

Appendix A - Children Under 8 Years

Statistics			
		Gender	Other_Allergies
N	Valid	63	63
	Missing	2	2

Frequency Table

Gender					
		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Boy	41	63.1	65.1	65.1
	Girl	22	33.8	34.9	100.0
	Total	63	96.9	100.0	
Missing	System	2	3.1		
Total		65	100.0		

Other Allergies					
		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	54	83.1	85.7	85.7
	No	9	13.8	14.3	100.0
	Total	63	96.9	100.0	
Missing	System	2	3.1		
Total		65	100.0		

CROSSTABS

/TABLES=Gender BY Other_Allergies

/SHOWDIM=2

/FORMAT=AVALUE TABLES

/STATISTICS=CHISQ D

/CELLS=COUNT COLUMN

/COUNT ROUND CELL.

Crosstabs

Case Processing Summary

	Valid		Cases Missing		Total	
	N	Percent	N	Percent	N	Percent
Gender * Other_Allergies	63	96.9%	2	3.1%	65	100.0%

Gender * Other Allergies Crosstabulation

		Other_Allergies		Total	
		Yes	No		
Gender	Boy	Count	37	4	41
		% within Other_Allergies	68.5%	44.4%	65.1%
	Girl	Count	17	5	22
		% within Other_Allergies	31.5%	55.6%	34.9%
Total		Count	54	9	63
		% within Other_Allergies	100.0%	100.0%	100.0%

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)	Exact Sig. (2-sided)	Exact Sig. (1-sided)
Pearson Chi-Square	1.967 ^a	1	.161		
Continuity Correction ^b	1.051	1	.305		
Likelihood Ratio	1.878	1	.171		
Fisher's Exact Test				.256	.153
Linear-by-Linear Association	1.936	1	.164		
N of Valid Cases	63				

a. 1 cells (25.0%) have expected count less than 5. The minimum expected count is 3.14.

b. Computed only for a 2x2 table

Directional Measures

			Value	Asymptotic Standard Error ^a	Approximate T ^b
Ordinal by Ordinal	Somers' d	Symmetric	.169	.126	1.282
		Gender Dependent	.241	.177	1.282
		Other_Allergies Dependent	.130	.101	1.282

Directional Measures

			Approximate Significance
Ordinal by Ordinal	Somers' d	Symmetric	.200
		Gender Dependent	.200
		Other_Allergies Dependent	.200

a. Not assuming the null hypothesis.

b. Using the asymptotic standard error assuming the null hypothesis.

Crosstabs - Co-Morbidity to All Three Allergic Conditions [hay fever ; asthma and eczema] by Gender

Comparison of proportions calculator

Sample 1

Proportion (%):

Sample size:

Sample 2

Proportion (%):

Sample size:

Results

Difference	10.8 %
95% CI	-11.5486% to 28.1867%
Chi-squared	1.004
DF	1
Significance level	P = 0.3164

Frequencies

		Statistics						
		Peanut	Tree_Nuts	Egg	Soy	Dairy	Seafood	Wheat
N	Valid	45	49	34	6	23	13	8
	Missing	20	16	31	59	42	52	57

Statistics

		Sesame_seeds	Kiwi	Other_Allergens	Number_of_Allergens
N	Valid	6	5	13	65
	Missing	59	60	52	0

Frequency Table

Peanut

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	45	69.2	100.0	100.0
Missing	System	20	30.8		
Total		65	100.0		

Tree Nuts

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	49	75.4	100.0	100.0
Missing	System	16	24.6		
Total		65	100.0		

Egg

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	34	52.3	100.0	100.0
Missing	System	31	47.7		
Total		65	100.0		

Soy

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	6	9.2	100.0	100.0
Missing	System	59	90.8		
Total		65	100.0		

Dairy

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	23	35.4	100.0	100.0
Missing	System	42	64.6		
Total		65	100.0		

Seafood

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	13	20.0	100.0	100.0
Missing	System	52	80.0		
Total		65	100.0		

Wheat

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	8	12.3	100.0	100.0
Missing	System	57	87.7		
Total		65	100.0		

Sesame seeds

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	6	9.2	100.0	100.0
Missing	System	59	90.8		
Total		65	100.0		

Kiwi

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	5	7.7	100.0	100.0
Missing	System	60	92.3		
Total		65	100.0		

Other Allergens

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	1 Other	9	13.8	69.2	69.2
	2 Other	2	3.1	15.4	84.6
	Other	1	1.5	7.7	92.3
	4+Other	1	1.5	7.7	100.0
	Total	13	20.0	100.0	
Missing	System	52	80.0		
Total		65	100.0		

Number of Allergens

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	1 Allergen	15	23.1	23.1	23.1
	2 - 3 Allergens	23	35.4	35.4	58.5
	4 Allergens or More	27	41.5	41.5	100.0
	Total	65	100.0	100.0	

Crosstabs

[Children Under 8 Years - Co-Morbidity to Allergic Rhinitis, Asthma and Eczema

Case Processing Summary

	Cases					
	Valid		Missing		Total	
	N	Percent	N	Percent	N	Percent
Gender * Atopic	63	98.4%	1	1.6%	64	100.0%

Gender * Atopic Crosstabulation

Count

		Atopic		Total
		yes	no	
Gender	Boys	10	31	41
	Girls	3	19	22
Total		13	50	63

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)	Exact Sig. (2-sided)	Exact Sig. (1-sided)
Pearson Chi-Square	1.011 ^a	1	.315		
Continuity Correction ^b	.461	1	.497		
Likelihood Ratio	1.064	1	.302		
Fisher's Exact Test				.515	.253
Linear-by-Linear Association	.995	1	.319		
N of Valid Cases	63				

a. 1 cells (25.0%) have expected count less than 5. The minimum expected count is 4.54.

b. Computed only for a 2x2 table

Directional Measures

			Value	Asymptotic Standard Error ^a	Approximate T ^b
Ordinal by Ordinal	Somers' d	Symmetric	.125	.114	1.079
		Gender Dependent	.149	.136	1.079
		Atopic Dependent	.108	.099	1.079

Directional Measures

			Approximate Significance
Ordinal by Ordinal	Somers' d	Symmetric	.280
		Gender Dependent	.280
		Atopic Dependent	.280

a. Not assuming the null hypothesis.

b. Using the asymptotic standard error assuming the null hypothesis.

Symmetric Measures

		Value	Asymptotic Standard Error ^a	Approximate T ^b	Approximate Significance
Interval by Interval	Pearson's R	.127	.115	.997	.322 ^c
Ordinal by Ordinal	Spearman Correlation	.127	.115	.997	.322 ^c
N of Valid Cases		63			

- a. Not assuming the null hypothesis.
- b. Using the asymptotic standard error assuming the null hypothesis.
- c. Based on normal approximation.

```
FREQUENCIES VARIABLES=Miss_Friends_BirthdaysParents_Fuss Feel_Different_to_Fr
iends
    Get_Upset_Not_Able_to_Eat_Certain_FoodAttend_Specialist
/BARCHART FREQ
/ORDER=ANALYSIS.
```

Frequencies

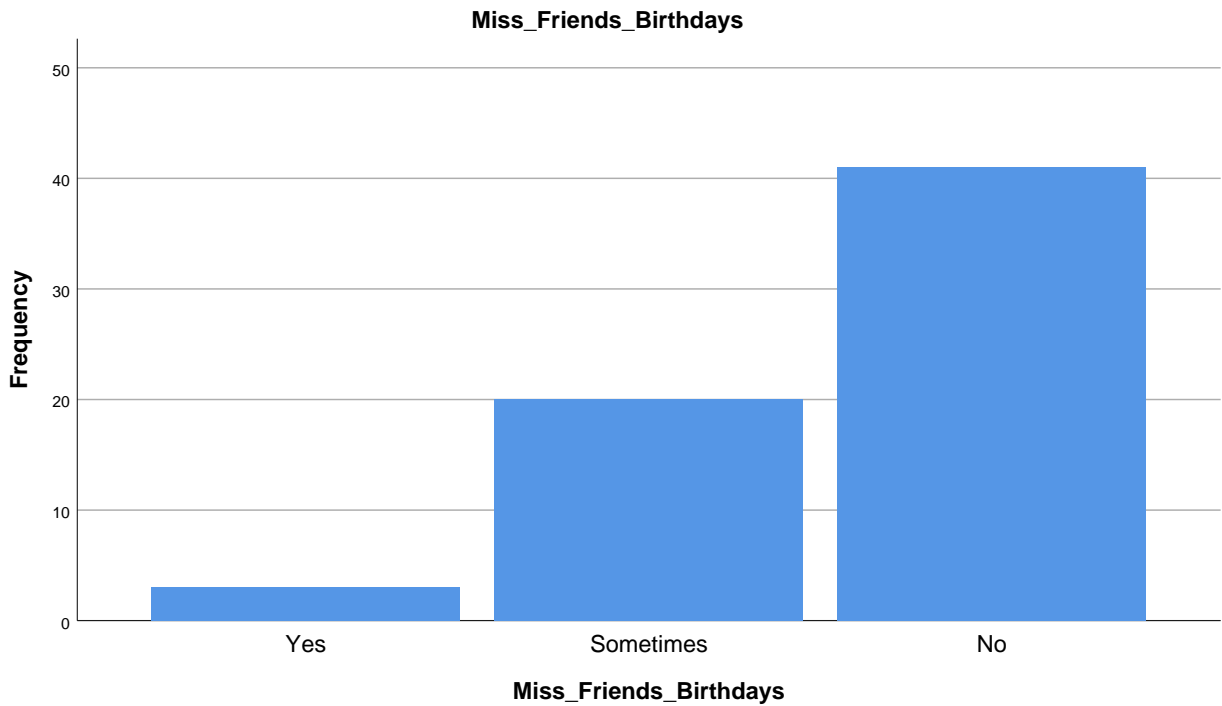
Statistics						
		Miss_Friends_ Birthdays	Parents_Fuss	Feel_Different_t o_Friends	Get_Upset_Not _Able_to_Eat_ Certain_Foods	Attend_Speciali st
N	Valid	64	64	64	64	65
	Missing	1	1	1	1	0

Frequency Table

Miss Friends Birthdays

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	3	4.6	4.7	4.7
	Sometimes	20	30.8	31.3	35.9
	No	41	63.1	64.1	100.0
	Total	64	98.5	100.0	
Missing	99	1	1.5		
Total		65	100.0		

Bar Chart



CROSSTABS

```

/TABLES=Miss_Friends_BirthdaysBY Gender BY Group
/SHOWDIM=2
/FORMAT=AVALUE TABLES
/STATISTICS=CHISQ D
/CELLS=COUNT COLUMN
/COUNT ROUND CELL.

```

Crosstabs

Case Processing Summary

	Valid		Cases Missing		Total	
	N	Percent	N	Percent	N	Percent
Miss_Friends_Birthdays * Gender * Group	63	96.9%	2	3.1%	65	100.0%

Miss Friends' Birthdays * Gender * Group Crosstabulation

Group: Total

			Gender		Total
			Boy	Girl	
Miss_Friends_Birthdays	Yes	Count	2	1	3
		% within Gender	4.9%	4.5%	4.8%
	Sometimes	Count	15	4	19
		% within Gender	36.6%	18.2%	30.2%
	No	Count	24	17	41
		% within Gender	58.5%	77.3%	65.1%
Total		Count	41	22	63
		% within Gender	100.0%	100.0%	100.0%

Chi-Square Tests

Group		Value	df	Asymptotic Significance (2- sided)
Under 8 Years	Pearson Chi-Square	2.384 ^a	2	.304
	Likelihood Ratio	2.503	2	.286
	Linear-by-Linear Association	1.531	1	.216
	N of Valid Cases	63		
Total	Pearson Chi-Square	2.384 ^a	2	.304
	Likelihood Ratio	2.503	2	.286
	Linear-by-Linear Association	1.531	1	.216
	N of Valid Cases	63		

a. 2 cells (33.3%) have expected count less than 5. The minimum expected count is 1.05.

Directional Measures

Group	Value	Asymptotic Standard Error ^a	Approximate T ^b	Approximate Significance
Under 8 Years	.174	.117	1.477	.140
	.180	.121	1.477	.140
	.169	.114	1.477	.140

a. Not assuming the null hypothesis.

b. Using the asymptotic standard error assuming the null hypothesis.

CROSSTABS

```

/TABLES=Miss_Friends_BirthdaysBY Feel_Different_to_Friends
/SHOWDIM=2
/FORMAT=AVALUE TABLES
/STATISTICS=CHISQ D
/CELLS=COUNT COLUMN
/COUNT ROUND CELL.

```

Crosstabs

Case Processing Summary

	Valid		Cases Missing		Total	
	N	Percent	N	Percent	N	Percent
Miss_Friends_Birthdays * Feel_Different_to_Friends	64	98.5%	1	1.5%	65	100.0%

Miss Friends Birthdays * Feel Different to Friends Crosstabulation

			Feel_Different_to_Friends		
			Yes	Sometimes	No
Miss_Friends_Birthdays	Yes	Count	3	0	0
		% within Feel_Different_to_Friends	12.5%	0.0%	0.0%
	Sometimes	Count	9	6	5
		% within Feel_Different_to_Friends	37.5%	31.6%	23.8%
	No	Count	12	13	16
		% within Feel_Different_to_Friends	50.0%	68.4%	76.2%
Total	Count	24	19	21	
	% within Feel_Different_to_Friends	100.0%	100.0%	100.0%	

Miss Friends Birthdays * Feel Different to Friends Crosstabulation

			Total
Miss_Friends_Birthdays	Yes	Count	3
		% within Feel_Different_to_Friends	4.7%
	Sometimes	Count	20
		% within Feel_Different_to_Friends	31.3%
	No	Count	41
		% within Feel_Different_to_Friends	64.1%
Total	Count	64	
	% within Feel_Different_to_Friends	100.0%	

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	6.952 ^a	4	.138
Likelihood Ratio	7.884	4	.096
Linear-by-Linear Association	5.052	1	.025
N of Valid Cases	64		

a. 3 cells (33.3%) have expected count less than 5. The minimum expected count is .89.

Directional Measures

			Value	Asymptotic Standard Error ^a	Approximate T ^b
Ordinal by Ordinal	Somers' d	Symmetric	.237	.109	2.119
		Miss_Friends_Birthdays Dependent	.206	.097	2.119
		Feel_Different_to_Friends Dependent	.279	.126	2.119

Directional Measures

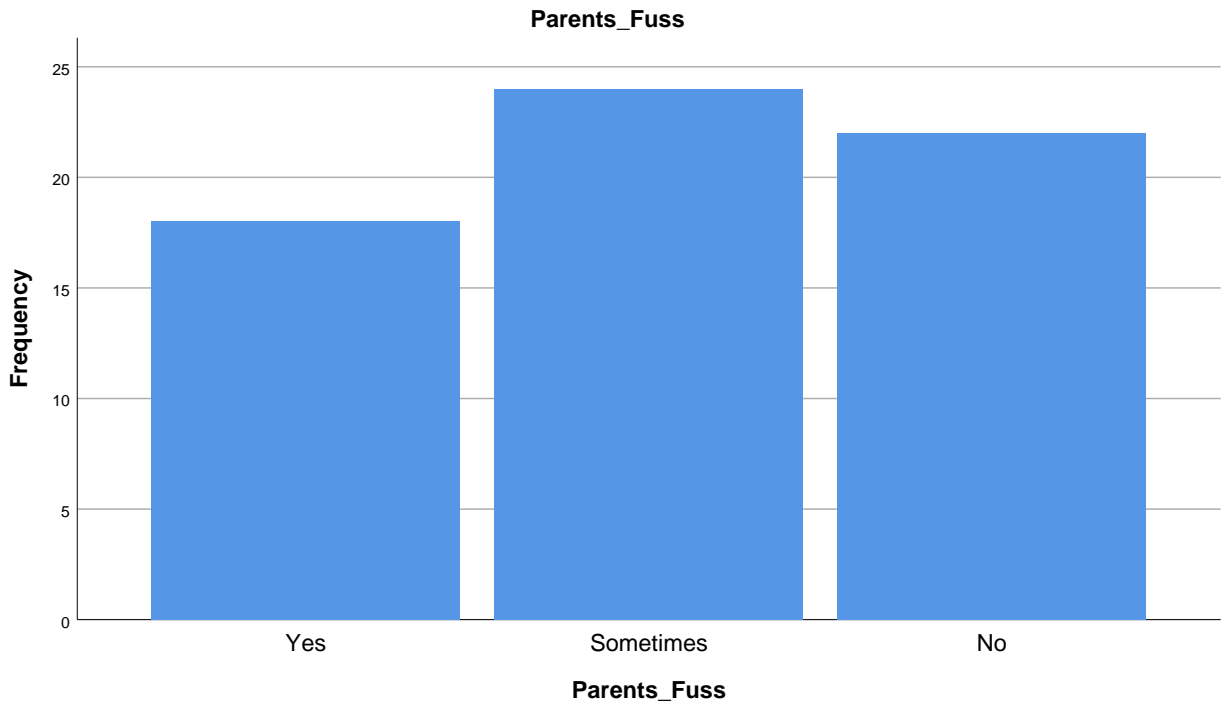
			Approximate Significance
Ordinal by Ordinal	Somers' d	Symmetric	.034
		Miss_Friends_Birthdays Dependent	.034
		Feel_Different_to_Friends Dependent	.034

a. Not assuming the null hypothesis.

b. Using the asymptotic standard error assuming the null hypothesis.

