000
Error	1694219.138	1125	1505.973		



Linear Mixed Model / Generalized Estimating Equation

- Linear Mixed Model
 - Outcome為數值型資料且需符合常態分佈
 - 資料型態為長資料，可容許遺漏值的存在
- GEE
 - 不需符合常態分佈假設，為半母數方法(semi-parametric)
 - Outcome可為連續、類別、順序或計數類型
 - 資料型態為長資料，可容許遺漏值的存在

GEE矩陣

- 獨立矩陣 (Independent)
- AR(1) 矩陣 (Auto-regressive first order)
- 可交換矩陣 (Exchangeable)
- 未結構化矩陣 (Unstructured)

獨立矩陣

	t_1	t_2	t_3	t_4	t_5
t_1	—				
t_2	0	—			
t_3	0	0	—		
t_4	0	0	0	—	
t_5	0	0	0	0	—

AR(1) 矩陣

	t_1	t_2	t_3	t_4	t_5
t_1	—				
t_2	ρ	—			
t_3	ρ^2	ρ	—		
t_4	ρ^3	ρ^2	ρ	—	
t_5	ρ^4	ρ^3	ρ^2	ρ	—

可交換矩陣

	t_1	t_2	t_3	t_4	t_5
t_1	—				
t_2	ρ	—			
t_3	ρ	ρ	—		
t_4	ρ	ρ	ρ	—	
t_5	ρ	ρ	ρ	ρ	—

未結構化矩陣

	t_1	t_2	t_3	t_4	t_5
t_1	—				
t_2	ρ_1	—			
t_3	ρ_2	ρ_5	—		
t_4	ρ_3	ρ_6	ρ_8	—	
t_5	ρ_4	ρ_7	ρ_9	ρ_{10}	—

GEE

Midlife overweight and obesity increase late-life dementia risk

A population-based twin study

W.L. Xu, MD, PhD
 A.R. Atti, MD, PhD
 M. Gatz, PhD
 N.L. Pedersen, PhD
 B. Johansson, PhD
 L. Fratiglioni, MD, PhD

ABSTRACT

Objective: The relation of overweight to dementia is controversial. We aimed to examine the association of midlife overweight and obesity with dementia, Alzheimer disease (AD), and vascular dementia (VaD) in late life, and to verify the hypothesis that genetic and early-life environmental factors contribute to the observed association.

Methods: From the Swedish Twin Registry, 8,534 twin individuals aged ≥ 65 (mean age 74.4) were assessed to detect dementia cases (DSM-IV criteria). Height and weight at midlife (mean age 43.1) were available in the Registry. Data were analyzed as follows: 1) unmatched case-

Table 2 Adjusted odds ratio (OR) and 95% confidence interval (CI) of dementia, Alzheimer disease, and vascular dementia related to midlife BMI (results from generalized estimating equation models)

Midlife BMI	No. of twins	All dementia			Alzheimer disease			Vascular dementia		
		No.	OR (95% CI) ^a	OR (95% CI) ^b	No.	OR (95% CI) ^a	OR (95% CI) ^b	No.	OR (95% CI) ^a	OR (95% CI) ^b
Continuous	8,534	464	1.09 (1.06-1.12)	1.06 (1.03-1.10)	232	1.09 (1.04-1.13)	1.06 (1.01-1.10)	74	1.14 (1.08-1.21)	1.11 (1.04-1.19)
Categorical										
<20	627	17	0.74 (0.44-1.25)	0.79 (0.45-1.38)	8	0.89 (0.64-1.23)	0.66 (0.31-1.41)	0	—	—
20-25	5,366	240	1 (Reference)	1 (Reference)	120	1 (Reference)	1 (Reference)	36	1 (Reference)	1 (Reference)
>25	2,541	207	1.50 (1.22-1.84)	1.80 (1.37-2.35)	104	1.52 (1.15-2.02)	1.98 (1.36-2.88)	38	1.62 (1.01-2.59)	1.35 (0.81-2.24)
25-30	2,297	177	1.37 (1.11-1.70)	1.71 (1.30-2.25)	90	1.41 (1.05-1.89)	1.91 (1.30-2.80)	31	1.39 (0.85-2.29)	1.17 (0.69-2.00)
>30	244	30	3.01 (1.95-4.64)	3.88 (2.12-7.11)	14	2.87 (1.57-5.26)	3.43 (1.49-7.90)	7	4.38 (1.89-10.14)	3.50 (1.36-8.99)

Abbreviations: BMI = body mass index; CI = confidence interval; OR = odds ratio.

^a Adjusted for age, sex, and education.

^b Adjusted for age, sex, education, diabetes, hypertension, stroke, and heart disease.

GEE

Summative Effects of Vascular Risk Factors on the Progression of Alzheimer Disease

Wei-Ju Lee, MD, PhD,^{*†‡§¶} Yi-Chu Liao, MD, PhD,^{†¶} Yen-Feng Wang, MD, PhD,^{†¶**}
Yung-Shuan Lin, MD,^{†***} Shuu-Jiun Wang, MD,^{†¶**} and Jong-Ling Fuh, MD,^{†¶**}

OBJECTIVES: To investigate the summative effects of vascular risk factors (VRFs) on the progression of Alzheimer disease (AD).

DESIGN: Longitudinal follow-up cohort study.

SETTING: AD patients from two teaching hospitals in Taiwan with 3-year follow-ups.

three or fewer VRFs (MMSE, $P = .009$; CDRSB, $P = .02$). Subsequent analyses of APOE ε4 carriers with more than three VRFs also showed their more rapid cognitive decline compared with patients without VRFs (MMSE, $P = .02$; CDRSB, $P = .001$) and patients with three or fewer VRFs (MMSE, $P = .009$; CDRSB, $P = .02$), but no significant difference was found in APOE ε4 noncarriers.

Vascular risk factors, No. (%)	
Coronary heart disease	63 (19.1)
Cardiac arrhythmia	30 (9.1)
Cerebrovascular disease	12 (3.6)
Hypertension	217 (65.8)
Diabetes mellitus	154 (46.7)
Obesity	18 (5.5)
Smoking	44 (13.3)
Physical inactivity	121 (36.7)

Table 2. Results of the generalized estimating equation analyzing the effect of 3-year MMSE changes in AD patients with and without APOE ε4 and different VRF indexes

