MULTIPLE LINEAR REGRESSION

EXAMPLE: BLOOD VISCOSITY AND PACKED CELL VOLUME

The following blood viscosity data studied earlier are a good example of where multiple regression could be used. Recall that the data blood viscosity in samples taken from 32 hospital patients. We wish to model viscosity (y) as a function three covariates

- Packed Cell Volume (PCV), *x*₁.
- Plasma Fibrinogen, x_2 .
- Plasma Protein, *x*₃.

Unit	Viscosity	PCV	Plasma Fib.	Plasma Pro.
	y	x_1	x_2	x_3
1	3.71	40.00	344	6.27
2	3.78	40.00	330	4.86
3	3.85	42.50	280	5.09
4	3.88	42.00	418	6.79
5	3.98	45.00	774	6.40
6	4.03	42.00	388	5.48
7	4.05	42.50	336	6.27
8	4.14	47.00	431	6.89
9	4.14	46.75	276	5.18
10	4.20	48.00	422	5.73
11	4.20	46.00	280	5.89
12	4.27	47.00	460	6.58
13	4.27	43.25	412	5.67
14	4.37	45.00	320	6.23
15	4.41	50.00	502	4.99
16	4.64	45.00	550	6.37
17	4.68	51.25	414	6.40
18	4.73	50.25	304	6.00
19	4.87	49.00	472	5.94
20	4.94	50.00	728	5.16
21	4.95	50.00	716	6.29
22	4.96	49.00	400	5.96
23	5.02	50.50	576	5.90
24	5.02	51.25	354	5.81
25	5.12	49.50	392	5.49
26	5.15	56.00	352	5.41
27	5.17	50.00	572	6.24
28	5.18	47.00	634	6.50
29	5.38	53.25	458	6.60
30	5.77	57.00	1070	4.82
31	5.90	54.00	488	5.70
32	5.90	54.00	488	5.70

We consider four analyses:

Model Summary^b

Model	R	R Square	Adjusted R Square	Std. Error of the Estimate
1	.885 ^a	.784	.761	.30370

a. Predictors: (Constant), Plasma Protein (g/100ml),
 Plasma Fibrinogen (mg/100ml), Packed Cell Volume (%)

ANOVAb

Model		Sum of Squares	df	Mean Square	F	Sig.
1	Regression	9.368	3	3.123	33.856	.000 ^a
	Residual	2.582	28	.092		
	Total	11.950	31			

- a. Predictors: (Constant), Plasma Protein (g/100ml), Plasma Fibrinogen (mg/100ml), Packed Cell Volume (%)
- b. Dependent Variable: Blood Viscosity (cP)

Multiple Regression: Parameter Estimates



a. Dependent Variable: Blood Viscosity (cP)

Only the packed cell volume coefficient is significantly different from zero (p < 0.001)

The other covariates do not seem to be significantly different from zero.

Regression on Packed Cell Volume only

Model Summary

Model	R	R Square	Adjusted R Square	Std. Error of the Estimate
1	.879 ^a	.772	.765	.30116

a. Predictors: (Constant), Packed Cell Volume (%)

\textbf{ANOVA}^{b}

Model		Sum of Squares	df	Mean Square	F	Sig.
1	Regression	9.230	1	9.230	101.764	.000 ^a
	Residual	2.721	30	.091		
	Total	11.950	31			

a. Predictors: (Constant), Packed Cell Volume (%)

b. Dependent Variable: Blood Viscosity (cP)

Coefficients^a

		Unstandardized Coefficients		Standardized Coefficients			95% Confidenc	e Interval for B
Model		В	Std. Error	Beta	t	Sig.	Lower Bound	Upper Bound
1	(Constant)	-1.223	.584		-2.094	.045	-2.416	030
	Packed Cell Volume (%)	.122	.012	.879	10.088	.000	.098	.147

a. Dependent Variable: Blood Viscosity (cP)

Regression on Plasma Protein only

Model Summary

Model	R	R Square	Adjusted R Square	Std. Error of the Estimate
1	.457 ^a	.209	.183	.56129

a. Predictors: (Constant), Plasma Fibrinogen (mg/100ml)

\textbf{ANOVA}^{b}

Model		Sum of Squares	df	Mean Square	F	Sig.
1	Regression	2.499	1	2.499	7.932	.009 ^a
	Residual	9.451	30	.315		
	Total	11.950	31			

a. Predictors: (Constant), Plasma Fibrinogen (mg/100ml)

b. Dependent Variable: Blood Viscosity (cP)

Coefficients^a

		Unstandardized Coefficients		Standardized Coefficients			95% Confidence	e Interval for B
Model		В	Std. Error	Beta	t	Sig.	Lower Bound	Upper Bound
1	(Constant)	3.871	.292		13.236	.000	3.274	4.468
	Plasma Fibrinogen (mg/100ml)	.002	.001	.457	2.816	.009	.000	.003

a. Dependent Variable: Blood Viscosity (cP)

Plasfib is a significant term in the model (p = 0.009)