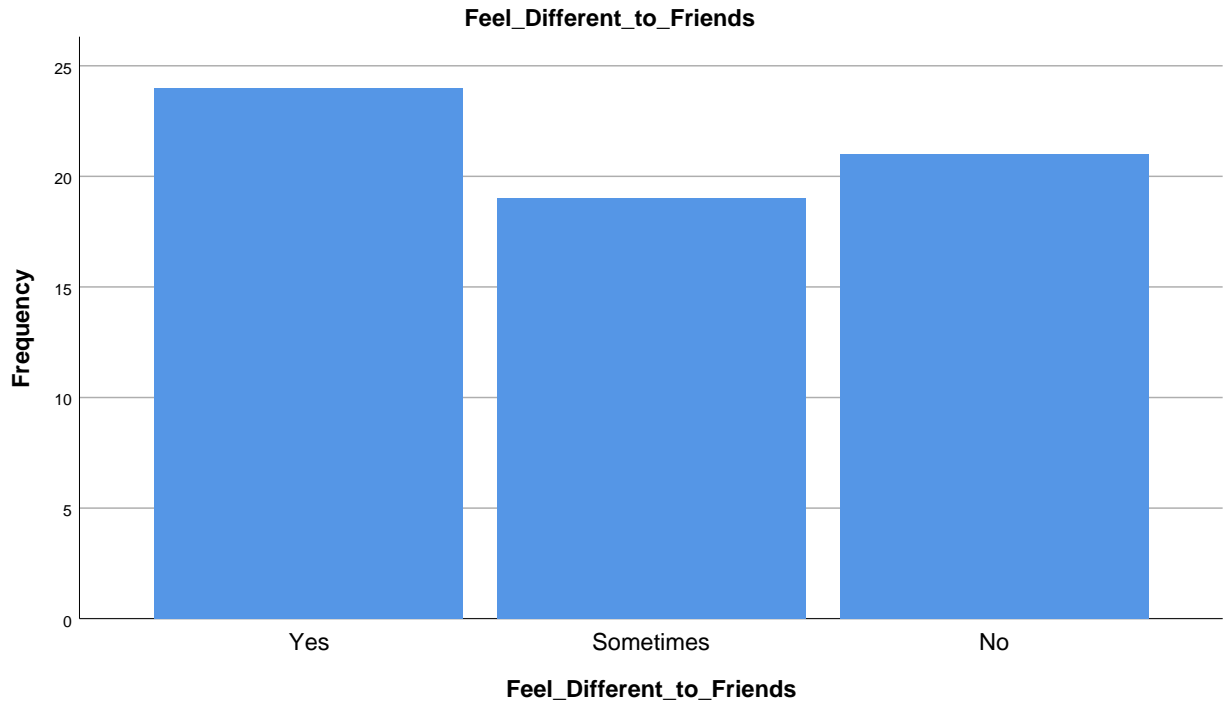
Parents Fuss

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	18	27.7	28.1	28.1
	Sometimes	24	36.9	37.5	65.6
	No	22	33.8	34.4	100.0
	Total	64	98.5	100.0	
Missing	99	1	1.5		
Total		65	100.0		



Feel Different to Friends

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	24	36.9	37.5	37.5
	Sometimes	19	29.2	29.7	67.2
	No	21	32.3	32.8	100.0
	Total	64	98.5	100.0	
Missing	99	1	1.5		
Total		65	100.0		



CROSSTABS

```

/TABLES=Feel_Different_to_FriendsBY Parents_Fuss
/SHOWDIM=2
/FORMAT=AVALUE TABLES
/STATISTICS=CHISQ D
/CELLS=COUNT COLUMN
/COUNT ROUND CELL.

```

Crosstabs

Case Processing Summary

	Valid		Cases Missing		Total	
	N	Percent	N	Percent	N	Percent
Feel_Different_to_Friends * Parents_Fuss	64	98.5%	1	1.5%	65	100.0%

Feel Different to Friends * Parents Fuss Crosstabulation

			Parents_Fuss		
			Yes	Sometimes	No
Feel_Different_to_Friends	Yes	Count	11	9	4
		% within Parents_Fuss	61.1%	37.5%	18.2%
	Sometimes	Count	6	8	5
		% within Parents_Fuss	33.3%	33.3%	22.7%
	No	Count	1	7	13
		% within Parents_Fuss	5.6%	29.2%	59.1%
Total		Count	18	24	22
		% within Parents_Fuss	100.0%	100.0%	100.0%

Feel Different to Friends * Parents Fuss Crosstabulation

			Total
Feel_Different_to_Friends	Yes	Count	24
		% within Parents_Fuss	37.5%
	Sometimes	Count	19
		% within Parents_Fuss	29.7%
	No	Count	21
		% within Parents_Fuss	32.8%
Total		Count	64
		% within Parents_Fuss	100.0%

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	14.215 ^a	4	.007
Likelihood Ratio	15.618	4	.004
Linear-by-Linear Association	13.007	1	.000
N of Valid Cases	64		

a. 0 cells (.0%) have expected count less than 5. The minimum expected count is 5.34.

Directional Measures

			Value	Asymptotic Standard Error ^a	Approximate T ^b
Ordinal by Ordinal	Somers' d	Symmetric	.405	.093	4.328
		Feel_Different_to_Friends Dependent	.406	.093	4.328
		Parents_Fuss Dependent	.405	.094	4.328

Directional Measures

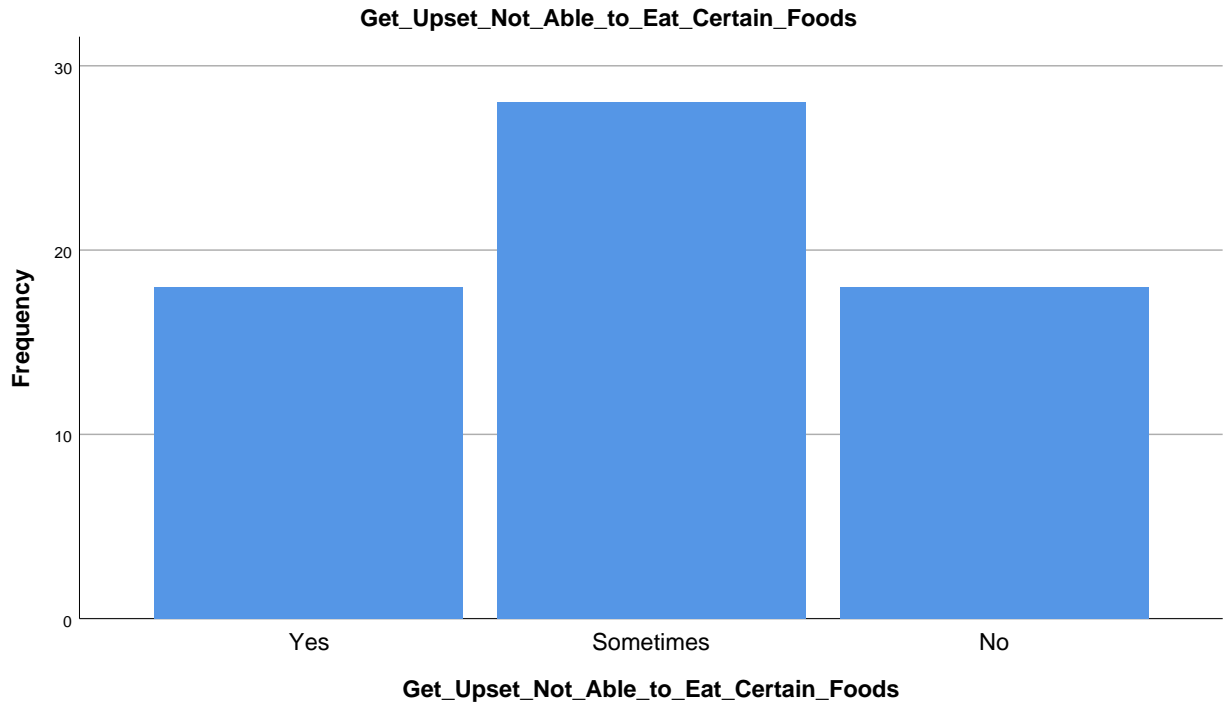
			Approximate Significance
Ordinal by Ordinal	Somers' d	Symmetric	.000
		Feel_Different_to_Friends Dependent	.000
		Parents_Fuss Dependent	.000

a. Not assuming the null hypothesis.

b. Using the asymptotic standard error assuming the null hypothesis.

Get Upset Not Able to Eat Certain Foods

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	18	27.7	28.1	28.1
	Sometimes	28	43.1	43.8	71.9
	No	18	27.7	28.1	100.0
	Total	64	98.5	100.0	
Missing	99	1	1.5		
Total		65	100.0		



CROSSTABS

```

/TABLES=Get_Upset_Not_Able_to_Eat_Certain_Foods BY Gender
/SHOWDIM=2
/FORMAT=AVALUE TABLES
/STATISTICS=CHISQ D
/CELLS=COUNT COLUMN
/COUNT ROUND CELL.

```

Crosstabs

Case Processing Summary

	Valid		Cases Missing		Total	
	N	Percent	N	Percent	N	Percent
Get_Upset_Not_Able_to_Eat_Certain_Foods * Gender	63	96.9%	2	3.1%	65	100.0%

Get Upset Not Able to Eat Certain Foods * Gender Crosstabulation

			Gender		
			Boy	Girl	Total
Get_Upset_Not_Able_to_Eat_Certain_Foods	Yes	Count	11	7	18
		% within Gender	26.8%	31.8%	28.6%
	Sometimes	Count	20	7	27
		% within Gender	48.8%	31.8%	42.9%
	No	Count	10	8	18
		% within Gender	24.4%	36.4%	28.6%
Total		Count	41	22	63
		% within Gender	100.0%	100.0%	100.0%

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	1.804 ^a	2	.406
Likelihood Ratio	1.825	2	.401
Linear-by-Linear Association	.120	1	.729
N of Valid Cases	63		

a. 0 cells (.0%) have expected count less than 5. The minimum expected count is 6.29.

Directional Measures

			Value	Asymptotic Standard Error ^a	Approximate T ^b
Ordinal by Ordinal	Somers' d	Symmetric	.041	.123	.332
		Get_Upset_Not_Able_to_Eat_Certain_Foods Dependent	.050	.150	.332
		Gender Dependent	.035	.104	.332

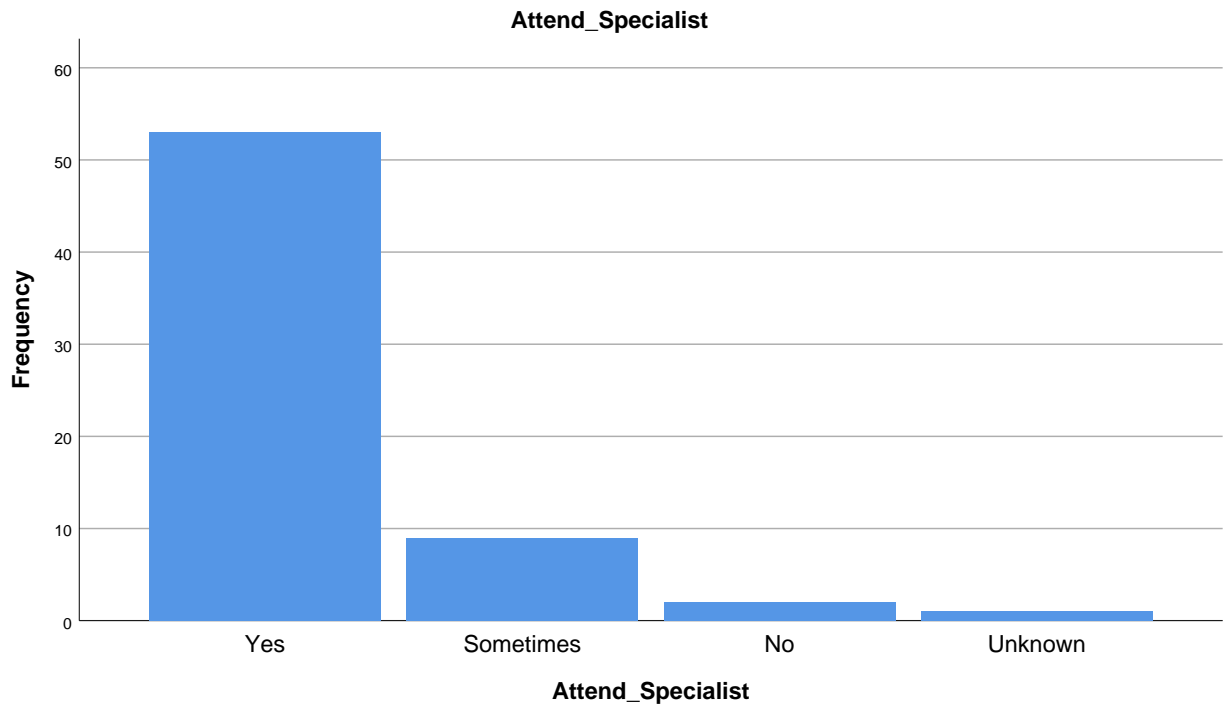
Directional Measures

			Approximate Significance
Ordinal by Ordinal	Somers' d	Symmetric	.740
		Get_Upset_Not_Able_to_E at_Certain_Foods Dependent	.740
		Gender Dependent	.740

- a. Not assuming the null hypothesis.
- b. Using the asymptotic standard error assuming the null hypothesis.

Attend Specialist

		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	53	81.5	81.5	81.5
	Sometimes	9	13.8	13.8	95.4
	No	2	3.1	3.1	98.5
	99	1	1.5	1.5	100.0
	Total	65	100.0	100.0	