Variable	All AD patients (n = 330)		AD patients with APOE ε4 (n = 129)		AD patients without APOE ε4 (n = 201)			
	β (95% CI)	P value ^a	β (95% CI)	P value ^a	β (95% CI)	P value ^a		
VRF groups								
>3 VRFs	-1.16 (-2.1 to -0.21)	.02	-1.99 (-3.62 to -0.36)	.02	-0.59 (-1.7 to 0.52)	.3		
3 VRFs	-0.36 (-1.24 to 0.51)	.42	-1.2 (-2.8 to 0.4)	.14	0.11 (-0.9 to 1.1)	.83		
2 VRFs	-0.15 (-0.92 to 0.61)	.69	-0.51 (-1.8 to 0.78)	.44	0.27 (-0.64 to 1.18)	.56		
1 VRF	-0.04 (-0.83 to 0.73)	.89	-0.48 (-1.76 to 0.81)	.47	0.3 (-0.68 to 1.28)	.55		
0 VRFs	Reference		Reference		Reference			
APOE ε4								
Carrier	-0.63 (-1.18 to -0.08)	.03						
Noncarrier	Reference							
Sex								
Female	0.35 (-0.15 to 0.85)	.17	0.64 (-0.21 to 1.5)	.14	0.19 (-0.42 to 0.79)	.55		
Male	Reference		Reference		Reference			
Age								
	-0.001 (-0.05 to 0.05)	.96	0.02 (-0.06 to 0.1)	.65	-0.01 (-0.07 to 0.05)	.7		
Education								
	-0.01 (-0.07 to 0.04)	.66	-0.06 (-0.15 to 0.03)	.2	<0.001 (-0.07 to 0.07)	.99		
Disease duration								
	0.001 (-0.004 to 0.01)	.76	0.005 (-0.003 to 0.01)	.2	-0.001 (-0.008 to 0.005)	.72		
Baseline MMSE								
	0.95 (0.9 to 0.99)	<.001	0.97 (0.9 to 1)	<.001	0.93 (0.87 to 1)	<.001		
Time								
	-1.17 (-1.4 to -1)	<.001	-1.34 (-1.65 to -1.03)	<.001	-1.08 (-1.3 to -0.86)	<.001		
Medication								
	-0.46 (-0.95 to 0.03)	.07	-0.85 (-1.77 to 0.07)	.07	-0.18 (-0.77 to 0.4)	.57		
Hospitalization rates								
	-0.24 (-0.66 to 0.17)	.25	-0.6 (-1.2 to 0.04)	.05	-0.18 (-0.71 to 0.34)	.49		

Abbreviations: AD, Alzheimer disease; APOE, apolipoprotein E; CI, confidence interval; MMSE, Mini-Mental State Examination; NMDA, N-methyl-D-aspartate; VRF, vascular risk factor.

^aP values were adjusted for APOE ε4 carrier status (in all AD patients), sex, age, years of education, disease duration, baseline MMSE score, time, use of medication (acetylcholinesterase inhibitors or NMDA receptor antagonists), and hospitalization rates.

GEE/LMM-資料轉置處理

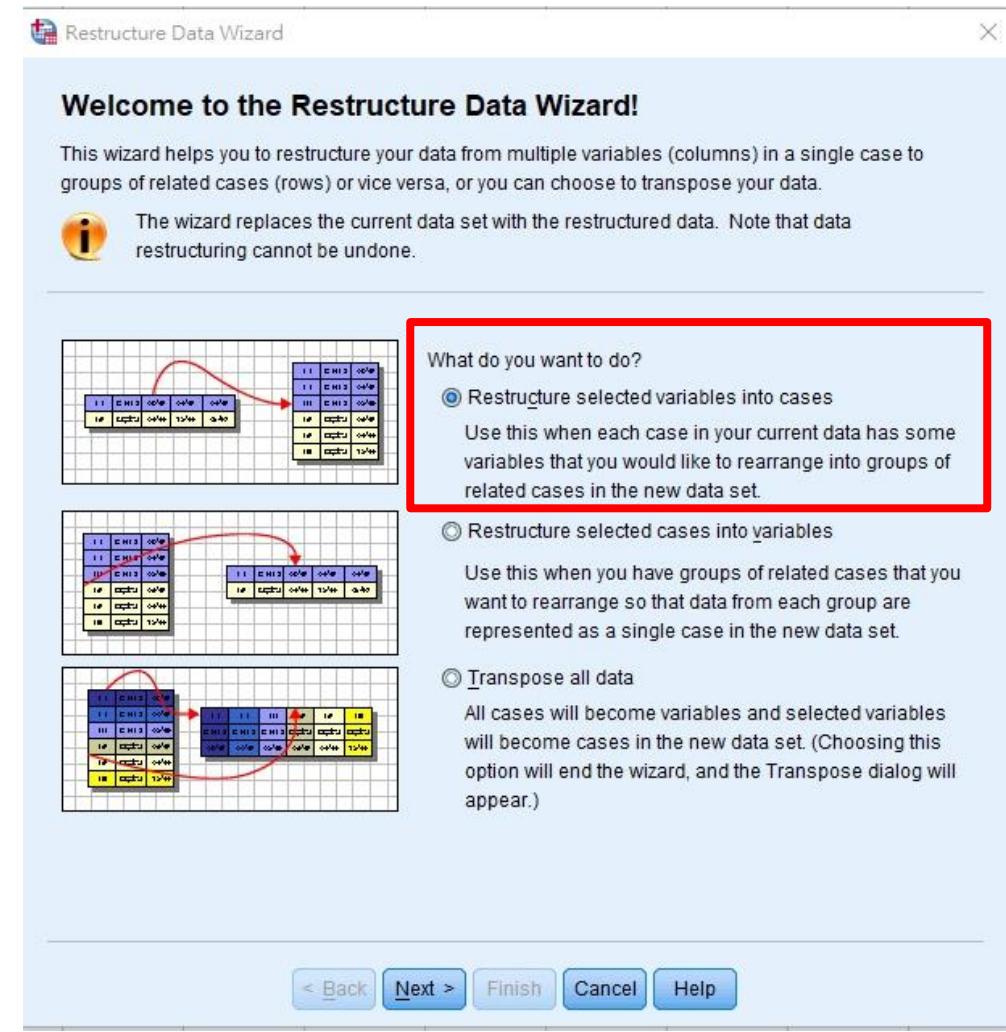
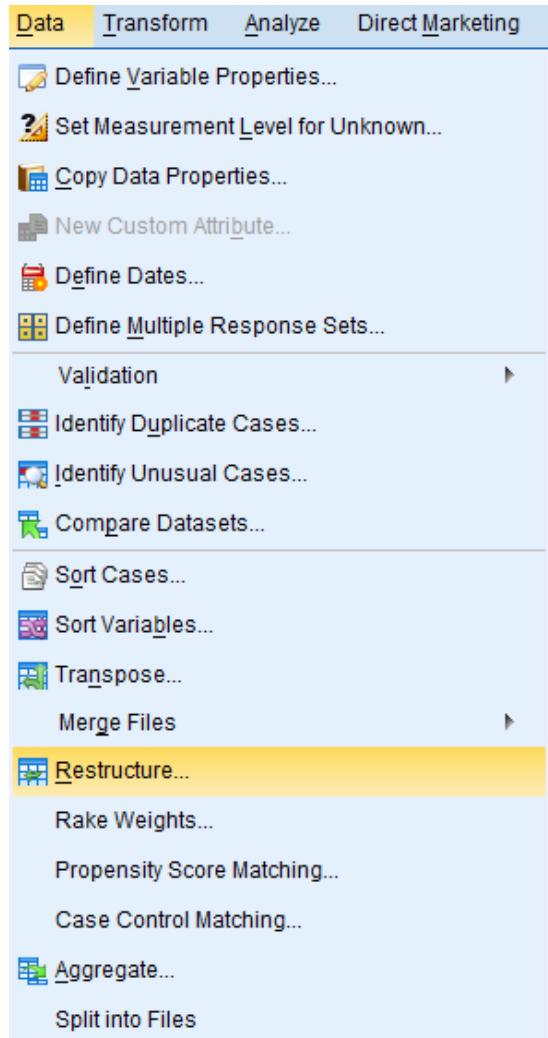
id	Gender	Treatment	Age	DepressionScale_1	DepressionScale_2	DepressionScale_3
1	2	1	29	4.0	32.0	13.0
2	2	1	32	29.0	14.0	8.0
3	1	1	32	19.0	27.0	9.0
4	2	2	35	19.0	29.0	33.0
5	1	2	31	20.0	27.0	9.0
6	2	1	33	8.0	12.0	12.0
7	1	2	34	20.0	2.0	19.0
8	1	1	34	23.0	6.0	23.0
9	2	1	29	9.0	.0	.0
10	1	1	34	4.0	33.0	10.0
11	2	1	25	24.0	15.0	10.0
12	2	2	34	10.0	31.0	31.0
13	1	2	26	26.0	27.0	8.0



id	Gender	Treatment	Age	Visit	DepressionScale
1	2	1	29	1	4.0
1	2	1	29	2	32.0
1	2	1	29	3	13.0
1	2	1	29	4	19.0
1	2	1	29	5	11.0
2	2	1	32	1	29.0
2	2	1	32	2	14.0
2	2	1	32	3	8.0
2	2	1	32	4	5.0
2	2	1	32	5	4.0
3	1	1	32	1	19.0
3	1	1	32	2	27.0
3	1	1	32	3	9.0
3	1	1	32	4	15.0

GEE/LMM-資料轉置處理

資料 > 重新架構



GEE/LMM-資料轉置處理

Variables to Cases: Number of Variable Groups

You have chosen to restructure selected variables into groups of related cases in the new file.