Regression on Plasma Fibrinogen only

Model Summary

Model	R	R Square	Adjusted R Square	Std. Error of the Estimate
1	.101 ^a	.010	023	.62791

a. Predictors: (Constant), Plasma Protein (g/100ml)

\textbf{ANOVA}^{b}

Model		Sum of Squares	df	Mean Square	F	Sig.
1	Regression	.122	1	.122	.310	.582 ^a
	Residual	11.828	30	.394		
	Total	11.950	31			

a. Predictors: (Constant), Plasma Protein (g/100ml)

b. Dependent Variable: Blood Viscosity (cP)

Coefficients^a

Unstandardized Coefficients		lardized cients	Standardized Coefficients			95% Confidence	e Interval for B	
Model		В	Std. Error	Beta	t	Sig.	Lower Bound	Upper Bound
1	(Constant)	5.296	1.174		4.510	.000	2.898	7.694
	Plasma Protein (g/100ml)	110	.198	101	556	.582	515	.295

a. Dependent Variable: Blood Viscosity (cP)

Plaspro is not a significant term in the model (p =0.582)

Blood Viscosity Data Set

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Select the Dependent variable (**viscosity**) and the three independent variables 3 (pcv, plasfib and plaspro)



Click the Statistics button: on the Statistics dialog, select Estimates, Confidence⁴ Intervals and Model fit. Click Continue.



Click the Plots button



Select ***ZRESID** for the *Y* variable and ***ZPRED** for the *X* variable.

Then click Next.



Select ***ZRESID** for the *Y* variable and ***ZPRED** for the *X* variable.

Then click Produce all partial Plots. Then Continue.



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Select the quantities to store as new variables in the data set.

Click Continue.



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Click OK and the output is generated.

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6	6	4.03	42.00	388	5.48	3.90369	.12631	.41590	3.68995	4.11743	3.24590	4.56148								
7	7	4.05	42.50	336	6.27	3.97283	.07717	.25409	3.78695	4.15872	3.32356	4.62211								
8	8	4.14	47.00	431	6.89	4.56154	42154	-1.38804	4.33621	4.78688	3.89990	5.22319								
9	9	4.14	46.75	276	5.18	4.40157	26157	86130	4.17972	4.62343	3.74110	5.06204		_	-	_				
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14	14	4.37	45.00	320	6.23	4.25686	.11314	.37255	4.09611	4.41761	3.61433	4.89939								
15	15	4.41	50.00	502	4.99	4.86448	45448	-1.49650	4.65413	5.07484	4.20779	5.52118								
16	16	4.64	45.00	550	6.37	4.35491	.28509	.93873	4.17363	4.53619	3.70694	5.00288								
17	17	4.68	51.25	414	6.40	5.03159	35159	-1.15771	4.83663	5.22655	4.37966	5.68352								
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20	19	4.07	49.00	472	5.94	4.77364	.09636	. 07287	4.00930	4.00791	4.14113	5.40014								<u> </u>
20	20	4.95	50.00	716	6.29	5.00255	05255	17303	4.78804	5.21706	4.34451	5.66059								<u> </u>
22	22	4.96	49.00	400	5.96	4.74550	.21450	.70631	4.61602	4.87497	4.11007	5.38092								
23	23	5.02	50.50	576	5.90	4.98907	.03093	.10184	4.85220	5.12594	4.35210	5.62605								
24	24	5.02	51.25	354	5.81	4.98385	.03615	.11902	4.80178	5.16593	4.33566	5.63205								
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31	31	5.90	54.00	488	5.70	5.35457	.54543	1.79595	5.15881	5.55034	4.70240	6.00675								
32	32	5.90	54.00	488	5.70	5.35457	.54543	1.79595	5.15881	5.55034	4.70240	6.00675								
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5 plaspro	Numeric	10	2	Plasma Protein (g/100ml)	None	None	8	Right	Scale
6 PRE_1	Numeric	11	5	Unstandardized Predicted Value	None	None	13	Right	Scale
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- a. Predictors: (Constant), Plasma Protein (g/100ml),
 Plasma Fibrinogen (mg/100ml), Packed Cell Volume (%)
- b. Dependent Variable: Blood Viscosity (cP)

Results: ANOVA

ANOVA^b

Model		Sum of Squares	df	Mean Square	F	Sig.
1	Regression	9.368	3	3.123	33.856	.000 ^a
	Residual	2.582	28	.092		
	Total	11.950	31			

a. Predictors: (Constant), Plasma Protein (g/100ml), Plasma Fibrinogen (mg/100ml), Packed Cell Volume (%)

b. Dependent Variable: Blood Viscosity (cP)

The ANOVA for the multiple regression has a highly significant F value, with a p-value < 0.001. Here H0 : E[Y] = beta.0H1 : E[Y] = beta.0 + beta.1 x1 + beta.2 x2 + beta.3 x3This result implies that the multiple regression (Ha) fits significantly better than the model with no dependence on any of the predictors (H0).