NPAR TESTS

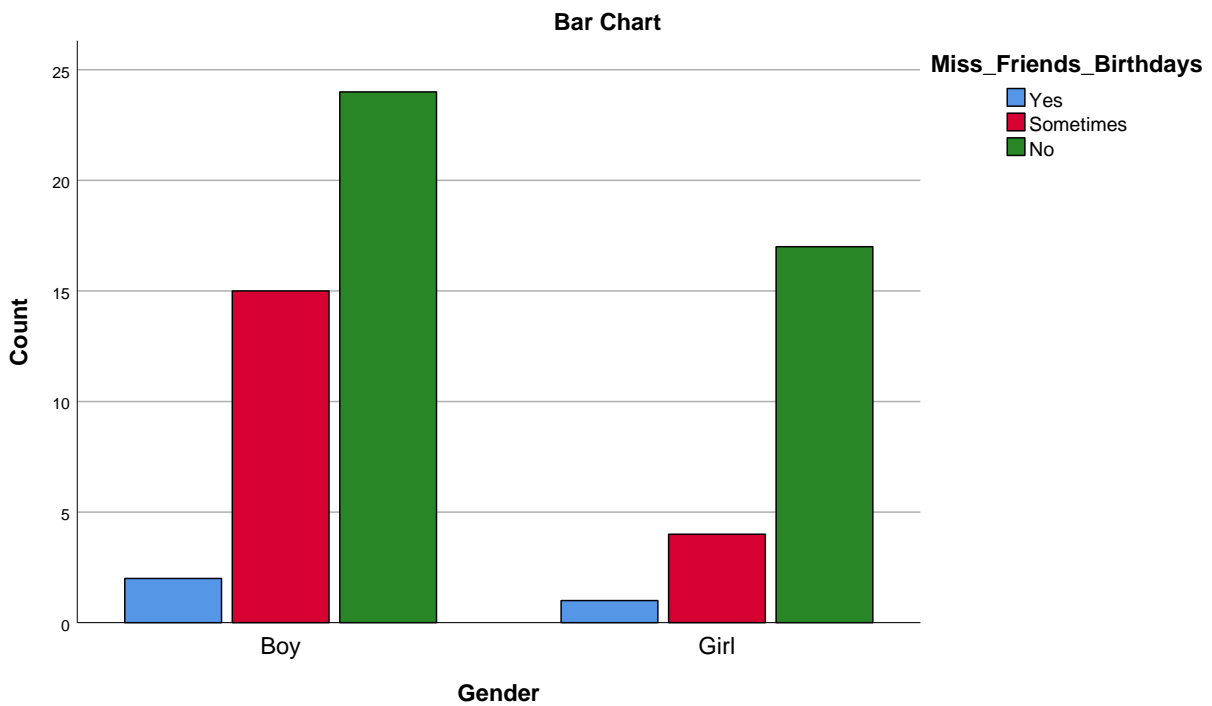
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/CHISQUARE=Miss_Friends_BirthdaysParents_Fuss Feel_Different_to_Friends  
Get_Upset_Not_Able_to_Eat_Certain_Foods  
/EXPECTED=EQUAL  
/MISSING ANALYSIS.
```

NPar Tests

Chi-Square Test - Equal Distribution

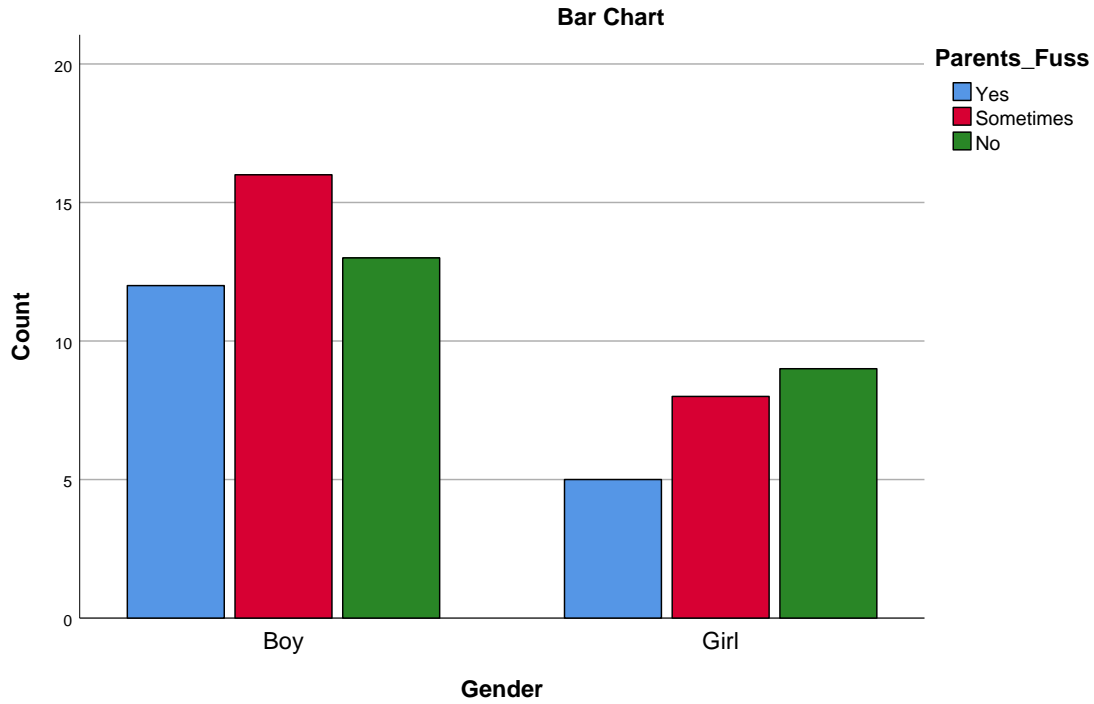
Frequencies

Miss Friends' Birthdays			
	Observed N	Expected N	Residual
Yes	3	21.3	-18.3
Sometimes	20	21.3	-1.3
No	41	21.3	19.7
Total	64		



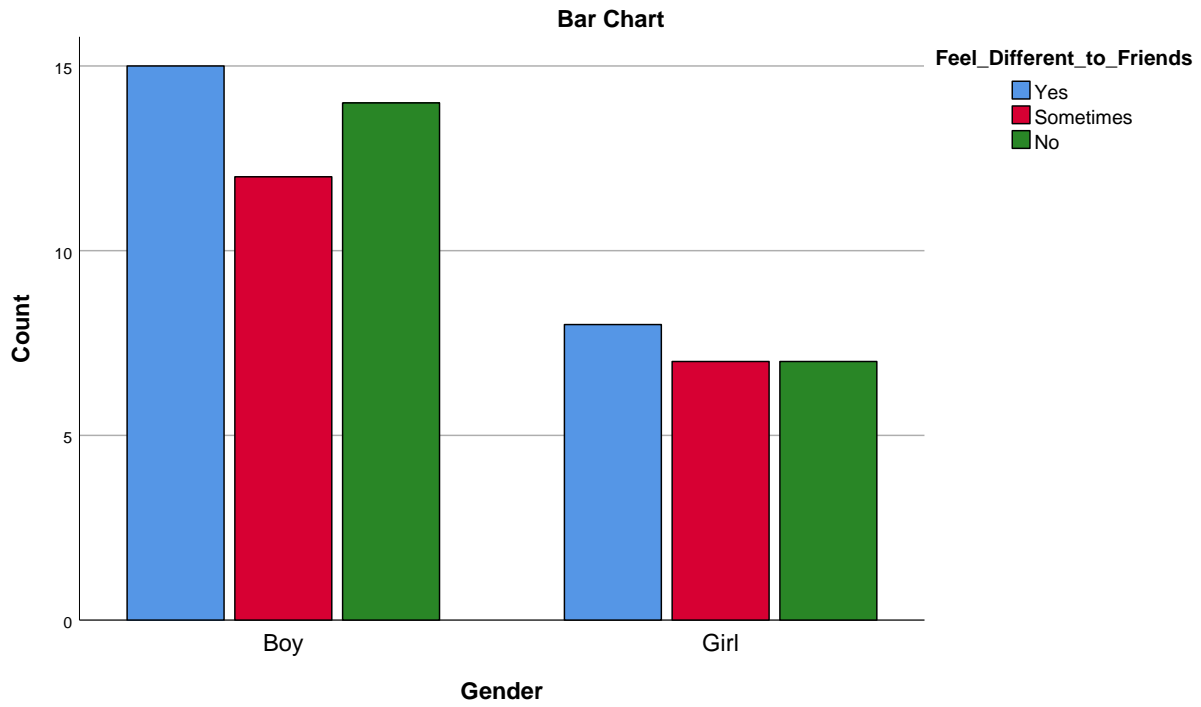
Parents Fuss

	Observed N	Expected N	Residual
Yes	18	21.3	-3.3
Sometimes	24	21.3	2.7
No	22	21.3	.7
Total	64		



Feel Different to Friends

	Observed N	Expected N	Residual
Yes	24	21.3	2.7
Sometimes	19	21.3	-2.3
No	21	21.3	-.3
Total	64		



Get Upset Not Able to Eat Certain Foods			
	Observed N	Expected N	Residual
Yes	18	21.3	-3.3
Sometimes	28	21.3	6.7
No	18	21.3	-3.3
Total	64		

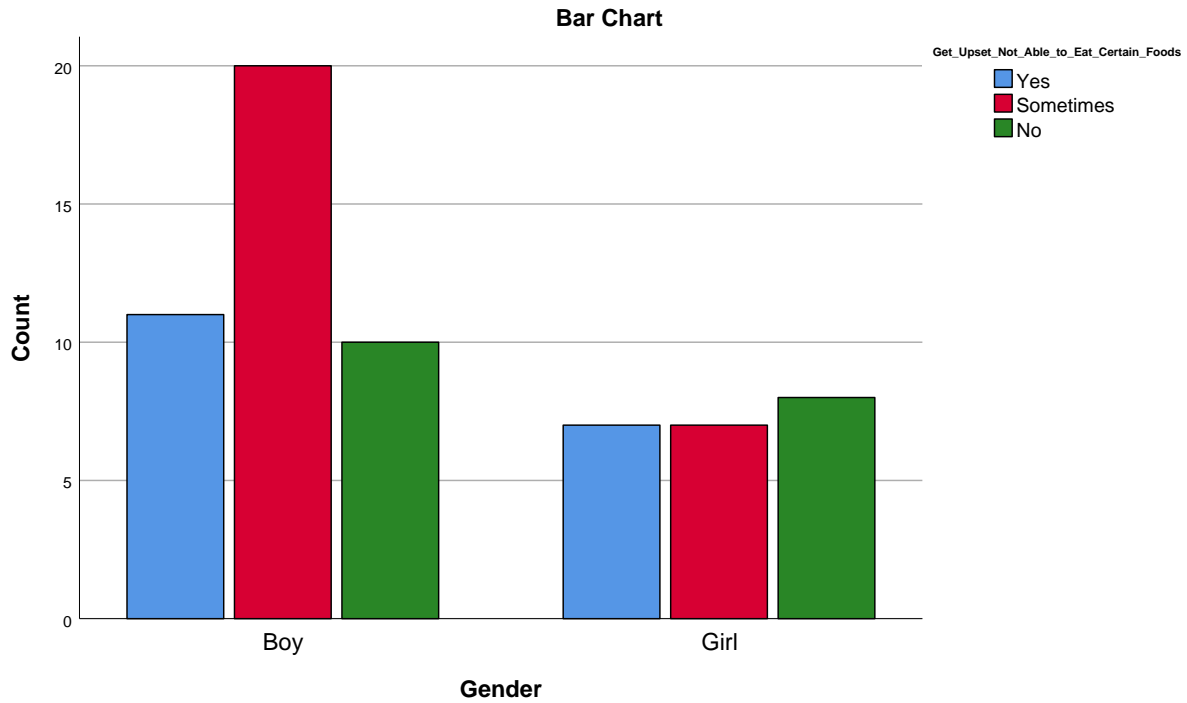


Table 4.1: Equal Distribution

	Miss Friend's Birthdays	Parents Fuss	Feel Different to Friends	Get Upset Not Able to Eat Certain Foods
Chi-Square	33.969 ^a	.875 ^a	.594 ^a	3.125 ^a
df	2	2	2	2
Asymp. Sig.	.000	.646	.743	.210

a. 0 cells (.0%) have expected frequencies less than 5. The minimum expected cell frequency is 21.3.

NPAR TESTS

```

/CHISQUARE=Miss_Friends_BirthdaysParents_Fuss Feel_Different_to_Friends
  Get_Upset_Not_Able_to_Eat_Certain_Foods
/EXPECTED=7 13 44
/MISSING ANALYSIS.

```

Chi-Square Test - QoL Impact

Frequencies

Miss Friends' Birthdays

	Observed N	Expected N	Residual
Yes	3	7.0	-4.0
Sometimes	20	13.0	7.0
No	41	44.0	-3.0
Total	64		

Parents Fuss

	Observed N	Expected N	Residual
Yes	18	7.0	11.0
Sometimes	24	13.0	11.0
No	22	44.0	-22.0
Total	64		

Feel Different to Friends

	Observed N	Expected N	Residual
Yes	24	7.0	17.0
Sometimes	19	13.0	6.0
No	21	44.0	-23.0
Total	64		

Get Upset Not Able to Eat Certain Foods

	Observed N	Expected N	Residual
Yes	18	7.0	11.0
Sometimes	28	13.0	15.0
No	18	44.0	-26.0
Total	64		

Test Statistics

	Miss_Friends_ Birthdays	Parents_Fuss	Feel_Different_t o_Friends	Get_Upset_Not _Able_to_Eat _Certain_Foods
Chi-Square	6.259 ^a	37.593 ^a	56.078 ^a	49.957 ^a
df	2	2	2	2
Asymp. Sig.	.044	.000	.000	.000

a. 0 cells (.0%) have expected frequencies less than 5. The minimum expected cell frequency is 7.0.

```

PLUM Get_Upset_Not_Able_to_Eat_Certain_Foods BY Gender
  /CRITERIA=CIN(95) DELTA(0) LCONVERGE(0) MXITER(100) MXSTEP(5) PCONVERGE(1.0E
-6) SINGULAR(1.0E-8)
  /LINK=LOGIT
  /PRINT=FIT PARAMETER SUMMARY TPARALLEL
  /SAVE=ESTPROB.

```

PLUM - Ordinal Regression - Get Upset Not Able to Eat Certain Foods

Case Processing Summary

		N	Marginal Percentage
Get_Upset_Not_Able_to_E at_Certain_Foods	Yes	18	28.6%
	Sometimes	27	42.9%
	No	18	28.6%
Gender	Boy	41	65.1%
	Girl	22	34.9%
Valid		63	100.0%
Missing		2	
Total		65	

Model Fitting Information

Model	-2 Log Likelihood	Chi-Square	df	Sig.
Intercept Only	16.137			
Final	16.010	.127	1	.721

Link function: Logit.

Goodness-of-Fit

	Chi-Square	df	Sig.
Pearson	1.671	1	.196
Deviance	1.698	1	.193

Link function: Logit.

Pseudo R-Square

Cox and Snell	.002
Nagelkerke	.002
McFadden	.001

Link function: Logit.

Parameter Estimates

		Estimate	Std. Error	Wald	df	Sig.
Threshold	[Get_Upset_Not_Able_to_E at_Certain_Foods = 1]	-1.039	.429	5.872	1	.015
	[Get_Upset_Not_Able_to_E at_Certain_Foods = 2]	.797	.419	3.617	1	.057
Location	[Gender=1 = Boy]	-.178	.490	.132	1	.716
	Girl	0 ^a	.	.	0	.

Parameter Estimates

		95% Confidence Interval	
		Lower Bound	Upper Bound
Threshold	[Get_Upset_Not_Able_to_E at_Certain_Foods = 1]	-1.879	-.199
	[Get_Upset_Not_Able_to_E at_Certain_Foods = 2]	-.024	1.618
Location	[Gender=1 = Boy]	-1.139	.782
	Girl	.	.

Link function: Logit.

- a. This parameter is set to zero because it is redundant.

Test of Parallel Lines^a

Model	-2 Log Likelihood	Chi-Square	df	Sig.
Null Hypothesis	16.010			
General	14.311	1.698	1	.193

The null hypothesis states that the location parameters (slope coefficients) are the same across response categories.

- a. Link function: Logit.

Get_Upset_Not_Able_to_Eat_Certain_Foods * Gender

Crosstab

		Gender		Total	
		Boy	Girl		
Get_Upset_Not_Able_to_Eat_Certain_Foods	Yes	Count	11	7	18
		Expected Count	11.7	6.3	18.0
		% within Get_Upset_Not_Able_to_Eat_Certain_Foods	61.1%	38.9%	100.0%
		% within Gender	26.8%	31.8%	28.6%
	Sometimes	Count	20	7	27
		Expected Count	17.6	9.4	27.0
		% within Get_Upset_Not_Able_to_Eat_Certain_Foods	74.1%	25.9%	100.0%
		% within Gender	48.8%	31.8%	42.9%
No	Count	10	8	18	
	Expected Count	11.7	6.3	18.0	

Crosstab

		Gender		
		Boy	Girl	Total
	% within Get_Upset_Not_Able_to_E at_Certain_Foods	55.6%	44.4%	100.0%
	% within Gender	24.4%	36.4%	28.6%
Total	Count	41	22	63
	Expected Count	41.0	22.0	63.0
	% within Get_Upset_Not_Able_to_E at_Certain_Foods	65.1%	34.9%	100.0%
	% within Gender	100.0%	100.0%	100.0%

Chi-Square Tests

	Value	df	Asymptotic Significance (2- sided)
Pearson Chi-Square	1.804 ^a	2	.406
Likelihood Ratio	1.825	2	.401
Linear-by-Linear Association	.120	1	.729
N of Valid Cases	63		

a. 0 cells (.0%) have expected count less than 5. The minimum expected count is 6.29.

Get_Upset_Not_Able_to_Eat_Certain_Foods * Other_Allergies

Crosstab

			Other_Allergies		Total
			Yes	No	
Get_Upset_Not_Able_to_E at_Certain_Foods	Yes	Count	17	1	18
		% within Get_Upset_Not_Able_to_E at_Certain_Foods	94.4%	5.6%	100.0%
		% within Other_Allergies	31.5%	11.1%	28.6%
		% of Total	27.0%	1.6%	28.6%
	Sometimes	Count	24	3	27
		% within Get_Upset_Not_Able_to_E at_Certain_Foods	88.9%	11.1%	100.0%
		% within Other_Allergies	44.4%	33.3%	42.9%
		% of Total	38.1%	4.8%	42.9%
	No	Count	13	5	18
		% within Get_Upset_Not_Able_to_E at_Certain_Foods	72.2%	27.8%	100.0%
		% within Other_Allergies	24.1%	55.6%	28.6%
		% of Total	20.6%	7.9%	28.6%
Total	Count	54	9	63	
	% within Get_Upset_Not_Able_to_E at_Certain_Foods	85.7%	14.3%	100.0%	
	% within Other_Allergies	100.0%	100.0%	100.0%	
	% of Total	85.7%	14.3%	100.0%	

Chi-Square Tests

	Value	df	Asymptotic Significance (2- sided)
Pearson Chi-Square	4.019 ^a	2	.134
Likelihood Ratio	3.843	2	.146
Linear-by-Linear Association	3.572	1	.059
N of Valid Cases	63		

a. 3 cells (50.0%) have expected count less than 5. The minimum expected count is 2.57.