A group of related variables, called a variable group, represents measurements on one variable.

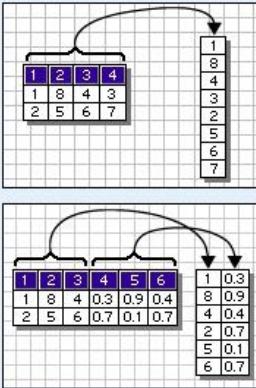
For example, the variable may be width. If it is recorded in three separate measurements, each one representing a different point in time—w1, w2, and w3, then the data are arranged in a group of variables.

If there is more than one variable in the file often it is also recorded in a variable group, for example height, recorded in h1, h2, and h3.

How many variable groups do you want to restructure?

One (for example, w1, w2, and w3)

More than one (for example, w1, w2, w3 and h1, h2, h3, etc.)
How Many?



< Back [Next >](#) [Finish](#) [Cancel](#) [Help](#)

Variables to Cases: Select Variables

For each variable group you have in the current data the restructured file will have one target variable.

In this step, choose how to identify case groups in the restructured data, and choose which variables belong with each target variable.

Optionally, you can also choose variables to copy to the new file as Fixed Variables.

Variables in the Current File:

- id
- 性別 [Gender]
- 治療方式 [Treatment]
- 年齡 [Age] (highlighted)
- DepressionScale_1
- DepressionScale_2
- DepressionScale_3
- DepressionScale_4
- DepressionScale_5

Case Group Identification

Use case number
Name: id1 [Label...](#)

Variables to be Transposed

Target Variable: DepressionScale (highlighted)

- DepressionScale_1
- DepressionScale_2
- DepressionScale_3
- DepressionScale_4

1

Fixed Variable(s):

- id
- 性別 [Gender]
- 治療方式 [Treatment]

2

3

< Back [Next >](#) [Finish](#) [Cancel](#) [Help](#)

GEE/LMM-資料轉置處理

 Restructure Data Wizard - Step 4 of 7

Variables to Cases: Create Index Variables

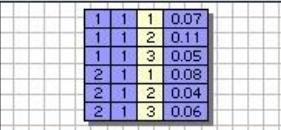
In the current data, values for a variable group appear in a single case in multiple variables. For example, a single case contains the values for w1, w2, and w3.

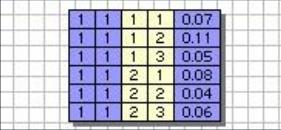
In the new data, values for a variable group will appear in multiple cases in a single variable. For example, there will be three cases, one each for w1, w2, and w3.

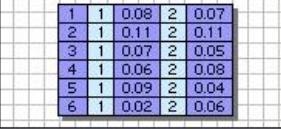
An index is a new variable that identifies the group of new cases that was created from the original case. For example, an index named "w" would have the values 1, 2, and 3.

How many index variables do you want to create?

- One
Use this when a variable group records the effects of a single factor, treatment or condition.
- More than one How many?
Use this when a variable group records the effects of more than one factor, treatment or condition.
- None
Use this if index information is stored in one of the sets of variables to be transposed.







< Back Next > Finish Cancel Help

 Restructure Data Wizard - Step 5 of 7

Variables to Cases: Create One Index Variable

You have chosen to create one index variable. The variable's values can be sequential numbers or the names of variables in a group.

In the table you can specify the name and label for the index variable.

What kind of index values?

- Sequential numbers
Index Values: 1, 2, 3, 4, 5
- Variable names
Index Values: DepressionScale_1, DepressionScale_2, DepressionScale_3, Depr...

Edit the Index Variable Name and Label:

Name	Label	Levels	Index Values
Visit		5	1, 2, 3, 4, 5

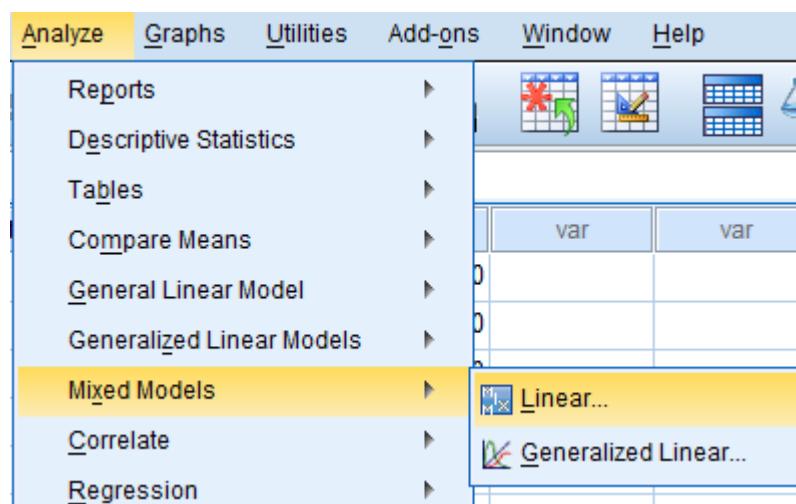
< Back Next > Finish Cancel Help

Mixed model

id	Gender	Treatment	Visit	DepressionScale
1	2	1	1	4.0
1	2	1	2	32.0
1	2	1	3	13.0
1	2	1	4	19.0
1	2	1	5	11.0
2	2	1	1	29.0
2	2	1	2	14.0
2	2	1	3	8.0
2	2	1	4	5.0
2	2	1	5	4.0
3	1	1	1	19.0
3	1	1	2	27.0
3	1	1	3	9.0
3	1	1	4	15.0
3	1	1	5	9.0

Mixed model analysis

分析>混合模式>線性



The screenshot shows the SPSS menu bar with 'Analyze' selected. Under 'Analyze', there is a tree view of statistical procedures. The 'Mixed Models' node is expanded, and its first child, 'Linear...', is highlighted with a yellow background.

Linear Mixed Models: Specify Subjects and Repeated

Click Continue for models with uncorrelated terms.
 Specify Subject variable for models with correlated random effects.
 Specify both Repeated and Subject variables for models with correlated residuals within the random effects.

Subjects: id

Repeated: Visit

Repeated Covariance Type: Compound Symmetry

Buttons: Continue, Reset, Cancel, Help

**重複共變異數類型:
複合對稱
(選取AIC及BIC最小)**

Mixed model analysis

Linear Mixed Models

Dependent Variable: 愛鬱分數 [Depression...]

Factor(s): 性別 [Gender], 治療方式 [Treatment], Visit

Covariate(s):

Residual Weight:

Fixed... Statistics... EM Means... Save...

OK Paste Reset Cancel Help

Linear Mixed Models: Fixed Effects

Fixed Effects: Build terms, Build nested terms

Factors and Covariates: Gender, Treatment, Visit

Model: Gender, Treatment, Visit, Treatment*Visit

Factorial, By*, (Within), Clear Term, Add, Remove

Build Term: Include intercept, Sum of squares: Type III

Continue Cancel Help

Linear Mixed Models: EM Means

Estimated Marginal Means of Fitted Models

Factors(s) and Factor Interactions: (OVERALL), Gender, Treatment, Visit, Treatment*Visit

Display Means for: Gender, Treatment, Visit, Treatment*Visit

Compare main effects, Confidence Interval Adjustment: Bonferroni, Reference Category

Linear Mixed Models: Statistics

Summary Statistics: Descriptive statistics, Case Processing Summary

Model Statistics: Parameter estimates, Tests for covariance parameters, Correlations of parameter estimates, Covariances of parameter estimates, Covariances of random effects, Covariances of residuals, Contrast coefficient matrix

Confidence interval: 95 %

Continue Cancel Help

Mixed model output

Information Criteria^a

-2 Restricted Log Likelihood	650.289
Akaike's Information Criterion (AIC)	654.289
Hurvich and Tsai's Criterion (AICC)	654.433
Bozdogan's Criterion (CAIC)	661.198
Schwarz's Bayesian Criterion (BIC)	659.198

The information criteria are displayed in smaller-is-better form.

a. Dependent Variable: 豐饒分數

Pairwise Comparisons^a

(I) Visit	(J) Visit	Mean Difference (I-J)	Std. Error	df	Sig. ^b	95% Confidence Interval for Difference ^b	
						Lower Bound	Upper Bound
1	2	-2.712	2.741	68.882	1.000	-10.662	5.238
	3	2.429	2.741	68.882	1.000	-5.521	10.380
	4	3.793	2.741	68.882	1.000	-4.157	11.743
	5	3.567	2.868	70.166	1.000	-4.744	11.879
2	1	2.712	2.741	68.882	1.000	-5.238	10.662
	3	5.141	2.741	68.882	.649	-2.809	13.092
	4	6.505	2.741	68.882	.204	-1.445	14.455
	5	6.279	2.868	70.166	.319	-2.032	14.591
3	1	-2.429	2.741	68.882	1.000	-10.380	5.521
	2	-5.141	2.741	68.882	.649	-13.092	2.809
	4	1.364	2.741	68.882	1.000	-6.587	9.314
	5	1.138	2.868	70.166	1.000	-7.174	9.450
4	1	-3.793	2.741	68.882	1.000	-11.743	4.157
	2	-6.505	2.741	68.882	.204	-14.455	1.445
	3	-1.364	2.741	68.882	1.000	-9.314	6.587
	5	.226	2.868	70.166	1.000	-8.537	8.086
5	1	-3.567	2.868	70.166	1.000	-11.879	4.744
	2	-6.279	2.868	70.166	.319	-14.591	2.032
	3	-1.138	2.868	70.166	1.000	-9.450	7.174
	4	.226	2.868	70.166	1.000	-8.086	8.537

Based on estimated marginal means

a. Dependent Variable: 豐饒分數.

Estimates of Fixed Effects^a

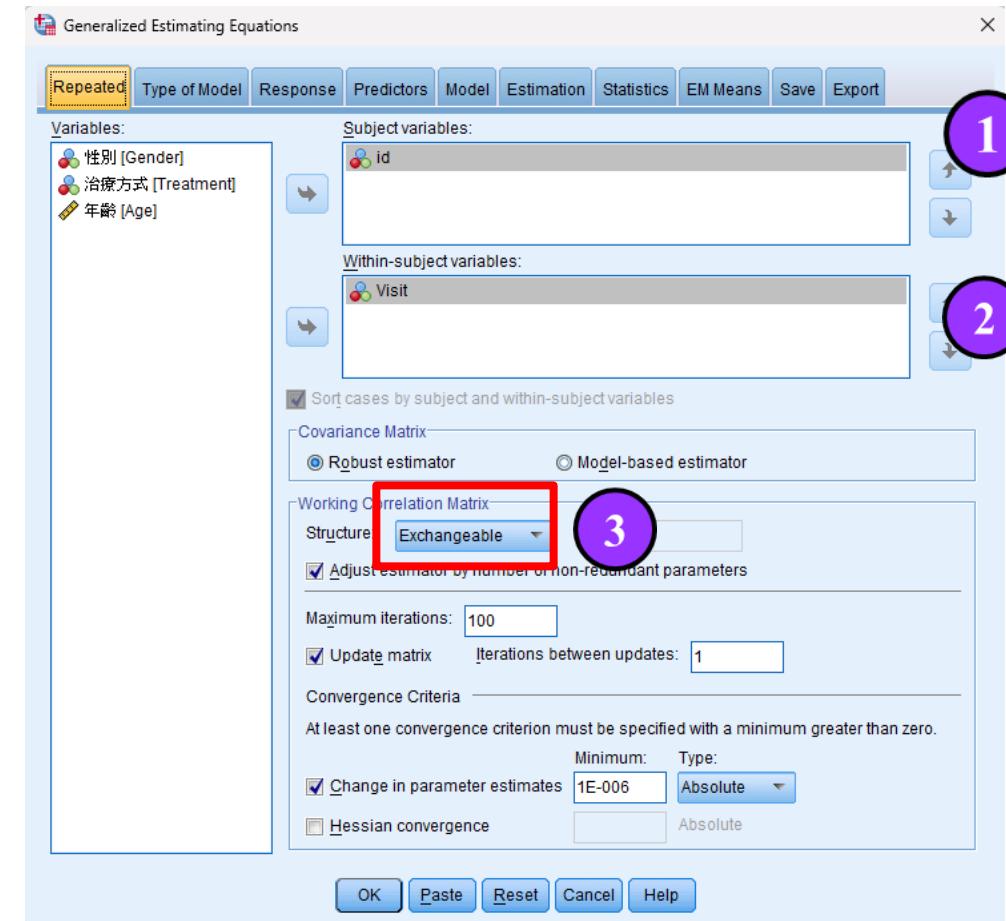
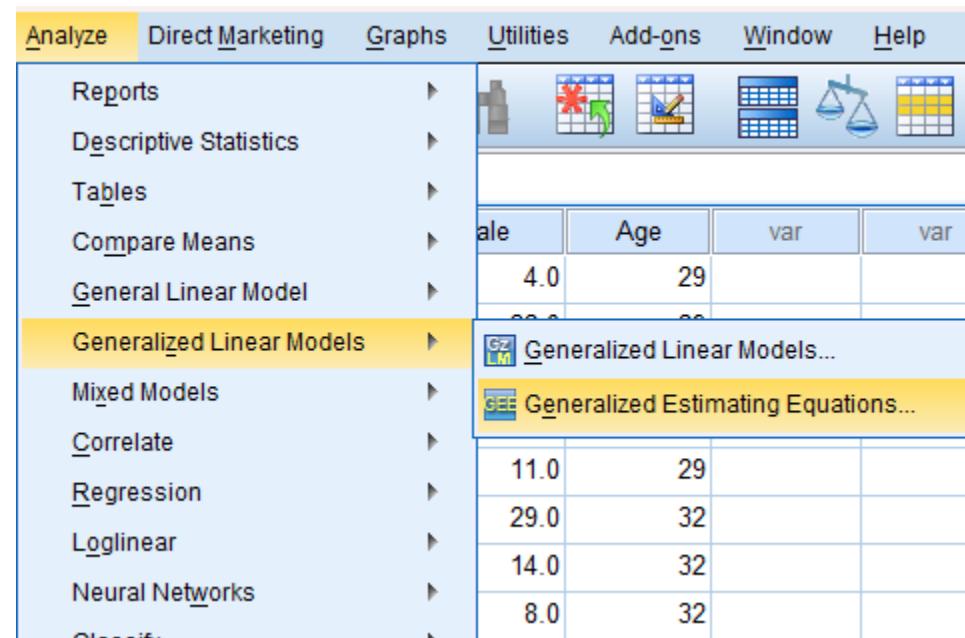
Parameter	Estimate	Std. Error	df	t	Sig.	95% Confidence Interval	
						Lower Bound	Upper Bound
Intercept	10.457603	3.313212	74.672	3.156	.002	3.856871	17.058335
[Gender=1]	2.029279	2.351907	16.727	.863	.400	-2.938983	6.997541
[Gender=2]	0 ^b	0
[Treatment=0]	3.590407	4.499750	81.501	.798	.427	-5.361851	12.542665
[Treatment=1]	0 ^b	0
[Visit=1]	3.529088	3.898315	70.588	.905	.368	-4.244717	11.302894
[Visit=2]	3.619997	3.898315	70.588	.929	.356	-4.153808	11.393803
[Visit=3]	-2.107276	3.898315	70.588	-.541	.591	-9.881081	5.666530
[Visit=4]	.165452	3.898315	70.588	.042	.966	-7.608354	7.939257
[Visit=5]	0 ^b	0
[Visit=1] * [Treatment=0]	.076556	5.731167	70.319	.013	.989	-11.352986	11.506098
[Visit=2] * [Treatment=0]	5.318980	5.731167	70.319	.928	.357	-6.110562	16.748522
[Visit=3] * [Treatment=0]	6.490697	5.731167	70.319	1.133	.261	-4.938845	17.920239
[Visit=4] * [Treatment=0]	-.782030	5.731167	70.319	-.136	.892	-12.211572	10.647512
[Visit=5] * [Treatment=0]	0 ^b	0
[Visit=1] * [Treatment=1]	0 ^b	0
[Visit=2] * [Treatment=1]	0 ^b	0
[Visit=3] * [Treatment=1]	0 ^b	0
[Visit=4] * [Treatment=1]	0 ^b	0
[Visit=5] * [Treatment=1]	0 ^b	0