```

PLUM Get_Upset_Not_Able_to_Eat_Certain_Foods BY Number_of_Allergens
  /CRITERIA=CIN(95) DELTA(0) LCONVERGE(0) MXITER(100) MXSTEP(5) PCONVERGE(1.0E
-6) SINGULAR(1.0E-8)
  /LINK=LOGIT
  /PRINT=FIT PARAMETER SUMMARY TPARALLEL

```

Case Processing Summary

		N	Marginal Percentage
Get_Upset_Not_Able_to_Eat_Certain_Foods	Yes	18	28.1%
	Sometimes	28	43.8%
	No	18	28.1%
Number_of_Allergens	1 Allergen	15	23.4%
	2 - 3 Allergens	22	34.4%
	4 Allergens or More	27	42.2%
Valid		64	100.0%
Missing		1	
Total		65	

Model Fitting Information

Model	-2 Log Likelihood	Chi-Square	df	Sig.
Intercept Only	26.208			
Final	21.086	5.123	2	.077

Link function: Logit.

Goodness-of-Fit

	Chi-Square	df	Sig.
Pearson	2.371	2	.306
Deviance	2.438	2	.295

Link function: Logit.

Pseudo R-Square

Cox and Snell	.077
Nagelkerke	.087
McFadden	.037

Link function: Logit.

Parameter Estimates

		Estimate	Std. Error	Wald	df	Sig.
Threshold	[Get_Upset_Not_Able_to_E at_Certain_Foods = 1]	-.806	.387	4.344	1	.037
	[Get_Upset_Not_Able_to_E at_Certain_Foods = 2]	1.197	.406	8.674	1	.003
Location	[Number_of_Allergens=1]	-.537	.608	.782	1	.377
	[Number_of_Allergens=2]	.844	.547	2.384	1	.123
	[Number_of_Allergens=3]	0 ^a	.	.	0	.

Parameter Estimates

		95% Confidence Interval	
		Lower Bound	Upper Bound
Threshold	[Get_Upset_Not_Able_to_E at_Certain_Foods = 1]	-1.563	-.048
	[Get_Upset_Not_Able_to_E at_Certain_Foods = 2]	.400	1.994
Location	[Number_of_Allergens=1]	-1.729	.654
	[Number_of_Allergens=2]	-.228	1.916
	[Number_of_Allergens=3]	.	.

Link function: Logit.

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

Test of Parallel Lines^a

Model	-2 Log Likelihood	Chi-Square	df	Sig.
Null Hypothesis	21.086			
General	18.647	2.438	2	.295

The null hypothesis states that the location parameters (slope coefficients) are the same across response categories.

a. Link function: Logit.

CROSSTABS

```

/TABLES=Get_Upset_Not_Able_to_Eat_Certain_FoodBY Number_of_Allergens
/SHOWDIM=2
/FORMAT=AVALUE TABLES
/STATISTICS=CHISQ
/CELLS=COUNT COLUMN
/COUNT ROUND CELL.

```

Crosstabs

Case Processing Summary

	Valid		Cases Missing		Total	
	N	Percent	N	Percent	N	Percent
Get_Upset_Not_Able_to_Eat_Certain_Foods * Number_of_Allergens	64	98.5%	1	1.5%	65	100.0%

Get_Upset_Not_Able_to_Eat_Certain_Foods * Number_of_Allergens Crosstabulation

			Number_of_Allergens	
			1 Allergen	2 - 3 Allergens
Get_Upset_Not_Able_to_Eat_Certain_Foods	Yes	Count	7	2
		% within Number_of_Allergens	46.7%	8.7%
	Sometimes	Count	5	12
		% within Number_of_Allergens	33.3%	52.2%
	No	Count	3	9
		% within Number_of_Allergens	20.0%	39.1%
Total	Count	15	23	
	% within Number_of_Allergens	100.0%	100.0%	

Get_Upset_Not_Able_to_Eat_Certain_Foods * Number_of_Allergens Crosstabulation

				Number_of_Aller...	
				4 Allergens or More	Total
Get_Upset_Not_Able_to_Eat_Certain_Foods	Yes	Count		9	18
		% within Number_of_Allergens		34.6%	28.1%
	Sometimes	Count		11	28
		% within Number_of_Allergens		42.3%	43.8%
	No	Count		6	18
		% within Number_of_Allergens		23.1%	28.1%
Total		Count		26	64
		% within Number_of_Allergens		100.0%	100.0%

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	7.646 ^a	4	.105
Likelihood Ratio	8.426	4	.077
Linear-by-Linear Association	.045	1	.832
N of Valid Cases	64		

a. 2 cells (22.2%) have expected count less than 5. The minimum expected count is 4.22.

```

PLUM Feel_Different_to_FriendsBY Gender
/CRITERIA=CIN(95) DELTA(0) LCONVERGE(0) MXITER(100) MXSTEP(5) PCONVERGE(1.0E
-6) SINGULAR(1.0E-8)
/LINK=LOGIT
/PRINT=FIT PARAMETER SUMMARY TPARALLEL
/SAVE=ESTPROB.
    
```

PLUM - Ordinal Regression - Feel Different to Friends

Case Processing Summary

		N	Marginal Percentage
Feel_Different_to_Friends	Yes	23	36.5%
	Sometimes	19	30.2%
	No	21	33.3%
Gender	Boy	41	65.1%
	Girl	22	34.9%
Valid		63	100.0%
Missing		2	
Total		65	

Model Fitting Information

Model	-2 Log Likelihood	Chi-Square	df	Sig.
Intercept Only	14.498			
Final	14.490	.009	1	.926

Link function: Logit.

Goodness-of-Fit

	Chi-Square	df	Sig.
Pearson	.046	1	.830
Deviance	.046	1	.831

Link function: Logit.

Pseudo R-Square

Cox and Snell	.000
Nagelkerke	.000
McFadden	.000

Link function: Logit.

Parameter Estimates

		Estimate	Std. Error	Wald	df	Sig.
Threshold	[Feel_Different_to_Friends = 1]	-.525	.410	1.637	1	.201
	[Feel_Different_to_Friends = 2]	.722	.415	3.028	1	.082
Location	[Gender=1 = Boys]	.045	.486	.008	1	.927
	Girls	0 ^a	.	.	0	.

Parameter Estimates

		95% Confidence Interval	
		Lower Bound	Upper Bound
Threshold	[Feel_Different_to_Friends = 1]	-1.328	.279
	[Feel_Different_to_Friends = 2]	-.091	1.536
Location	[Gender=1 = Boys]	-.908	.998
	Girls	.	.

Link function: Logit.

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

Test of Parallel Lines^a

Model	-2 Log Likelihood	Chi-Square	df	Sig.
Null Hypothesis	14.490			
General	14.444	.046	1	.831

The null hypothesis states that the location parameters (slope coefficients) are the same across response categories.

a. Link function: Logit.

Feel_Different_to_Friends * Gender

Crosstab

Count

		Gender		Total
		Boy	Girl	
Feel_Different_to_Friends	Yes	15	8	23
	Sometimes	12	7	19
	No	14	7	21
Total		41	22	63

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	.054 ^a	2	.973
Likelihood Ratio	.054	2	.973
Linear-by-Linear Association	.009	1	.925
N of Valid Cases	63		

a. 0 cells (.0%) have expected count less than 5. The minimum expected count is 6.63.

Feel_Different_to_Friends * Other_Allergies

Crosstab

			Other_Allergies		Total
			Yes	No	
Feel_Different_to_Friends	Yes	Count	21	2	23
		% within Feel_Different_to_Friends	91.3%	8.7%	100.0%
		% within Other_Allergies	38.9%	22.2%	36.5%
		% of Total	33.3%	3.2%	36.5%
	Sometimes	Count	17	2	19
		% within Feel_Different_to_Friends	89.5%	10.5%	100.0%
		% within Other_Allergies	31.5%	22.2%	30.2%
		% of Total	27.0%	3.2%	30.2%
	No	Count	16	5	21
		% within Feel_Different_to_Friends	76.2%	23.8%	100.0%
		% within Other_Allergies	29.6%	55.6%	33.3%
		% of Total	25.4%	7.9%	33.3%
Total	Count	54	9	63	
	% within Feel_Different_to_Friends	85.7%	14.3%	100.0%	
	% within Other_Allergies	100.0%	100.0%	100.0%	
	% of Total	85.7%	14.3%	100.0%	

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	2.362 ^a	2	.307
Likelihood Ratio	2.245	2	.325
Linear-by-Linear Association	1.975	1	.160
N of Valid Cases	63		

a. 3 cells (50.0%) have expected count less than 5. The minimum expected count is 2.71.

```

PLUM Feel_Different_to_FriendsBY Number_of_Allergens
  /CRITERIA=CIN(95) DELTA(0) LCONVERGE(0) MXITER(100) MXSTEP(5) PCONVERGE(1.0E
-6) SINGULAR(1.0E-8)
  /LINK=LOGIT
  /PRINT=FIT PARAMETER SUMMARY TPARALLEL

```

Case Processing Summary

		N	Marginal Percentage
Feel_Different_to_Friends	Yes	24	37.5%
	Sometimes	19	29.7%
	No	21	32.8%
Number_of_Allergens	1 Allergen	15	23.4%
	2 - 3 Allergens	22	34.4%
	4 Allergens or More	27	42.2%
Valid		64	100.0%
Missing		1	
Total		65	

Model Fitting Information

Model	-2 Log Likelihood	Chi-Square	df	Sig.
Intercept Only	21.923			
Final	20.173	1.750	2	.417

Link function: Logit.

Goodness-of-Fit

	Chi-Square	df	Sig.
Pearson	.917	2	.632
Deviance	.928	2	.629

Link function: Logit.

Pseudo R-Square

Cox and Snell	.027
Nagelkerke	.030
McFadden	.012

Link function: Logit.

Parameter Estimates

		Estimate	Std. Error	Wald	df	Sig.
Threshold	[Feel_Different_to_Friends = 1]	-.545	.374	2.123	1	.145
	[Feel_Different_to_Friends = 2]	.710	.379	3.504	1	.061
Location	[Number_of_Allergens=1]	-.559	.604	.855	1	.355
	[Number_of_Allergens=2]	.274	.529	.267	1	.605
	[Number_of_Allergens=3]	0 ^a	.	.	0	.

Parameter Estimates

		95% Confidence Interval	
		Lower Bound	Upper Bound
Threshold	[Feel_Different_to_Friends = 1]	-1.279	.188
	[Feel_Different_to_Friends = 2]	-.033	1.453
Location	[Number_of_Allergens=1]	-1.743	.626
	[Number_of_Allergens=2]	-.764	1.311
	[Number_of_Allergens=3]	.	.

Link function: Logit.

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

Test of Parallel Lines^a

Model	-2 Log Likelihood	Chi-Square	df	Sig.
Null Hypothesis	20.173			
General	19.245	.928	2	.629

The null hypothesis states that the location parameters (slope coefficients) are the same across response categories.

a. Link function: Logit.

CROSSTABS

```

/TABLES=Feel_Different_to_FriendsBY Number_of_Allergens
/SHOWDIM=2
/FORMAT=AVALUE TABLES
/STATISTICS=CHISQ
/CELLS=COUNT COLUMN
/COUNT ROUND CELL.

```

Crosstabs

Case Processing Summary

	Valid		Cases Missing		Total	
	N	Percent	N	Percent	N	Percent
Feel_Different_to_Friends * Number_of_Allergens	64	98.5%	1	1.5%	65	100.0%

Feel_Different_to_Friends * Number_of_Allergens Crosstabulation

				Number_of_Allergens	
				1 Allergen	2 - 3 Allergens
Feel_Different_to_Friends	Yes	Count	8	6	
		% within Number_of_Allergens	53.3%	26.1%	
	Sometimes	Count	3	8	
		% within Number_of_Allergens	20.0%	34.8%	
	No	Count	4	9	
		% within Number_of_Allergens	26.7%	39.1%	
Total	Count	15	23		
	% within Number_of_Allergens	100.0%	100.0%		

Feel_Different_to_Friends * Number_of_Allergens Crosstabulation

		Number_of_Aller...		
		4 Allergens or More	Total	
Feel_Different_to_Friends	Yes	Count	10	24
		% within Number_of_Allergens	38.5%	37.5%
	Sometimes	Count	8	19
		% within Number_of_Allergens	30.8%	29.7%
	No	Count	8	21
		% within Number_of_Allergens	30.8%	32.8%
Total		Count	26	64
		% within Number_of_Allergens	100.0%	100.0%

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	2.979 ^a	4	.561
Likelihood Ratio	3.006	4	.557
Linear-by-Linear Association	.227	1	.634
N of Valid Cases	64		

a. 2 cells (22.2%) have expected count less than 5. The minimum expected count is 4.45.

```

PLUM Miss_Friends_BirthdaysBY Gender
  /CRITERIA=CIN(95) DELTA(0) LCONVERGE(0) MXITER(100) MXSTEP(5) PCONVERGE(1.0E
-6) SINGULAR(1.0E-8)
  /LINK=LOGIT
  /PRINT=CELLINFO FIT PARAMETER SUMMARY
  /SAVE=ESTPROB PREDCAT PCPROB ACPROB.
    
```

PLUM - Ordinal Regression - Missing Friends' Birthdays

Case Processing Summary

		N	Marginal Percentage
Miss_Friends_Birthdays	Yes	3	4.8%
	Sometimes	19	30.2%
	No	41	65.1%
Gender	Boy	41	65.1%
	Girl	22	34.9%
Valid		63	100.0%
Missing		2	
Total		65	

Model Fitting Information

Model	-2 Log Likelihood	Chi-Square	df	Sig.
Intercept Only	14.151			
Final	12.059	2.092	1	.148

Link function: Logit.

Goodness-of-Fit

	Chi-Square	df	Sig.
Pearson	.460	1	.498
Deviance	.411	1	.521

Link function: Logit.

Pseudo R-Square

Cox and Snell	.033
Nagelkerke	.041
McFadden	.021

Link function: Logit.

Parameter Estimates

		Estimate	Std. Error	Wald	df	Sig.
Threshold	[Miss_Friends_Birthdays = 1]	-3.610	.750	23.148	1	.000
	[Miss_Friends_Birthdays = 2]	-1.198	.504	5.655	1	.017
Location	[Gender=1]	-0.835	.593	1.983	1	.159
	[Gender=2]	0 ^a	.	.	0	.

Parameter Estimates

		95% Confidence Interval	
		Lower Bound	Upper Bound
Threshold	[Miss_Friends_Birthdays = 1]	-5.081	-2.140
	[Miss_Friends_Birthdays = 2]	-2.185	-.211
Location	[Gender=1]	-1.998	.327
	[Gender=2]	.	.

Link function: Logit.

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

Miss_Friends_Birthdays * Gender

Cell Information

Frequency

Gender		Miss_Friends_Birthdays		
		Yes	Sometimes	No
Boy	Observed	2	15	24
	Expected	2.407	14.416	24.177
	Pearson Residual	-.270	.191	-.056
Girl	Observed	1	4	17
	Expected	.579	4.521	16.900
	Pearson Residual	.560	-.275	.051

Link function: Logit.

Crosstab

		Gender			
		Boy	Girl	Total	
Miss_Friends_Birthdays	Yes	Count	2	1	3
		Expected Count	2.0	1.0	3.0
		% within Miss_Friends_Birthdays	66.7%	33.3%	100.0%
		% within Gender	4.9%	4.5%	4.8%
	Sometimes	Count	15	4	19
		Expected Count	12.4	6.6	19.0
		% within Miss_Friends_Birthdays	78.9%	21.1%	100.0%
		% within Gender	36.6%	18.2%	30.2%
	No	Count	24	17	41
		Expected Count	26.7	14.3	41.0
		% within Miss_Friends_Birthdays	58.5%	41.5%	100.0%
		% within Gender	58.5%	77.3%	65.1%
Total	Count	41	22	63	
	Expected Count	41.0	22.0	63.0	
	% within Miss_Friends_Birthdays	65.1%	34.9%	100.0%	
	% within Gender	100.0%	100.0%	100.0%	

Chi-Square Tests

	Value	df	Asymptotic Significance (2- sided)
Pearson Chi-Square	2.384 ^a	2	.304
Likelihood Ratio	2.503	2	.286
Linear-by-Linear Association	1.531	1	.216
N of Valid Cases	63		

a. 2 cells (33.3%) have expected count less than 5. The minimum expected count is 1.05.

Miss_Friends_Birthdays * Other_Allergies

Crosstab

			Other_Allergies		
			Yes	No	Total
Miss_Friends_Birthdays	Yes	Count	3	0	3
		% within Miss_Friends_Birthdays	100.0%	0.0%	100.0%
		% within Other_Allergies	5.6%	0.0%	4.8%
		% of Total	4.8%	0.0%	4.8%
	Sometimes	Count	17	2	19
		% within Miss_Friends_Birthdays	89.5%	10.5%	100.0%
		% within Other_Allergies	31.5%	22.2%	30.2%
		% of Total	27.0%	3.2%	30.2%
	No	Count	34	7	41
		% within Miss_Friends_Birthdays	82.9%	17.1%	100.0%
		% within Other_Allergies	63.0%	77.8%	65.1%
		% of Total	54.0%	11.1%	65.1%
Total	Count	54	9	63	
	% within Miss_Friends_Birthdays	85.7%	14.3%	100.0%	
	% within Other_Allergies	100.0%	100.0%	100.0%	
	% of Total	85.7%	14.3%	100.0%	

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	.979 ^a	2	.613
Likelihood Ratio	1.410	2	.494
Linear-by-Linear Association	.942	1	.332
N of Valid Cases	63		

a. 3 cells (50.0%) have expected count less than 5. The minimum expected count is .43.

PLUM Miss_Friends_BirthdaysBY Number_of_Allergens
 /CRITERIA=CIN(95) DELTA(0) LCONVERGE(0) MXITER(100) MXSTEP(5) PCONVERGE(1.0E

-6) SINGULAR(1.0E-8)
 /LINK=LOGIT
 /PRINT=FIT PARAMETER SUMMARY TPARALLEL
 /SAVE=ESTPROB.

Case Processing Summary

		N	Marginal Percentage
Miss_Friends_Birthdays	Yes	3	4.7%
	Sometimes	20	31.3%
	No	41	64.1%
Number_of_Allergens	1 Allergen	15	23.4%
	2 - 3 Allergens	22	34.4%
	4 Allergens or More	27	42.2%
Valid		64	100.0%
Missing		1	
Total		65	

Model Fitting Information

Model	-2 Log Likelihood	Chi-Square	df	Sig.
Intercept Only	17.545			
Final	17.371	.174	2	.916

Link function: Logit.

Goodness-of-Fit

	Chi-Square	df	Sig.
Pearson	2.014	2	.365
Deviance	2.884	2	.236

Link function: Logit.

Pseudo R-Square

Cox and Snell	.003
Nagelkerke	.003
McFadden	.002

Link function: Logit.

Parameter Estimates

		Estimate	Std. Error	Wald	df	Sig.
Threshold	[Miss_Friends_Birthdays = 1]	-3.083	.669	21.211	1	.000
	[Miss_Friends_Birthdays = 2]	-.645	.403	2.562	1	.109
Location	[Number_of_Allergens=1]	-.258	.655	.155	1	.694
	[Number_of_Allergens=2]	-.014	.598	.001	1	.981
	[Number_of_Allergens=3]	0 ^a	.	.	0	.

Parameter Estimates

		95% Confidence Interval	
		Lower Bound	Upper Bound
Threshold	[Miss_Friends_Birthdays = 1]	-4.394	-1.771
	[Miss_Friends_Birthdays = 2]	-1.434	.145
Location	[Number_of_Allergens=1]	-1.542	1.026
	[Number_of_Allergens=2]	-1.186	1.157
	[Number_of_Allergens=3]	.	.

Link function: Logit.

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

Test of Parallel Lines^a

Model	-2 Log Likelihood	Chi-Square	df	Sig.
Null Hypothesis	17.371			
General	14.487	2.884	2	.236

The null hypothesis states that the location parameters (slope coefficients) are the same across response categories.

a. Link function: Logit.

CROSSTABS

```

/TABLES=Miss_Friends_BirthdaysBY Number_of_Allergens
/SHOWDIM=2
/FORMAT=AVALUE TABLES
/STATISTICS=CHISQ

```

/CELLS=COUNT COLUMN
 /COUNT ROUND CELL.

Crosstabs

Case Processing Summary

	Valid		Cases Missing		Total	
	N	Percent	N	Percent	N	Percent
Miss_Friends_Birthdays * Number_of_Allergens	64	98.5%	1	1.5%	65	100.0%

Miss_Friends_Birthdays * Number_of_Allergens Crosstabulation

				Number_of_Allergens	
				1 Allergen	2 - 3 Allergens
Miss_Friends_Birthdays	Yes	Count	1	0	
		% within Number_of_Allergens	6.7%	0.0%	
	Sometimes	Count	5	8	
		% within Number_of_Allergens	33.3%	34.8%	
	No	Count	9	15	
		% within Number_of_Allergens	60.0%	65.2%	
Total		Count	15	23	
		% within Number_of_Allergens	100.0%	100.0%	

Miss_Friends_Birthdays * Number_of_Allergens Crosstabulation

		Number_of_Aller...		
		4 Allergens or More	Total	
Miss_Friends_Birthdays	Yes	Count	2	3
		% within Number_of_Allergens	7.7%	4.7%
	Sometimes	Count	7	20
		% within Number_of_Allergens	26.9%	31.3%
	No	Count	17	41
		% within Number_of_Allergens	65.4%	64.1%
Total		Count	26	64
		% within Number_of_Allergens	100.0%	100.0%

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	2.023 ^a	4	.731
Likelihood Ratio	3.009	4	.556
Linear-by-Linear Association	.017	1	.898
N of Valid Cases	64		

a. 4 cells (44.4%) have expected count less than 5. The minimum expected count is .70.

PLUM - Ordinal Regression - Parents Fuss

```

PLUM Parents_Fuss BY Gender
  /CRITERIA=CIN(95) DELTA(0) LCONVERGE(0) MXITER(100) MXSTEP(5) PCONVERGE(1.0E
-6) SINGULAR(1.0E-8)
  /LINK=LOGIT
  /PRINT=CELLINFO FIT PARAMETER SUMMARY
  /SAVE=ESTPROB PREDCAT PCPROB ACPROB.
    
```

Case Processing Summary

		N	Marginal Percentage
Parents_Fuss	Yes	17	27.0%
	Sometimes	24	38.1%
	No	22	34.9%
Gender	Boy	41	65.1%
	Girl	22	34.9%
Valid		63	100.0%
Missing		2	
Total		65	

Model Fitting Information

Model	-2 Log Likelihood	Chi-Square	df	Sig.
Intercept Only	14.953			
Final	14.362	.590	1	.442

Link function: Logit.

Goodness-of-Fit

	Chi-Square	df	Sig.
Pearson	.009	1	.925
Deviance	.009	1	.925

Link function: Logit.

Pseudo R-Square

Cox and Snell	.009
Nagelkerke	.011
McFadden	.004

Link function: Logit.

Parameter Estimates

		Estimate	Std. Error	Wald	df	Sig.
Threshold	[Parents_Fuss = 1]	-1.249	.438	8.119	1	.004
	[Parents_Fuss = 2]	.381	.409	.870	1	.351
Location	[Gender=1] = Boy	-.376	.491	.589	1	.443
	Girl	0 ^a	.	.	0	.

Parameter Estimates

		95% Confidence Interval	
		Lower Bound	Upper Bound
Threshold	[Parents_Fuss = 1]	-2.107	-.390
	[Parents_Fuss = 2]	-.420	1.183
Location	[Gender=1] = Boy	-1.338	.585
	Girl	.	.

Link function: Logit.

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

Parents_Fuss * Gender

Cell Information

Frequency

Gender		Parents_Fuss		
		Yes	Sometimes	No
Boy	Observed	12	16	13
	Expected	12.087	15.830	13.084
	Pearson Residual	-.030	.055	-.028
Girl	Observed	5	8	9
	Expected	4.905	8.168	8.927
	Pearson Residual	.049	-.074	.032

Link function: Logit.

Crosstab

			Gender		
			Boy	Girl	Total
Parents_Fuss	Yes	Count	12	5	17
		Expected Count	11.1	5.9	17.0
		% within Parents_Fuss	70.6%	29.4%	100.0%
		% within Gender	29.3%	22.7%	27.0%
	Sometimes	Count	16	8	24
		Expected Count	15.6	8.4	24.0
		% within Parents_Fuss	66.7%	33.3%	100.0%
		% within Gender	39.0%	36.4%	38.1%
	No	Count	13	9	22
		Expected Count	14.3	7.7	22.0
		% within Parents_Fuss	59.1%	40.9%	100.0%
		% within Gender	31.7%	40.9%	34.9%
Total	Count	41	22	63	
	Expected Count	41.0	22.0	63.0	
	% within Parents_Fuss	65.1%	34.9%	100.0%	
	% within Gender	100.0%	100.0%	100.0%	

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	.601 ^a	2	.741
Likelihood Ratio	.599	2	.741
Linear-by-Linear Association	.570	1	.450
N of Valid Cases	63		

a. 0 cells (.0%) have expected count less than 5. The minimum expected count is 5.94.

Parents_Fuss * Other_Allergies

Crosstab

			Other_Allergies		
			Yes	No	Total
Parents_Fuss	Yes	Count	16	1	17
		% within Parents_Fuss	94.1%	5.9%	100.0%
		% within Other_Allergies	29.6%	11.1%	27.0%
		% of Total	25.4%	1.6%	27.0%
	Sometimes	Count	20	4	24
		% within Parents_Fuss	83.3%	16.7%	100.0%
		% within Other_Allergies	37.0%	44.4%	38.1%
		% of Total	31.7%	6.3%	38.1%
	No	Count	18	4	22
		% within Parents_Fuss	81.8%	18.2%	100.0%
		% within Other_Allergies	33.3%	44.4%	34.9%
		% of Total	28.6%	6.3%	34.9%
Total	Count	54	9	63	
	% within Parents_Fuss	85.7%	14.3%	100.0%	
	% within Other_Allergies	100.0%	100.0%	100.0%	
	% of Total	85.7%	14.3%	100.0%	

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	1.364 ^a	2	.506
Likelihood Ratio	1.579	2	.454
Linear-by-Linear Association	1.088	1	.297
N of Valid Cases	63		

a. 3 cells (50.0%) have expected count less than 5. The minimum expected count is 2.43.

```

PLUM Parents_Fuss BY Number_of_Allergens
  /CRITERIA=CIN(95) DELTA(0) LCONVERGE(0) MXITER(100) MXSTEP(5) PCONVERGE(1.0E
-6) SINGULAR(1.0E-8)
  /LINK=LOGIT
  /PRINT=FIT PARAMETER SUMMARY TPARALLEL
    
```

/SAVE=ESTPROB.

Case Processing Summary

		N	Marginal Percentage
Parents_Fuss	Yes	18	28.1%
	Sometimes	24	37.5%
	No	22	34.4%
Number_of_Allergens	1 Allergen	15	23.4%
	2 - 3 Allergens	22	34.4%
	4 Allergens or More	27	42.2%
Valid		64	100.0%
Missing		1	
Total		65	

Model Fitting Information

Model	-2 Log Likelihood	Chi-Square	df	Sig.
Intercept Only	24.903			
Final	23.071	1.832	2	.400

Link function: Logit.

Goodness-of-Fit

	Chi-Square	df	Sig.
Pearson	3.939	2	.140
Deviance	4.095	2	.129

Link function: Logit.

Pseudo R-Square

Cox and Snell	.028
Nagelkerke	.032
McFadden	.013

Link function: Logit.

Parameter Estimates

		Estimate	Std. Error	Wald	df	Sig.
Threshold	[Parents_Fuss = 1]	-.899	.387	5.404	1	.020
	[Parents_Fuss = 2]	.723	.380	3.623	1	.057
Location	[Number_of_Allergens=1]	-.448	.597	.565	1	.452
	[Number_of_Allergens=2]	.403	.533	.572	1	.450
	[Number_of_Allergens=3]	0 ^a	.	.	0	.

Parameter Estimates

		95% Confidence Interval	
		Lower Bound	Upper Bound
Threshold	[Parents_Fuss = 1]	-1.657	-.141
	[Parents_Fuss = 2]	-.021	1.468
Location	[Number_of_Allergens=1]	-1.618	.721
	[Number_of_Allergens=2]	-.642	1.449
	[Number_of_Allergens=3]	.	.

Link function: Logit.

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

Test of Parallel Lines^a

Model	-2 Log Likelihood	Chi-Square	df	Sig.
Null Hypothesis	23.071			
General	18.976	4.095	2	.129

The null hypothesis states that the location parameters (slope coefficients) are the same across response categories.

a. Link function: Logit.

CROSSTABS

```

/TABLES=Parents_Fuss BY Number_of_Allergens
/SHOWDIM=2
/FORMAT=AVALUE TABLES
/STATISTICS=CHISQ
/CELLS=COUNT COLUMN
/COUNT ROUND CELL.

```

Crosstabs

Case Processing Summary

	Valid		Cases Missing		Total	
	N	Percent	N	Percent	N	Percent
Parents_Fuss * Number_of_Allergens	64	98.5%	1	1.5%	65	100.0%

Parents_Fuss * Number_of_Allergens Crosstabulation

			Number_of_Allergens		
			1 Allergen	2 - 3 Allergens	4 Allergens or More
Parents_Fuss	Yes	Count	7	3	8
		% within Number_of_Allergens	46.7%	13.0%	30.8%
	Sometimes	Count	3	12	9
		% within Number_of_Allergens	20.0%	52.2%	34.6%
	No	Count	5	8	9
		% within Number_of_Allergens	33.3%	34.8%	34.6%
Total	Count	15	23	26	
	% within Number_of_Allergens	100.0%	100.0%	100.0%	

Parents_Fuss * Number_of_Allergens Crosstabulation

			Total
Parents_Fuss	Yes	Count	18
		% within Number_of_Allergens	28.1%
	Sometimes	Count	24
		% within Number_of_Allergens	37.5%
	No	Count	22
		% within Number_of_Allergens	34.4%
Total	Count	64	
	% within Number_of_Allergens	100.0%	

Chi-Square Tests

	Value	df	Asymptotic Significance (2-sided)
Pearson Chi-Square	6.368 ^a	4	.173
Likelihood Ratio	6.636	4	.156
Linear-by-Linear Association	.217	1	.642
N of Valid Cases	64		

a. 1 cells (11.1%) have expected count less than 5. The minimum expected count is 4.22.

RELIABILITY

```

/VARIABLES=Miss_Friends_BirthdaysParents_Fuss Feel_Different_to_Friends
  Get_Upset_Not_Able_to_Eat_Certain_Foods
/SCALE('ALL VARIABLES') ALL
/MODEL=ALPHA
/STATISTICS=DESCRIPTIVE
/SUMMARY=CORR.

```

Reliability

Scale: ALL VARIABLES

Case Processing Summary

		N	%
Cases	Valid	64	98.5
	Excluded ^a	1	1.5
	Total	65	100.0

a. Listwise deletion based on all variables in the procedure.

Reliability Statistics

Cronbach's Alpha	Cronbach's Alpha Based on Standardized Items	N of Items
.625	.618	4

Item Statistics

	Mean	Std. Deviation	N
Miss_Friends_Birthdays	2.59	.583	64
Parents_Fuss	2.06	.794	64
Feel_Different_to_Friends	1.95	.844	64
Get_Upset_Not_Able_to_Eat_Certain_Foods	2.00	.756	64

Summary Item Statistics

	Mean	Minimum	Maximum	Range	Maximum / Minimum	Variance
Inter-Item Correlations	.288	.124	.454	.330	3.659	.011

Summary Item Statistics

	N of Items
Inter-Item Correlations	4

FACTOR

```

/VARIABLES Miss_Friends_BirthdaysParents_Fuss Feel_Different_to_Friends
  Get_Upset_Not_Able_to_Eat_Certain_Foods
/MISSING LISTWISE
/ANALYSIS Miss_Friends_BirthdaysParents_Fuss Feel_Different_to_Friends
  Get_Upset_Not_Able_to_Eat_Certain_Foods
/PRINT INITIAL CORRELATION EXTRACTION
/CRITERIA MINEIGEN(1) ITERATE(25)
/EXTRACTION PC
/ROTATION NOROTATE
/METHOD=CORRELATION

```

Factor Analysis

Correlation Matrix

		Miss_Friends_ Birthdays	Parents_Fuss	Feel_Different_t o_Friends
Correlation	Miss_Friends_Birthdays	1.000	.124	.283
	Parents_Fuss	.124	1.000	.454
	Feel_Different_to_Friends	.283	.454	1.000
	Get_Upset_Not_Able_to_E at_Certain_Foods	.252	.264	.348

Correlation Matrix

		Get_Upset_Not _Able_to_Eat_ Certain_Foods
Correlation	Miss_Friends_Birthdays	.252
	Parents_Fuss	.264
	Feel_Different_to_Friends	.348
	Get_Upset_Not_Able_to_E at_Certain_Foods	1.000

Communalities

	Initial	Extraction
Miss_Friends_Birthdays	1.000	.300
Parents_Fuss	1.000	.480
Feel_Different_to_Friends	1.000	.641
Get_Upset_Not_Able_to_E at_Certain_Foods	1.000	.462

Extraction Method: Principal Component Analysis.

Total Variance Explained

Component	Total	Initial Eigenvalues		Extraction Sums of Squared Loadings		
		% of Variance	Cumulative %	Total	% of Variance	Cumulative %
1	1.883	47.072	47.072	1.883	47.072	47.072
2	.897	22.425	69.497			
3	.709	17.717	87.214			
4	.511	12.786	100.000			

Extraction Method: Principal Component Analysis.

Component Matrix^a

	Component 1
Miss_Friends_Birthdays	.548
Parents_Fuss	.693
Feel_Different_to_Friends	.800
Get_Upset_Not_Able_to_E at_Certain_Foods	.680

Extraction Method: Principal Component Analysis.

a. 1 components extracted.