a. Dependent Variable: 豐饒分數.

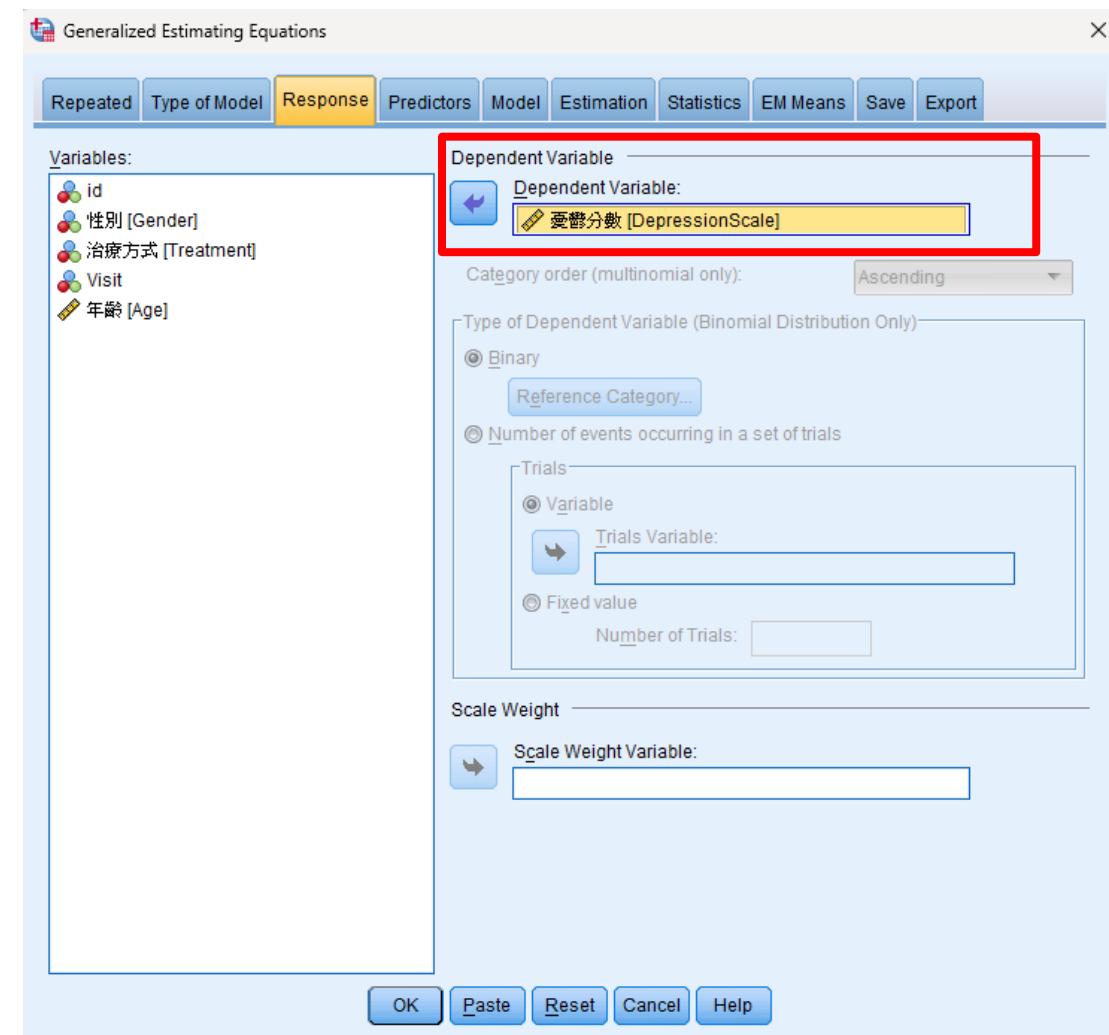
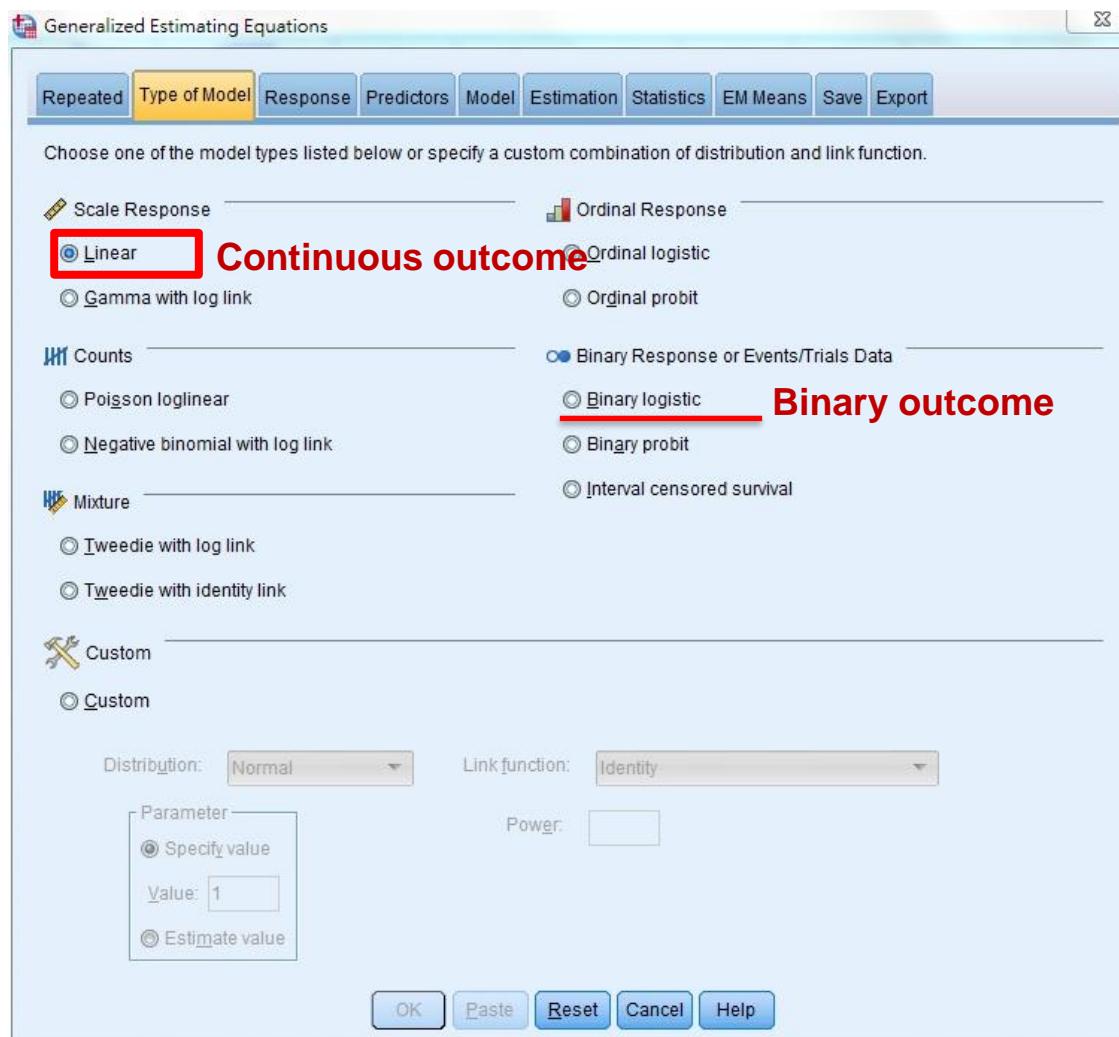
b. This parameter is set to zero because it is redundant.