```
ONEWAY Median_QoL BY Number_of_Allergens
  /STATISTICS HOMOGENEITY
  /MISSING ANALYSIS.
```

Oneway

Test of Homogeneity of Variances

		Levene Statistic	df1	df2	Sig.
Median_QoL	Based on Mean	.677	2	61	.512
	Based on Median	.306	2	61	.738
	Based on Median and with adjusted df	.306	2	49.387	.738
	Based on trimmed mean	.622	2	61	.540

ANOVA

Median_QoL

	Sum of Squares	df	Mean Square	F	Sig.
Between Groups	3.370	2	1.685	3.058	.054
Within Groups	33.614	61	.551		
Total	36.984	63			

```
T-TEST GROUPS=Number_of_Allergen$1 2)
/MISSING=ANALYSIS
/VARIABLES=Median_QoL
/CRITERIA=CI(.95).
```

T-Test

Group Statistics

	Number_of_Allergens	N	Mean	Std. Deviation	Std. Error Mean
Median_QoL	1 Allergen	15	2.33	.816	.211
	2 - 3 Allergens	23	1.74	.689	.144

Independent Samples Test

		Levene's Test for Equality of Variances		t-test for Equality of Means	
		F	Sig.	t	df
Median_QoL	Equal variances assumed	1.282	.265	2.416	36
	Equal variances not assumed			2.329	26.390

Independent Samples Test

		t-test for Equality of Means		
		Sig. (2-tailed)	Mean Difference	Std. Error Difference
Median_QoL	Equal variances assumed	.021	.594	.246
	Equal variances not assumed	.028	.594	.255

Independent Samples Test

		t-test for Equality of Means	
		95% Confidence Interval of the Difference	
		Lower	Upper
Median_QoL	Equal variances assumed	.095	1.093
	Equal variances not assumed	.070	1.118

```
T-TEST GROUPS=Number_of_Allergen$1 3)
/MISSING=ANALYSIS
/VARIABLES=Median_QoL
/CRITERIA=CI(.95).
```

T-Test

Group Statistics

		N	Mean	Std. Deviation	Std. Error Mean
Median_QoL	1 Allergen	15	2.33	.816	.211
	4 Allergens or More	26	2.08	.744	.146

Independent Samples Test

		Levene's Test for Equality of Variances		t-test for Equality of Means	
		F	Sig.	t	df
Median_QoL	Equal variances assumed	1.054	.311	1.026	39
	Equal variances not assumed			1.000	27.144

Independent Samples Test

		t-test for Equality of Means		
		Sig. (2-tailed)	Mean Difference	Std. Error Difference
Median_QoL	Equal variances assumed	.311	.256	.250
	Equal variances not assumed	.326	.256	.256

Independent Samples Test

		t-test for Equality of Means	
		95% Confidence Interval of the Difference	
		Lower	Upper
Median_QoL	Equal variances assumed	-.249	.762
	Equal variances not assumed	-.270	.782

```
T-TEST GROUPS=Number_of_Allergen$2)
/MISSING=ANALYSIS
/VARIABLES=Median_QoL
/CRITERIA=CI(.95).
```

T-Test

Group Statistics

	Number_of_Allergens	N	Mean	Std. Deviation	Std. Error Mean
Median_QoL	>= 2	49	1.92	.731	.104
	< 2	15	2.33	.816	.211

Independent Samples Test

		Levene's Test for Equality of Variances		t-test for Equality of Means	
		F	Sig.	t	df
Median_QoL	Equal variances assumed	1.322	.255	-1.871	62
	Equal variances not assumed			-1.764	21.346

Independent Samples Test

		t-test for Equality of Means		
		Sig. (2-tailed)	Mean Difference	Std. Error Difference
Median_QoL	Equal variances assumed	.066	-.415	.222
	Equal variances not assumed	.092	-.415	.235

Independent Samples Test

		t-test for Equality of Means	
		95% Confidence Interval of the Difference	
		Lower	Upper
Median_QoL	Equal variances assumed	-.858	.028
	Equal variances not assumed	-.904	.074

FREQUENCIES VARIABLES=KSA_Check_Adult KSA_Swap_Lunch KSA_Tell_Adult_if_Sick_After_Eating
 /ORDER=ANALYSIS.

Frequencies

		KSA_Check_Adult	KSA_Swap_Lunch	KSA_Tell_Adult_if_Sick_After_Eating
N	Valid	65	59	64
	Missing	0	6	1

Frequency Table

KSA Check Adult					
		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	65	100.0	100.0	100.0

KSA Swap Lunch					
		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	No	59	90.8	100.0	100.0
Missing	99	6	9.2		
Total		65	100.0		

KSA Tell Adult if Sick After Eating					
		Frequency	Percent	Valid Percent	Cumulative Percent
Valid	Yes	63	96.9	98.4	98.4
	No	1	1.5	1.6	100.0
	Total	64	98.5	100.0	
Missing	99	1	1.5		
Total		65	100.0		

Additional Statistics - ANOVA, linear model