GEE analysis

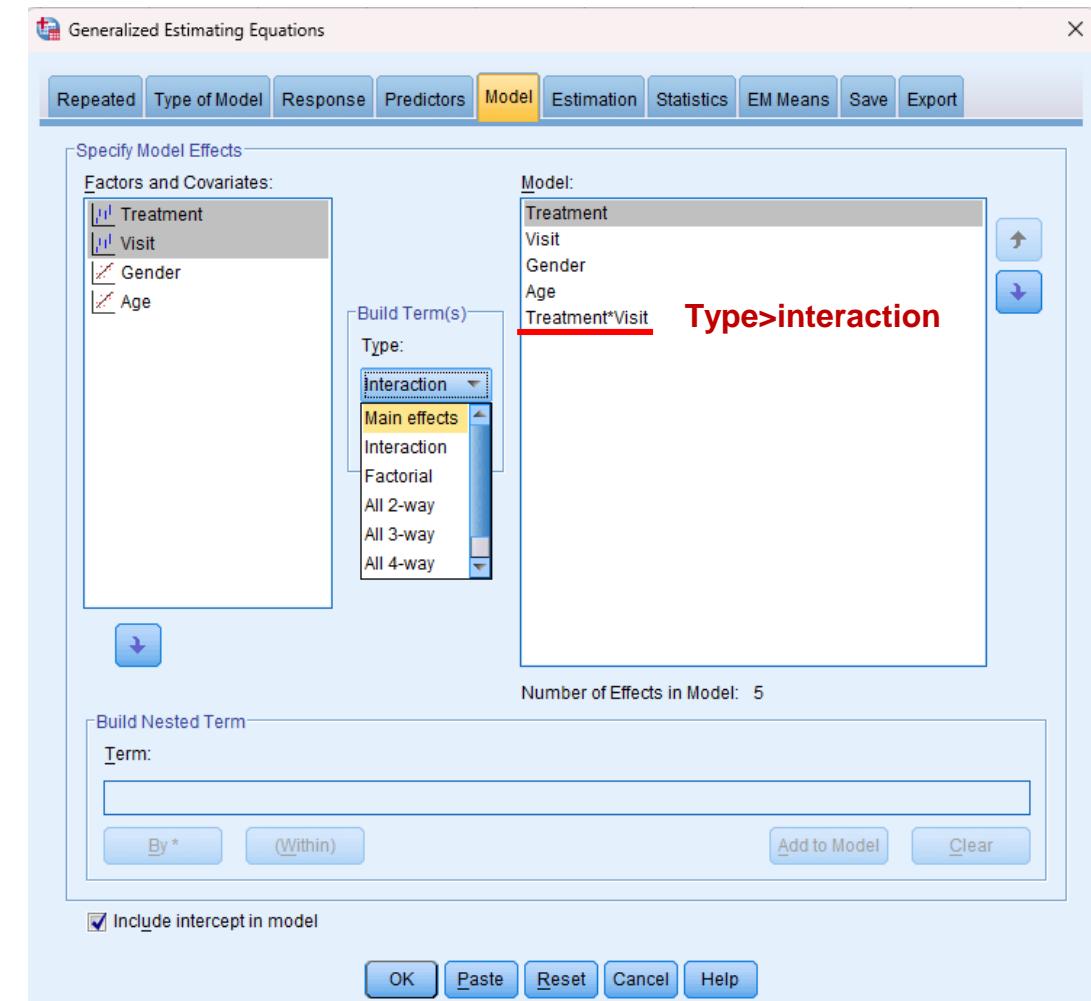
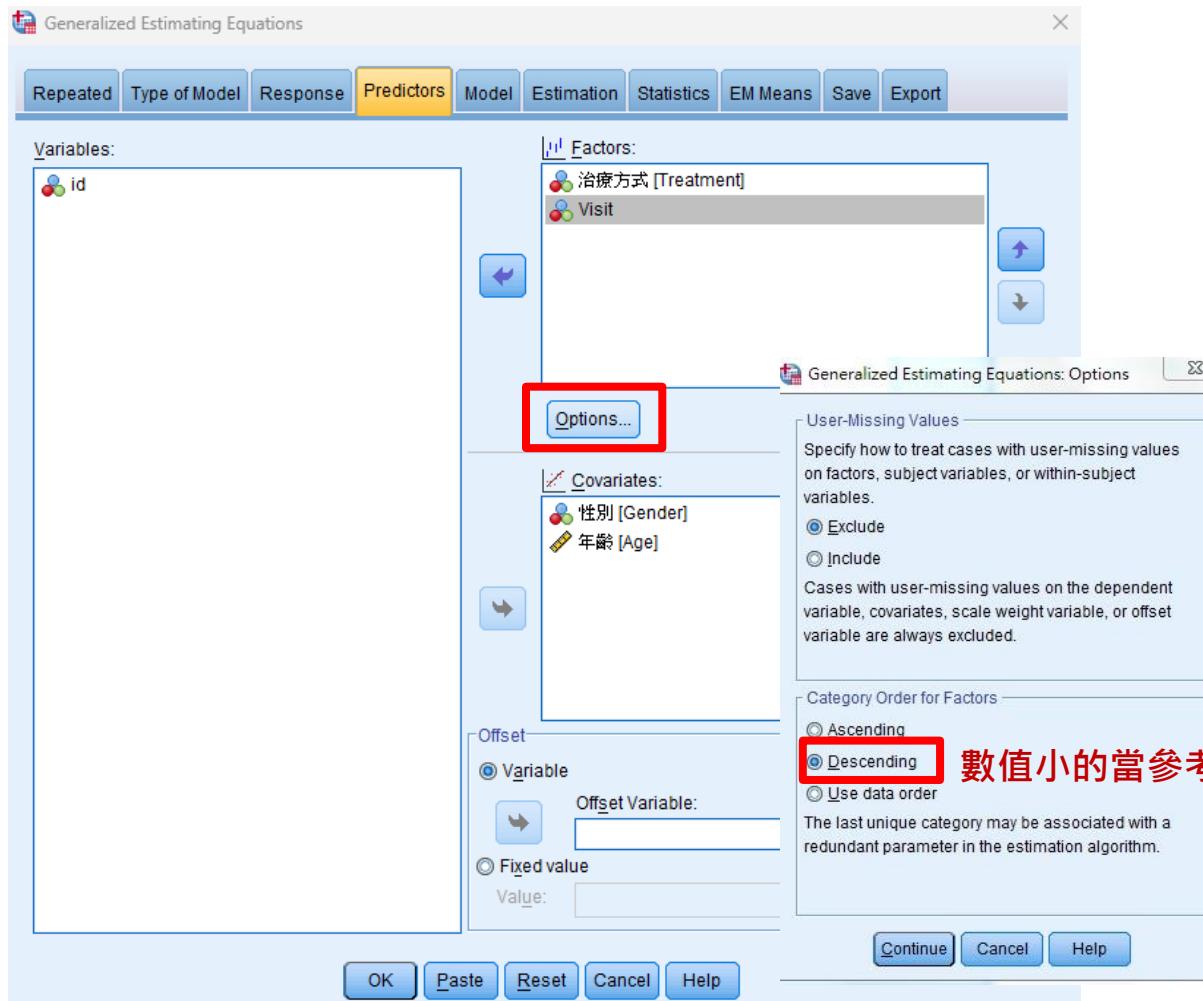
分析>廣義線性模型>廣義估計方程式(GEE)