```

GLM Miss_Friends_BirthdaysParents_Fuss Feel_Different_to_Friends
  Get_Upset_Not_Able_to_Eat_Certain_Foods BY Number_of_Allergens
/METHOD=SSTYPE(3)
/INTERCEPT=INCLUDE
/POSTHOC=Number_of_Allergen$TUKEY LSD BONFERRONI GT2 GH)
/EMMEANS=TABLES(Number_of_Allergen$ COMPARE ADJ(LSD)
/PRINT=DESCRIPTIVE ETASQ
/CRITERIA=ALPHA(.05)
/DESIGN= Number_of_Allergens

```

General Linear Model

Between-Subjects Factors

		Value Label	N
Number_of_Allergens	1	1 Allergen	15
	2	2 - 3 Allergens	22
	3	4 Allergens or More	27

Descriptive Statistics

	Number_of_Allergens	Mean	Std. Deviation	N
Miss_Friends_Birthdays	1 Allergen	2.53	.640	15
	2 - 3 Allergens	2.64	.492	22
	4 Allergens or More	2.59	.636	27
	Total	2.59	.583	64
Parents_Fuss	1 Allergen	1.87	.915	15
	2 - 3 Allergens	2.23	.685	22
	4 Allergens or More	2.04	.808	27
	Total	2.06	.794	64
Feel_Different_to_Friends	1 Allergen	1.73	.884	15
	2 - 3 Allergens	2.09	.811	22
	4 Allergens or More	1.96	.854	27
	Total	1.95	.844	64
Get_Upset_Not_Able_to_Eat_Certain_Foods	1 Allergen	1.73	.799	15
	2 - 3 Allergens	2.27	.631	22
	4 Allergens or More	1.93	.781	27
	Total	2.00	.756	64

Multivariate Tests^a

Effect		Value	F	Hypothesis df	Error df
Intercept	Pillai's Trace	.961	359.312 ^b	4.000	58.000
	Wilks' Lambda	.039	359.312 ^b	4.000	58.000
	Hotelling's Trace	24.780	359.312 ^b	4.000	58.000
	Roy's Largest Root	24.780	359.312 ^b	4.000	58.000
Number_of_Allergens	Pillai's Trace	.094	.724	8.000	118.000
	Wilks' Lambda	.907	.728 ^b	8.000	116.000
	Hotelling's Trace	.102	.730	8.000	114.000
	Roy's Largest Root	.099	1.458 ^c	4.000	59.000

Multivariate Tests^a

Effect		Sig.	Partial Eta Squared
Intercept	Pillai's Trace	.000	.961
	Wilks' Lambda	.000	.961
	Hotelling's Trace	.000	.961
	Roy's Largest Root	.000	.961
Number_of_Allergens	Pillai's Trace	.670	.047
	Wilks' Lambda	.667	.048
	Hotelling's Trace	.664	.049
	Roy's Largest Root	.227	.090

a. Design: Intercept + Number_of_Allergens

b. Exact statistic

c. The statistic is an upper bound on F that yields a lower bound on the significance level.

Estimated Marginal Means

Number_of_Allergens

Estimates

Dependent Variable	Number_of_Allergens	Mean	Std. Error	95% Confidence Interval	
				Lower Bound	Upper Bound
Miss_Friends_Birthdays	1 Allergen	2.533	.153	2.228	2.839
	2 - 3 Allergens	2.636	.126	2.384	2.889
	4 Allergens or More	2.593	.114	2.365	2.820
Parents_Fuss	1 Allergen	1.867	.205	1.456	2.277
	2 - 3 Allergens	2.227	.170	1.888	2.566
	4 Allergens or More	2.037	.153	1.731	2.343
Feel_Different_to_Friends	1 Allergen	1.733	.219	1.296	2.170
	2 - 3 Allergens	2.091	.180	1.730	2.452
	4 Allergens or More	1.963	.163	1.637	2.289
Get_Upset_Not_Able_to_Eat_Certain_Foods	1 Allergen	1.733	.190	1.353	2.114
	2 - 3 Allergens	2.273	.157	1.958	2.587
	4 Allergens or More	1.926	.142	1.642	2.210

Pairwise Comparisons

Dependent Variable	(I) Number_of_Allergens	(J) Number_of_Allergens	Mean
			Difference (I-J)
Miss_Friends_Birthdays	1 Allergen	2 - 3 Allergens	-.103
		4 Allergens or More	-.059
	2 - 3 Allergens	1 Allergen	.103
		4 Allergens or More	.044
	4 Allergens or More	1 Allergen	.059
Parents_Fuss	1 Allergen	2 - 3 Allergens	-.361
		4 Allergens or More	-.170
	2 - 3 Allergens	1 Allergen	.361
		4 Allergens or More	.190
	4 Allergens or More	1 Allergen	.170
Feel_Different_to_Friends	1 Allergen	2 - 3 Allergens	-.358
		4 Allergens or More	-.230
	2 - 3 Allergens	1 Allergen	.358
		4 Allergens or More	.128

Pairwise Comparisons

Dependent Variable	(I) Number_of_Allergens	(J) Number_of_Allergens	Std. Error	Sig. ^b
Miss_Friends_Birthdays	1 Allergen	2 - 3 Allergens	.198	.605
		4 Allergens or More	.190	.757
	2 - 3 Allergens	1 Allergen	.198	.605
		4 Allergens or More	.170	.798
	4 Allergens or More	1 Allergen	.190	.757
		2 - 3 Allergens	.170	.798
Parents_Fuss	1 Allergen	2 - 3 Allergens	.266	.181
		4 Allergens or More	.256	.508
	2 - 3 Allergens	1 Allergen	.266	.181
		4 Allergens or More	.228	.408
	4 Allergens or More	1 Allergen	.256	.508
		2 - 3 Allergens	.228	.408
Feel_Different_to_Friends	1 Allergen	2 - 3 Allergens	.283	.212
		4 Allergens or More	.273	.403
	2 - 3 Allergens	1 Allergen	.283	.212
		4 Allergens or More	.243	.601

Pairwise Comparisons

Dependent Variable	(I) Number_of_Allergens	(J) Number_of_Allergens	95% Confidence b...
			Lower Bound
Miss_Friends_Birthdays	1 Allergen	2 - 3 Allergens	-.499
		4 Allergens or More	-.440
	2 - 3 Allergens	1 Allergen	-.293
		4 Allergens or More	-.296
	4 Allergens or More	1 Allergen	-.322
		2 - 3 Allergens	-.383
Parents_Fuss	1 Allergen	2 - 3 Allergens	-.893
		4 Allergens or More	-.682
	2 - 3 Allergens	1 Allergen	-.172
		4 Allergens or More	-.266
	4 Allergens or More	1 Allergen	-.342
		2 - 3 Allergens	-.647
Feel_Different_to_Friends	1 Allergen	2 - 3 Allergens	-.924
		4 Allergens or More	-.775
	2 - 3 Allergens	1 Allergen	-.209
		4 Allergens or More	-.358

Pairwise Comparisons

Dependent Variable	(I) Number_of_Allergens	(J) Number_of_Allergens	95% Confidence Interval for ...
			Upper Bound
Miss_Friends_Birthdays	1 Allergen	2 - 3 Allergens	.293
		4 Allergens or More	.322
	2 - 3 Allergens	1 Allergen	.499
		4 Allergens or More	.383
	4 Allergens or More	1 Allergen	.440
		2 - 3 Allergens	.296
Parents_Fuss	1 Allergen	2 - 3 Allergens	.172
		4 Allergens or More	.342
	2 - 3 Allergens	1 Allergen	.893
		4 Allergens or More	.647
	4 Allergens or More	1 Allergen	.682
		2 - 3 Allergens	.266
Feel_Different_to_Friends	1 Allergen	2 - 3 Allergens	.209
		4 Allergens or More	.315
	2 - 3 Allergens	1 Allergen	.924
		4 Allergens or More	.614

Pairwise Comparisons

Dependent Variable	(I) Number_of_Allergens	(J) Number_of_Allergens	Mean Difference (I-J)
Get_Upset_Not_Able_to_Eat_Certain_Foods	4 Allergens or More	1 Allergen	.230
		2 - 3 Allergens	-.128
	1 Allergen	2 - 3 Allergens	-.539*
		4 Allergens or More	-.193
	2 - 3 Allergens	1 Allergen	.539*
		4 Allergens or More	.347
4 Allergens or More	1 Allergen	.193	
	2 - 3 Allergens	-.347	

Pairwise Comparisons

Dependent Variable	(I) Number_of_Allergens	(J) Number_of_Allergens	Std. Error	Sig. ^b
Get_Upset_Not_Able_to_E at_Certain_Foods	4 Allergens or More	1 Allergen	.273	.403
		2 - 3 Allergens	.243	.601
	1 Allergen	2 - 3 Allergens	.247	.033
		4 Allergens or More	.237	.420
	2 - 3 Allergens	1 Allergen	.247	.033
		4 Allergens or More	.212	.107
	4 Allergens or More	1 Allergen	.237	.420
		2 - 3 Allergens	.212	.107

Pairwise Comparisons

Dependent Variable	(I) Number_of_Allergens	(J) Number_of_Allergens	95% Confidence Int...
			Lower Bound
Get_Upset_Not_Able_to_E at_Certain_Foods	4 Allergens or More	1 Allergen	-.315
		2 - 3 Allergens	-.614
	1 Allergen	2 - 3 Allergens	-1.033
		4 Allergens or More	-.667
	2 - 3 Allergens	1 Allergen	.046
		4 Allergens or More	-.077
	4 Allergens or More	1 Allergen	-.282
		2 - 3 Allergens	-.770

Pairwise Comparisons

Dependent Variable	(I) Number_of_Allergens	(J) Number_of_Allergens	95% Confidence Interval for ...
			Upper Bound
Get_Upset_Not_Able_to_E at_Certain_Foods	4 Allergens or More	1 Allergen	.775
		2 - 3 Allergens	.358
	1 Allergen	2 - 3 Allergens	-.046
		4 Allergens or More	.282
	2 - 3 Allergens	1 Allergen	1.033
		4 Allergens or More	.770
	4 Allergens or More	1 Allergen	.667
		2 - 3 Allergens	.077

Based on estimated marginal means

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

b. Adjustment for multiple comparisons: Least Significant Difference (equivalent to no adjustments).

Multivariate Tests

	Value	F	Hypothesis df	Error df	Sig.	Partial Eta Squared
Pillai's trace	.094	.724	8.000	118.000	.670	.047
Wilks' lambda	.907	.728 ^a	8.000	116.000	.667	.048
Hotelling's trace	.102	.730	8.000	114.000	.664	.049
Roy's largest root	.099	1.458 ^b	4.000	59.000	.227	.090

Each F tests the multivariate effect of Number_of_Allergens. These tests are based on the linearly independent pairwise comparisons among the estimated marginal means.

a. Exact statistic

b. The statistic is an upper bound on F that yields a lower bound on the significance level.

Univariate Tests

Dependent Variable		Sum of Squares	df	Mean Square	F	Sig.
Miss_Friends_Birthdays	Contrast	.095	2	.047	.135	.874
	Error	21.343	61	.350		
Parents_Fuss	Contrast	1.190	2	.595	.941	.396
	Error	38.560	61	.632		
Feel_Different_to_Friends	Contrast	1.145	2	.572	.799	.455
	Error	43.714	61	.717		
Get_Upset_Not_Able_to_E at_Certain_Foods	Contrast	2.851	2	1.426	2.623	.081
	Error	33.149	61	.543		

Univariate Tests

Dependent Variable		Partial Eta Squared
Miss_Friends_Birthdays	Contrast	.004
	Error	
Parents_Fuss	Contrast	.030
	Error	
Feel_Different_to_Friends	Contrast	.026
	Error	
Get_Upset_Not_Able_to_E at_Certain_Foods	Contrast	.079
	Error	

The F tests the effect of Number_of_Allergens. This test is based on the linearly independent pairwise comparisons among the estimated marginal means.

Post Hoc Tests

Number_of_Allergens

Homogeneous Subsets

Miss_Friends_Birthdays

	Number_of_Allergens	N	Subset 1
Tukey HSD ^{a,b,c}	1 Allergen	15	2.53
	4 Allergens or More	27	2.59
	2 - 3 Allergens	22	2.64
	Sig.		.846
Hochberg ^{a,b,c}	1 Allergen	15	2.53
	4 Allergens or More	27	2.59
	2 - 3 Allergens	22	2.64
	Sig.		.926

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = .350.

- a. Uses Harmonic Mean Sample Size = 20.113.
- b. The group sizes are unequal. The harmonic mean of the group sizes is used. Type I error levels are not guaranteed.
- c. Alpha = .05.

Parents_Fuss

	Number_of_Allergens	N	Subset 1
Tukey HSD ^{a,b,c}	1 Allergen	15	1.87
	4 Allergens or More	27	2.04
	2 - 3 Allergens	22	2.23
	Sig.		.328
Hochberg ^{a,b,c}	1 Allergen	15	1.87
	4 Allergens or More	27	2.04
	2 - 3 Allergens	22	2.23
	Sig.		.394

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = .632.

- a. Uses Harmonic Mean Sample Size = 20.113.
- b. The group sizes are unequal. The harmonic mean of the group sizes is used. Type I error levels are not guaranteed.
- c. Alpha = .05.

Feel_Different_to_Friends

	Number_of_Allergens	N	Subset 1
Tukey HSD ^{a,b,c}	1 Allergen	15	1.73
	4 Allergens or More	27	1.96
	2 - 3 Allergens	22	2.09
	Sig.		.379
Hochberg ^{a,b,c}	1 Allergen	15	1.73
	4 Allergens or More	27	1.96
	2 - 3 Allergens	22	2.09
	Sig.		.456

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = .717.

- a. Uses Harmonic Mean Sample Size = 20.113.
- b. The group sizes are unequal. The harmonic mean of the group sizes is used. Type I error levels are not guaranteed.
- c. Alpha = .05.

Get_Upset_Not_Able_to_Eat_Certain_Foods

	Number_of_Allergens	N	Subset 1
Tukey HSD ^{a,b,c}	1 Allergen	15	1.73
	4 Allergens or More	27	1.93
	2 - 3 Allergens	22	2.27
	Sig.		.060
Hochberg ^{a,b,c}	1 Allergen	15	1.73
	4 Allergens or More	27	1.93
	2 - 3 Allergens	22	2.27
	Sig.		.069

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = .543.

a. Uses Harmonic Mean Sample Size = 20.113.

b. The group sizes are unequal. The harmonic mean of the group sizes is used. Type I error levels are not guaranteed.

c. Alpha = .05.

Regression

[DataSet3] E:\MASTER_Under 8years.sav

Variables Entered/Removed^a

Model	Variables Entered	Variables Removed	Method
1	Gender ^b	.	Enter

a. Dependent Variable: Miss_Friends_Birthdays

b. All requested variables entered.

Model Summary^b

Model	R	R Square	Adjusted R Square	Std. Error of the Estimate
1	.157 ^a	.025	.009	.581

a. Predictors: (Constant), Gender

b. Dependent Variable: Miss_Friends_Birthdays

ANOVA^a

Model		Sum of Squares	df	Mean Square	F	Sig.
1	Regression	.521	1	.521	1.545	.219 ^b
	Residual	20.559	61	.337		
	Total	21.079	62			

a. Dependent Variable: Miss_Friends_Birthdays

b. Predictors: (Constant), Gender

Coefficients^a

Model		Unstandardized Coefficients		Standardized Coefficients	t	Sig.
		B	Std. Error	Beta		
1	(Constant)	2.346	.220		10.685	.000
	Gender	.191	.153	.157	1.243	.219

Coefficients^a

Model		95.0% Confidence Interval for B		Collinearity Statistics	
		Lower Bound	Upper Bound	Tolerance	VIF
1	(Constant)	1.907	2.785		
	Gender	-.116	.497	1.000	1.000

a. Dependent Variable: Miss_Friends_Birthdays

Collinearity Diagnostics^a

Model	Dimension	Eigenvalue	Condition Index	Variance Proportions	
				(Constant)	Gender
1	1	1.943	1.000	.03	.03
	2	.057	5.832	.97	.97

a. Dependent Variable: Miss_Friends_Birthdays

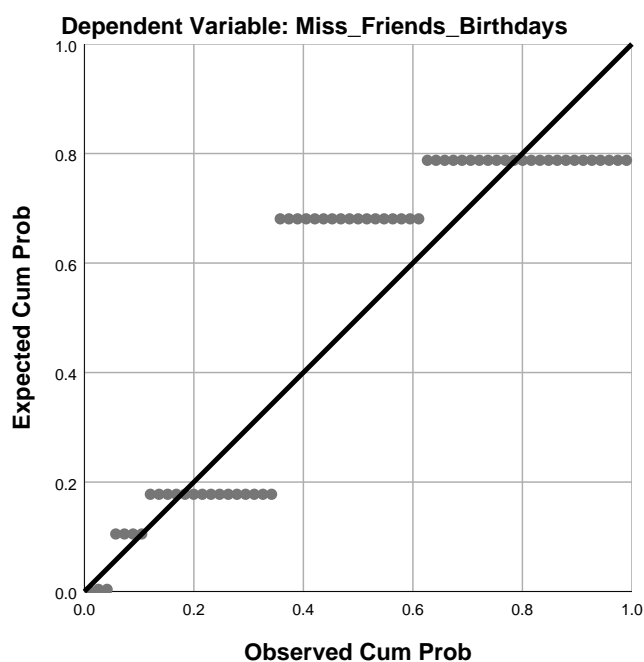
Residuals Statistics^a

	Minimum	Maximum	Mean	Std. Deviation	N
Predicted Value	2.54	2.73	2.60	.092	63
Residual	-1.727	.463	.000	.576	63
Std. Predicted Value	-.727	1.354	.000	1.000	63
Std. Residual	-2.975	.798	.000	.992	63

a. Dependent Variable: Miss_Friends_Birthdays

Charts

Normal P-P Plot of Regression Standardized Residual



REGRESSION

/MISSING LISTWISE

/STATISTICS COEFF OUTS CI(95) R ANOVA COLLIN TOL

/CRITERIA=PIN(.05) POUT(.10)

/NOORIGIN

/DEPENDENT Parents_Fuss

/METHOD=ENTER Gender

/SCATTERPLOT=(*ZRESID ,*ZPRED)

/RESIDUALS NORMPROB(ZRESID) .

Regression

Variables Entered/Removed^a

Model	Variables Entered	Variables Removed	Method
1	Gender ^b	.	Enter

a. Dependent Variable: Parents_Fuss

b. All requested variables entered.

Model Summary^b

Model	R	R Square	Adjusted R Square	Std. Error of the Estimate
1	.096 ^a	.009	-.007	.792

a. Predictors: (Constant), Gender

b. Dependent Variable: Parents_Fuss

ANOVA^a

Model		Sum of Squares	df	Mean Square	F	Sig.
1	Regression	.355	1	.355	.566	.455 ^b
	Residual	38.248	61	.627		
	Total	38.603	62			

a. Dependent Variable: Parents_Fuss

b. Predictors: (Constant), Gender

Coefficients^a

Model		Unstandardized Coefficients		Standardized Coefficients	t	Sig.
		B	Std. Error	Beta		
1	(Constant)	1.867	.299		6.235	.000
	Gender	.157	.209	.096	.752	.455

Coefficients^a

Model		95.0% Confidence Interval for B		Collinearity Statistics	
		Lower Bound	Upper Bound	Tolerance	VIF
1	(Constant)	1.268	2.466		
	Gender	-.261	.576	1.000	1.000

a. Dependent Variable: Parents_Fuss

Collinearity Diagnostics^a

Model	Dimension	Eigenvalue	Condition Index	Variance Proportions	
				(Constant)	Gender
1	1	1.943	1.000	.03	.03
	2	.057	5.832	.97	.97

a. Dependent Variable: Parents_Fuss

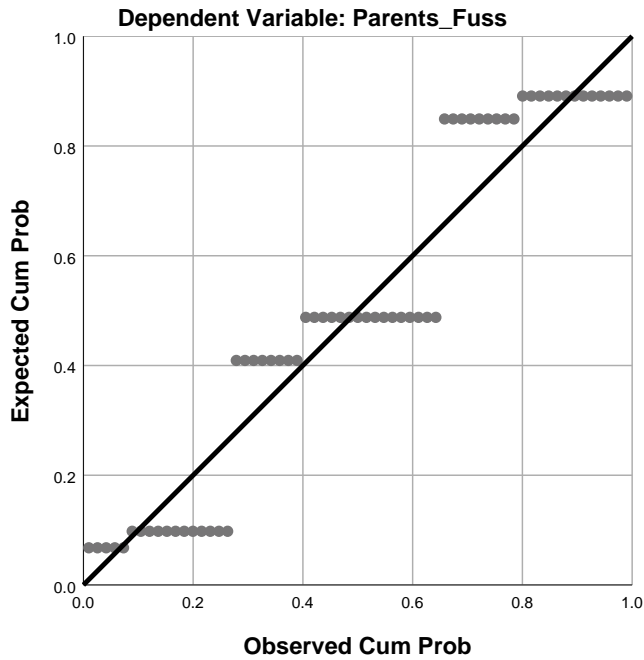
Residuals Statistics^a

	Minimum	Maximum	Mean	Std. Deviation	N
Predicted Value	2.02	2.18	2.08	.076	63
Residual	-1.182	.976	.000	.785	63
Std. Predicted Value	-.727	1.354	.000	1.000	63
Std. Residual	-1.492	1.232	.000	.992	63

a. Dependent Variable: Parents_Fuss

Charts

Normal P-P Plot of Regression Standardized Residual



```

REGRESSION
  /MISSING LISTWISE
  /STATISTICS COEFF OUTS CI(95) R ANOVA COLLIN TOL
  /CRITERIA=PIN(.05) POUT(.10)
  /NOORIGIN
  /DEPENDENT Feel_Different_to_Friends
  /METHOD=ENTER Gender
  /SCATTERPLOT=(*ZRESID ,*ZPRED)
  /RESIDUALS NORMPROB(ZRESID) .
  
```

Regression

Variables Entered/Removed^a

Model	Variables Entered	Variables Removed	Method
1	Gender ^b	.	Enter

a. Dependent Variable: Feel_Different_to_Friends

b. All requested variables entered.

Model Summary^b

Model	R	R Square	Adjusted R Square	Std. Error of the Estimate
1	.012 ^a	.000	-.016	.849

a. Predictors: (Constant), Gender

b. Dependent Variable: Feel_Different_to_Friends

ANOVA^a

Model		Sum of Squares	df	Mean Square	F	Sig.
1	Regression	.006	1	.006	.009	.925 ^b
	Residual	43.930	61	.720		
	Total	43.937	62			

a. Dependent Variable: Feel_Different_to_Friends

b. Predictors: (Constant), Gender

Coefficients^a

Model		Unstandardized Coefficients		Standardized Coefficients	t	Sig.
		B	Std. Error	Beta		
1	(Constant)	1.997	.321		6.222	.000
	Gender	-.021	.224	-.012	-.094	.925

Coefficients^a

Model		95.0% Confidence Interval for B		Collinearity Statistics	
		Lower Bound	Upper Bound	Tolerance	VIF
1	(Constant)	1.355	2.638		
	Gender	-.470	.427	1.000	1.000

a. Dependent Variable: Feel_Different_to_Friends

Collinearity Diagnostics^a

Model	Dimension	Eigenvalue	Condition Index	Variance Proportions	
				(Constant)	Gender
1	1	1.943	1.000	.03	.03
	2	.057	5.832	.97	.97

a. Dependent Variable: Feel_Different_to_Friends

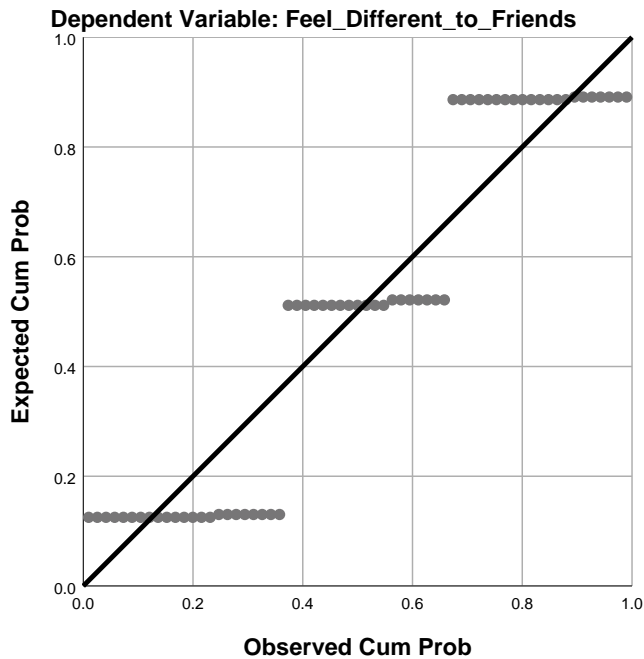
Residuals Statistics^a

	Minimum	Maximum	Mean	Std. Deviation	N
Predicted Value	1.95	1.98	1.97	.010	63
Residual	-.976	1.045	.000	.842	63
Std. Predicted Value	-1.354	.727	.000	1.000	63
Std. Residual	-1.150	1.232	.000	.992	63

a. Dependent Variable: Feel_Different_to_Friends

Charts

Normal P-P Plot of Regression Standardized Residual



```

/MISSING LISTWISE
/STATISTICS COEFF OUTS CI(95) R ANOVA COLLIN TOL
/CRITERIA=PIN(.05) POUT(.10)
/NOORIGIN
/DEPENDENT Get_Upset_Not_Able_to_Eat_Certain_Foods
/METHOD=ENTER Gender
/SCATTERPLOT=(*ZRESID ,*ZPRED)
/RESIDUALS NORMPROB(ZRESID) .

```

Regression

Variables Entered/Removed^a

Model	Variables Entered	Variables Removed	Method
1	Gender ^b	.	Enter

a. Dependent Variable: Get_Upset_Not_Able_to_Eat_Certain_Foods

b. All requested variables entered.

Model Summary^b

Model	R	R Square	Adjusted R Square	Std. Error of the Estimate
1	.044 ^a	.002	-.014	.767

a. Predictors: (Constant), Gender

b. Dependent Variable: Get_Upset_Not_Able_to_Eat_Certain_Foods

ANOVA^a

Model		Sum of Squares	df	Mean Square	F	Sig.
1	Regression	.070	1	.070	.119	.732 ^b
	Residual	35.930	61	.589		
	Total	36.000	62			

a. Dependent Variable: Get_Upset_Not_Able_to_Eat_Certain_Foods

b. Predictors: (Constant), Gender

Coefficients^a

Model		Unstandardized Coefficients		Standardized Coefficients	t	Sig.
		B	Std. Error	Beta		
1	(Constant)	1.906	.290		6.566	.000
	Gender	.070	.203	.044	.344	.732

Coefficients^a

Model		95.0% Confidence Interval for B		Collinearity Statistics	
		Lower Bound	Upper Bound	Tolerance	VIF
1	(Constant)	1.325	2.486		
	Gender	-.336	.475	1.000	1.000

a. Dependent Variable: Get_Upset_Not_Able_to_Eat_Certain_Foods

Collinearity Diagnostics^a

Model	Dimension	Eigenvalue	Condition Index	Variance Proportions	
				(Constant)	Gender
1	1	1.943	1.000	.03	.03
	2	.057	5.832	.97	.97

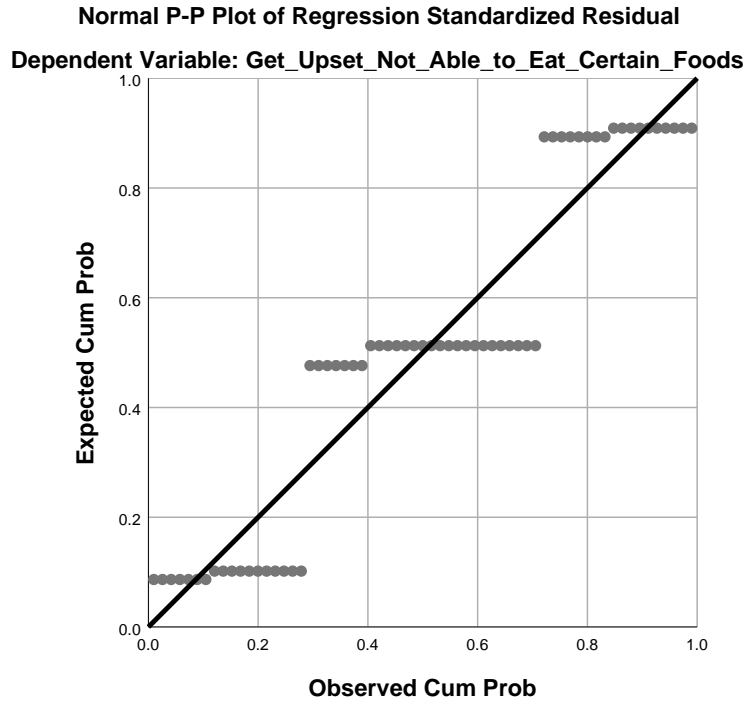
a. Dependent Variable: Get_Upset_Not_Able_to_Eat_Certain_Foods

Residuals Statistics^a

	Minimum	Maximum	Mean	Std. Deviation	N
Predicted Value	1.98	2.05	2.00	.034	63
Residual	-1.045	1.024	.000	.761	63
Std. Predicted Value	-.727	1.354	.000	1.000	63
Std. Residual	-1.362	1.335	.000	.992	63

a. Dependent Variable: Get_Upset_Not_Able_to_Eat_Certain_Foods

Charts



```

REGRESSION
  /MISSING LISTWISE
  /STATISTICS COEFF OUTS CI(95) R ANOVA COLLIN TOL
  /CRITERIA=PIN(.05) POUT(.10)
  /NOORIGIN
  /DEPENDENT Miss_Friends_Birthdays
  /METHOD=ENTER Number_of_Allergens
  /SCATTERPLOT=(*ZRESID ,*ZPRED)
  /RESIDUALS NORMPROB(ZRESID) .

```

Regression

Variables Entered/Removed^a

Model	Variables Entered	Variables Removed	Method
1	Number_of_Allergens ^b	.	Enter

a. Dependent Variable: Miss_Friends_Birthdays

b. All requested variables entered.

Model Summary^b

Model	R	R Square	Adjusted R Square	Std. Error of the Estimate
1	.030 ^a	.001	-.015	.588

a. Predictors: (Constant), Number_of_Allergens

b. Dependent Variable: Miss_Friends_Birthdays

ANOVA^a

Model		Sum of Squares	df	Mean Square	F	Sig.
1	Regression	.019	1	.019	.056	.814 ^b
	Residual	21.418	62	.345		
	Total	21.438	63			

a. Dependent Variable: Miss_Friends_Birthdays

b. Predictors: (Constant), Number_of_Allergens

Coefficients^a

Model		Unstandardized Coefficients		Standardized Coefficients	t	Sig.
		B	Std. Error	Beta		
1	(Constant)	2.546	.217		11.744	.000
	Number_of_Allergens	.022	.093	.030	.236	.814

Coefficients^a

Model		95.0% Confidence Interval for B		Collinearity Statistics	
		Lower Bound	Upper Bound	Tolerance	VIF
1	(Constant)	2.112	2.979		
	Number_of_Allergens	-.164	.208	1.000	1.000

a. Dependent Variable: Miss_Friends_Birthdays

Collinearity Diagnostics^a

Model	Dimension	Eigenvalue	Condition Index	Variance Proportions	
				(Constant)	Number_of_All ergens
1	1	1.941	1.000	.03	.03
	2	.059	5.726	.97	.97

a. Dependent Variable: Miss_Friends_Birthdays

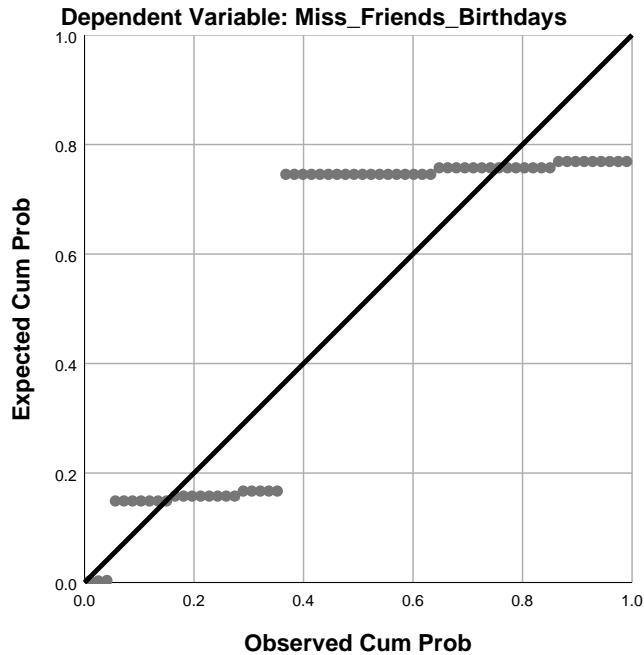
Residuals Statistics^a

	Minimum	Maximum	Mean	Std. Deviation	N
Predicted Value	2.57	2.61	2.59	.017	64
Residual	-1.612	.432	.000	.583	64
Std. Predicted Value	-1.495	1.023	.000	1.000	64
Std. Residual	-2.742	.736	.000	.992	64

a. Dependent Variable: Miss_Friends_Birthdays

Charts

Normal P-P Plot of Regression Standardized Residual



```

REGRESSION
/MISSING LISTWISE
/STATISTICS COEFF OUTS CI(95) R ANOVA COLLIN TOL
/CRITERIA=PIN(.05) POUT(.10)
/NOORIGIN
/DEPENDENT Parents_Fuss
/METHOD=ENTER Number_of_Allergens
/SCATTERPLOT=(*ZRESID ,*ZPRED)
/RESIDUALS NORMPROB(ZRESID) .

```

Regression

Variables Entered/Removed^a

Model	Variables Entered	Variables Removed	Method
1	Number_of_Allergens ^b	.	Enter

a. Dependent Variable: Parents_Fuss

b. All requested variables entered.

Model Summary^b

Model	R	R Square	Adjusted R Square	Std. Error of the Estimate
1	.057 ^a	.003	-.013	.799

a. Predictors: (Constant), Number_of_Allergens

b. Dependent Variable: Parents_Fuss

ANOVA^a

Model		Sum of Squares	df	Mean Square	F	Sig.
1	Regression	.127	1	.127	.199	.657 ^b
	Residual	39.623	62	.639		
	Total	39.750	63			

a. Dependent Variable: Parents_Fuss

b. Predictors: (Constant), Number_of_Allergens

Coefficients^a

Model		Unstandardized Coefficients		Standardized Coefficients	t	Sig.
		B	Std. Error	Beta		
1	(Constant)	1.939	.295		6.576	.000
	Number_of_Allergens	.057	.127	.057	.446	.657

Coefficients^a

Model		95.0% Confidence Interval for B		Collinearity Statistics	
		Lower Bound	Upper Bound	Tolerance	VIF
1	(Constant)	1.349	2.528		
	Number_of_Allergens	-.197	.310	1.000	1.000

a. Dependent Variable: Parents_Fuss

Collinearity Diagnostics^a

Model	Dimension	Eigenvalue	Condition Index	Variance Proportions	
				(Constant)	Number_of_Allergens
1	1	1.941	1.000	.03	.03
	2	.059	5.726	.97	.97

a. Dependent Variable: Parents_Fuss

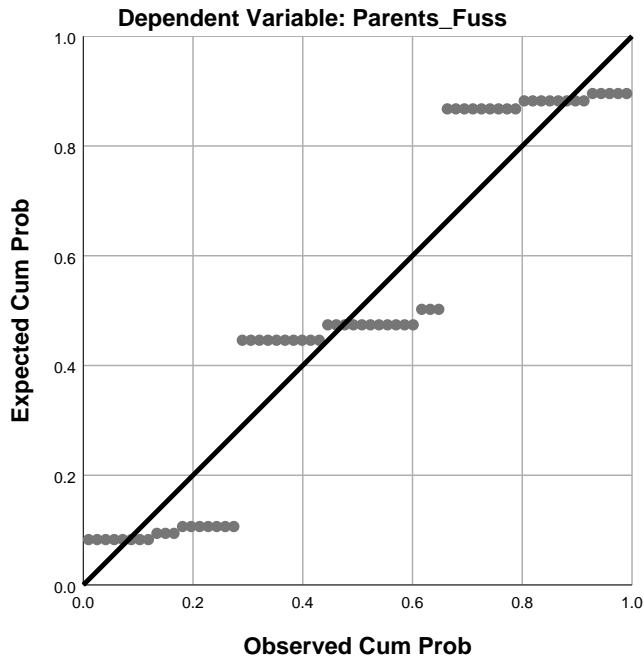
Residuals Statistics^a

	Minimum	Maximum	Mean	Std. Deviation	N
Predicted Value	2.00	2.11	2.06	.045	64
Residual	-1.108	1.005	.000	.793	64
Std. Predicted Value	-1.495	1.023	.000	1.000	64
Std. Residual	-1.387	1.257	.000	.992	64

a. Dependent Variable: Parents_Fuss

Charts

Normal P-P Plot of Regression Standardized Residual



REGRESSION

```

/MISSING LISTWISE
/STATISTICS COEFF OUTS CI(95) R ANOVA COLLIN TOL
/CRITERIA=PIN(.05) POUT(.10)
/NOORIGIN
/DEPENDENT Feel_Different_to_Friends
/METHOD=ENTER Number_of_Allergens
/SCATTERPLOT=(*ZRESID ,*ZPRED)
/RESIDUALS NORMPROB(ZRESID) .
    
```

Regression

Variables Entered/Removed^a

Model	Variables Entered	Variables Removed	Method
1	Number_of_Allergens ^b	.	Enter

a. Dependent Variable: Feel_Different_to_Friends

b. All requested variables entered.

Model Summary^b

Model	R	R Square	Adjusted R Square	Std. Error of the Estimate
1	.084 ^a	.007	-.009	.848

a. Predictors: (Constant), Number_of_Allergens

b. Dependent Variable: Feel_Different_to_Friends

ANOVA^a

Model		Sum of Squares	df	Mean Square	F	Sig.
1	Regression	.319	1	.319	.444	.507 ^b
	Residual	44.540	62	.718		
	Total	44.859	63			

a. Dependent Variable: Feel_Different_to_Friends

b. Predictors: (Constant), Number_of_Allergens

Coefficients^a

Model		Unstandardized Coefficients		Standardized Coefficients	t	Sig.
		B	Std. Error	Beta		
1	(Constant)	1.757	.313		5.621	.000
	Number_of_Allergens	.090	.134	.084	.667	.507

Coefficients^a

Model		95.0% Confidence Interval for B		Collinearity Statistics	
		Lower Bound	Upper Bound	Tolerance	VIF
1	(Constant)	1.132	2.382		
	Number_of_Allergens	-.179	.358	1.000	1.000

a. Dependent Variable: Feel_Different_to_Friends

Collinearity Diagnostics^a

Model	Dimension	Eigenvalue	Condition Index	Variance Proportions	
				(Constant)	Number_of_All ergens
1	1	1.941	1.000	.03	.03
	2	.059	5.726	.97	.97

a. Dependent Variable: Feel_Different_to_Friends

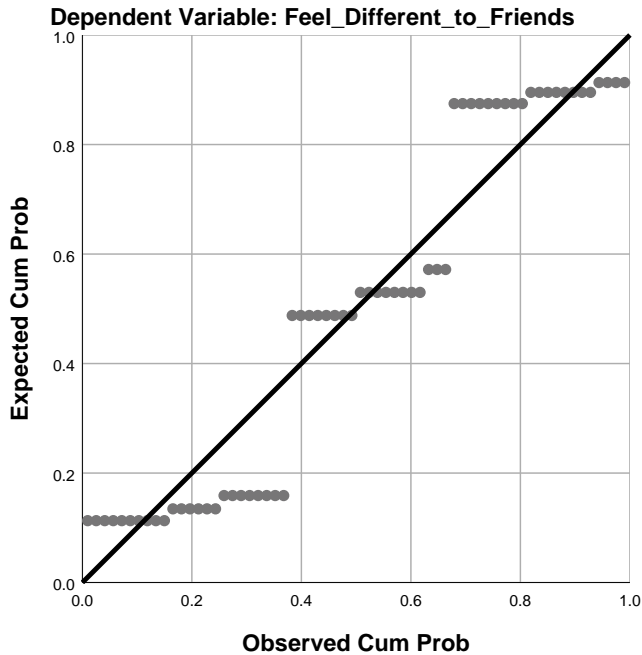
Residuals Statistics^a

	Minimum	Maximum	Mean	Std. Deviation	N
Predicted Value	1.85	2.03	1.95	.071	64
Residual	-1.026	1.153	.000	.841	64
Std. Predicted Value	-1.495	1.023	.000	1.000	64
Std. Residual	-1.210	1.361	.000	.992	64

a. Dependent Variable: Feel_Different_to_Friends

Charts

Normal P-P Plot of Regression Standardized Residual



```

REGRESSION
/MISSING LISTWISE
/STATISTICS COEFF OUTS CI(95) R ANOVA COLLIN TOL
/CRITERIA=PIN(.05) POUT(.10)
/NOORIGIN
/DEPENDENT Get_Upset_Not_Able_to_Eat_Certain_Foods
/METHOD=ENTER Number_of_Allergens
/SCATTERPLOT=(*ZRESID ,*ZPRED)
/RESIDUALS NORMPROB(ZRESID) .

```

Regression

Variables Entered/Removed^a

Model	Variables Entered	Variables Removed	Method
1	Number_of_Allergens ^b	.	Enter

a. Dependent Variable: Get_Upset_Not_Able_to_Eat_Certain_Foods

b. All requested variables entered.

Model Summary^b

Model	R	R Square	Adjusted R Square	Std. Error of the Estimate
1	.053 ^a	.003	-.013	.761

a. Predictors: (Constant), Number_of_Allergens

b. Dependent Variable: Get_Upset_Not_Able_to_Eat_Certain_Foods

ANOVA^a

Model		Sum of Squares	df	Mean Square	F	Sig.
1	Regression	.101	1	.101	.174	.678 ^b
	Residual	35.899	62	.579		
	Total	36.000	63			

a. Dependent Variable: Get_Upset_Not_Able_to_Eat_Certain_Foods

b. Predictors: (Constant), Number_of_Allergens

Coefficients^a

Model		Unstandardized Coefficients		Standardized Coefficients	t	Sig.
		B	Std. Error	Beta		
1	(Constant)	1.890	.281		6.735	.000
	Number_of_Allergens	.050	.121	.053	.417	.678

Coefficients^a

Model		95.0% Confidence Interval for B		Collinearity Statistics	
		Lower Bound	Upper Bound	Tolerance	VIF
1	(Constant)	1.329	2.451		
	Number_of_Allergens	-.191	.292	1.000	1.000

a. Dependent Variable: Get_Upset_Not_Able_to_Eat_Certain_Foods

Collinearity Diagnostics^a

Model	Dimension	Eigenvalue	Condition Index	Variance Proportions	
				(Constant)	Number_of_Allergens
1	1	1.941	1.000	.03	.03
	2	.059	5.726	.97	.97

a. Dependent Variable: Get_Upset_Not_Able_to_Eat_Certain_Foods

Residuals Statistics^a

	Minimum	Maximum	Mean	Std. Deviation	N
Predicted Value	1.94	2.04	2.00	.040	64
Residual	-1.041	1.060	.000	.755	64
Std. Predicted Value	-1.495	1.023	.000	1.000	64
Std. Residual	-1.368	1.393	.000	.992	64

a. Dependent Variable: Get_Upset_Not_Able_to_Eat_Certain_Foods