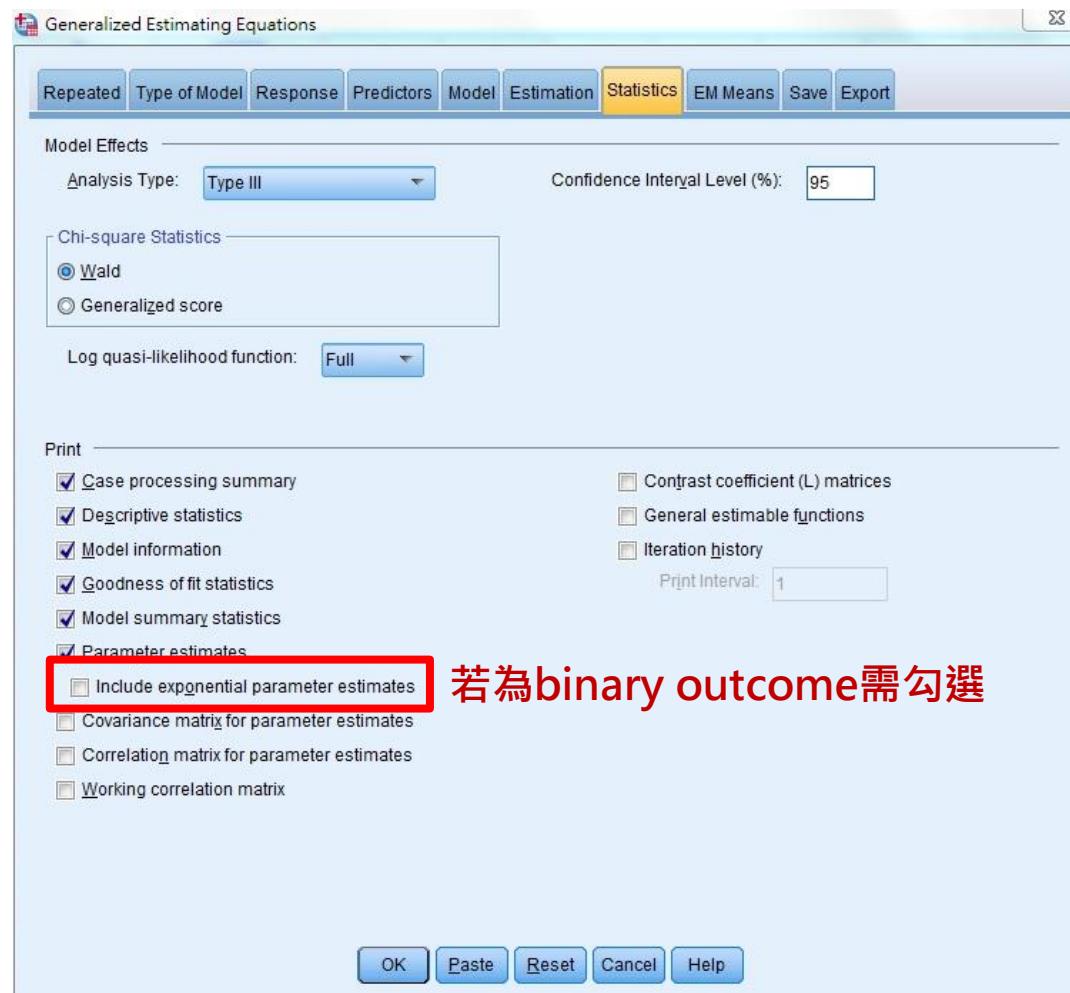
GEE analysis



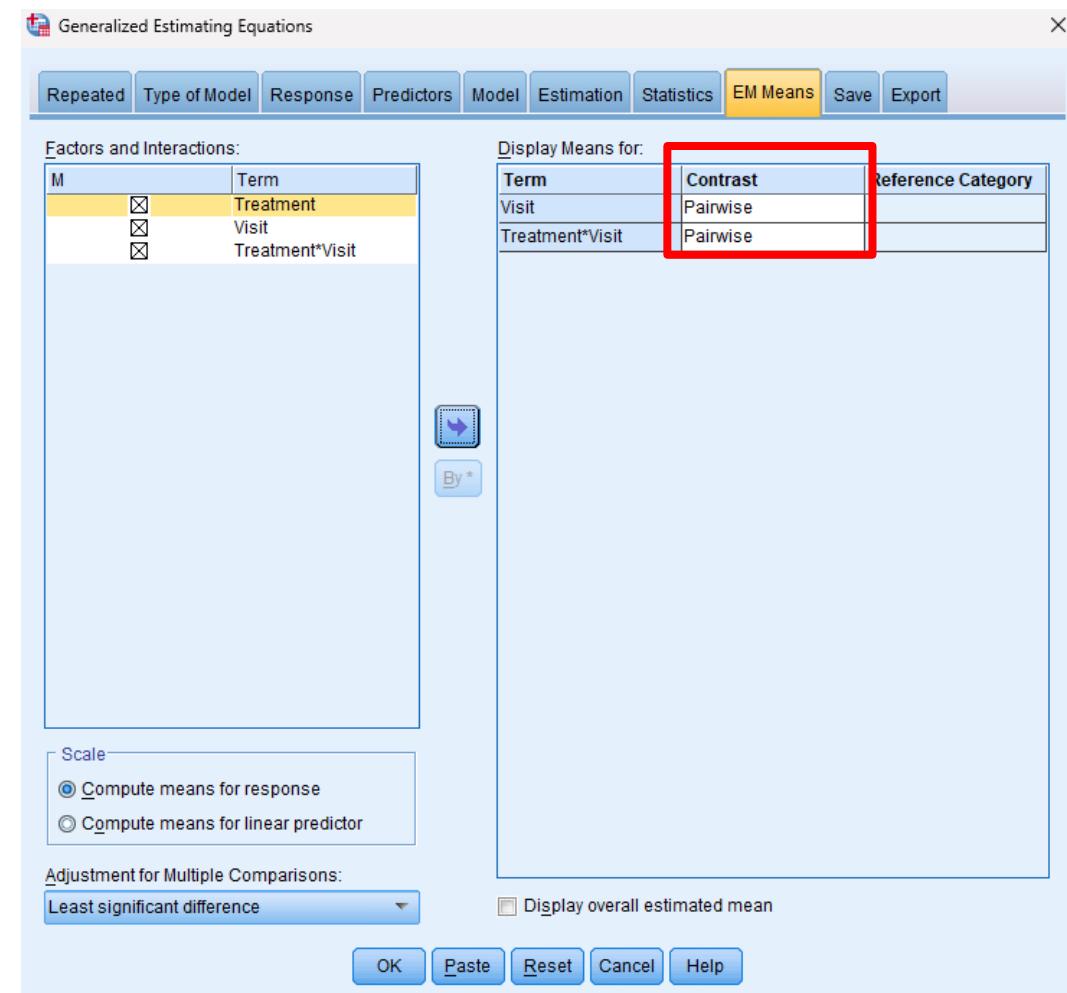
GEE analysis



GEE analysis



若為binary outcome需勾選



GEE-Output

Model Information

Dependent Variable	憂鬱分數
Probability Distribution	Normal
Link Function	Identity
Subject Effect	1 id
Within-Subject Effect	1 Visit
Working Correlation Matrix Structure	Exchangeable

Case Processing Summary

	N	Percent
Included	97	97.0%
Excluded	3	3.0%
Total	100	100.0%

Categorical Variable Information

		N	Percent
Factor	治療方式	Placebo	44 45.4%
		Treatment	53 54.6%
		Total	97 100.0%
	Visit	5	17 17.5%
		4	20 20.6%
		3	20 20.6%
		2	20 20.6%
		1	20 20.6%
	Total	97	100.0%

Goodness of Fit^a

	Value
Quasi Likelihood under Independence Model Criterion (QIC) ^b	7393.042
Corrected Quasi Likelihood under Independence Model Criterion (QICC) ^b	7389.820

a. Information criteria are in smaller-is-better form.
 b. Computed using the full log quasi-likelihood function.

QIC/QICC數值越小越好

Parameter Estimates

Parameter	B	Std. Error	95% Wald Confidence Interval		Hypothesis Test		
			Lower	Upper	Wald Chi-Square	df	Sig.
(Intercept)	23.711	17.2393	-10.078	57.499	1.892	1	.169
[Treatment=2]	3.616	3.3682	-2.986	10.218	1.153	1	.283
[Treatment=1]	0 ^a
[Visit=5]	-3.502	3.4075	-10.180	3.177	1.056	1	.304
[Visit=4]	-3.364	3.7090	-10.633	3.906	.822	1	.364
[Visit=3]	-5.636	2.7732	-11.072	-.201	4.131	1	.042
[Visit=2]	.091	4.6859	-9.093	9.275	.000	1	.985
[Visit=1]	0 ^a
Gender	-2.306	2.2536	-6.723	2.111	1.047	1	.306
Age	-.166	.4758	-1.099	.767	.122	1	.727
[Treatment=2] * [Visit=5]	-.113	4.8978	-9.712	9.487	.001	1	.982
[Treatment=2] * [Visit=4]	-.859	5.5842	-11.803	10.086	.024	1	.878
[Treatment=2] * [Visit=3]	6.414	5.4913	-4.349	17.177	1.364	1	.243
[Treatment=2] * [Visit=2]	5.242	6.0191	-6.555	17.040	.759	1	.384
[Treatment=2] * [Visit=1]	0 ^a
[Treatment=1] * [Visit=5]	0 ^a
[Treatment=1] * [Visit=4]	0 ^a
[Treatment=1] * [Visit=3]	0 ^a
[Treatment=1] * [Visit=2]	0 ^a
[Treatment=1] * [Visit=1]	0 ^a
(Scale)	86.657

Dependent Variable: 憂鬱分數
 Model: (Intercept), Treatment, Visit, Gender, Age, Treatment * Visit
 a. Set to zero because this parameter is redundant.

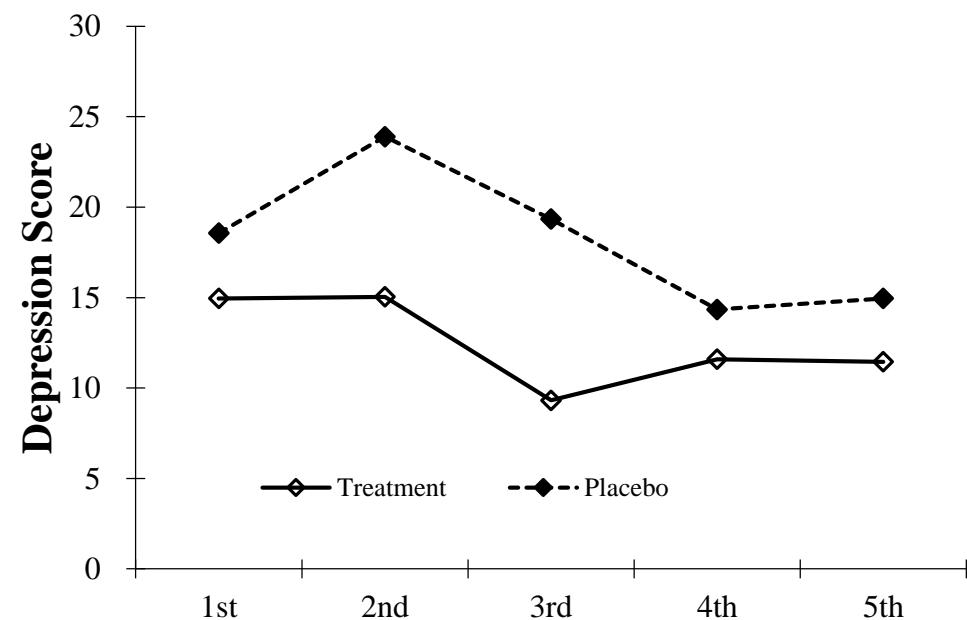
GEE-Output

Estimates

治療方式	Visit	Mean	Std. Error	95% Wald Confidence Interval	
				Lower	Upper
Placebo	5	14.953	45.0687	-73.380	103.286
	4	14.345	44.2540	-72.391	101.081
	3	19.345	44.3947	-67.667	106.357
	2	23.901	44.2926	-62.911	110.712
	1	18.567	44.0893	-67.846	104.981
Treatment	5	11.449	42.0461	-70.959	93.858
	4	11.587	43.0089	-72.708	95.883
	3	9.315	43.0638	-75.089	93.718
	2	15.042	43.0606	-69.355	99.439
	1	14.951	42.8486	-69.031	98.933

Covariates appearing in the model are fixed at the following values:

Gender=1.54; Age=31.42



感謝您的聆聽！

Thank you !