Results: Parameter Estimates

Coefficients^a

		Unstar Coet	dardized ficients	Standardized Coefficients			95% Confidence	e Interval for B
Model		В	Std. Error	Beta	t	Sig.	Lower Bound	Upper Bound
1	(Constant)	-1.378	.897		-1.537	.136	-3.215	.458
	Packed Cell Volume (%)	.117	.014	.839	8.584	.000	.089	.145
	Plasma Fibrinogen (mg/100ml)	.000	.000	.111	1.147	.261	.000	.001
	Plasma Protein (g/100ml)	.040	.097	.037	.412	.683	159	.239
a. De	ependent Variable: Blood Visc	cosity (cP)	/					

The coefficient and standard error for Plasma Fibrinogen are not exactly zero, but are zero to three decimal places.

Only the Packed Cell Volume coefficient is significantly different from zero (p < 0.001). The intercept (Constant), Plasma Fibrinogen, and Plasma Protein coefficients are not significantly different from zero (p=0.136, 0.261, 0.683 respectively)

Results: Scatterplot of Standardized Residual vs Predicted Value

Scatterplot

Dependent Variable: Blood Viscosity (cP)



Obtaining: Plots of Residuals vs Covariates

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15 15	4.41	50.00	P-P		4.86448	45448	-1.49650	4.65413	5.07484	4.20779	5.52118									
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25 25	5.12	49.50	392	5.49	4.78188	.33812	1.11336	4.62664	4.93711	4.14070	5.42305									
26 26	5.15	56.00	352	5.41	5.52195	37195	-1.22473	5.22653	5.81736	4.83327	6.21062									
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Main Effect Model: Significant Factor Effect (different intercept in both groups, slope=0)



28

Main Effect Model: Significant Covariate Effect (intercept, slope same in both groups)



Main Effect Model: Significant Covariate and Factor Effect (intercept different, slope same in the two groups)



Interaction Model: Covariate, Factor and Interaction Effect (different intercept and slope in the two groups)



31





glutest

Log-scale Data



Log-scale Data



Subgroups













Group 3

log(glutest)

Fits to the three subgroups



Projection back to the axis



log(glufast)

FACTOR PREDICTOR REGRESSION

We need to take some care when combining factor predictors and covariates in the regression model. Suppose that we have only two predictors

- A covariate, x_1
- A factor predictor, x_2 , now taking L levels, with the levels being indexed by l = 1, 2, ..., L.

We want to build a model that takes into account both x_1 and x_2 .

Example : Binary Factor L = 2

Suppose that factor predictor x_2 takes two levels, labelled 0 and 1, that identify two data subgroups. Five models can be considered, that correspond to different straight-line models

- MODEL 0 : Same intercept, slope zero, in the two subgroups
- MODEL 1 : Different intercept, slope zero, in the two subgroups
- MODEL 2 : Same intercept, same non-zero slope, in the two subgroups
- MODEL 3 : Different intercept, same non-zero slope, in the two subgroups
- MODEL 4 : Different intercept, different non-zero slopes, in the two subgroups

We can write out the models in terms of the usual slope and intercept parameters. The general model can be written

$$y = \begin{cases} \beta_{00} + \beta_{01}x_1 + \epsilon & \text{GROUP 0} \quad (l = 0) \\ \beta_{10} + \beta_{11}x_1 + \epsilon & \text{GROUP 1} \quad (l = 1) \end{cases}$$

• MODEL 0 :

$$\beta_{00} = \beta_{10} = \beta_0, \ \beta_{01} = \beta_{11} = 0$$

$$\beta_{00} \neq \beta_{10}, \ \beta_{01} = \beta_{11} = 0$$

- MODEL 1 :
- MODEL 2 :
- MODEL 3 :
- MODEL 4 :

$$\beta_{00} \neq \beta_{10}, \ \beta_{01} = \beta_{11} = 0$$

- $\beta_{00} = \beta_{10} = \beta_0, \ \beta_{01} = \beta_{11} = \beta_1 \neq 0$
- $\beta_{00} \neq \beta_{10}, \ \beta_{01} = \beta_{11} = \beta_1 \neq 0$
- $\beta_{00} \neq \beta_{10}, \ \beta_{01} \neq \beta_{11}$

The numbers of parameters, p, in each model are as follows:

MODEL 0	•	p = 1	eta_0
MODEL 1	•	p=2	eta_{00},eta_{10}
MODEL 2	•	p=2	eta_0,eta_1
MODEL 3	•	p = 3	eta_{00},eta_{10},eta_1
MODEL 4	:	p = 4	$eta_{00},eta_{10},eta_{10},eta_{11}$

SPSS Parameterization: The default parameterization used by SPSS is different from the one described above. SPSS takes a baseline group, and looks for **differences** in the parameters compared to the baseline group. The baseline group is taken to be the last listed subgroup for the factor predictor; in the binary example above, the baseline group would be Group 1.

The interaction model is therefore written

$$y = [\beta_0 + (1 - x_2)\delta_{00}] + [(\beta_1 + (1 - x_2)\delta_{01})x_1] + \epsilon$$

- δ_{00} is the **change in intercept** from Group 1 to Group 0
- δ_{01} is the **change in slope** from Group 1 to Group 0

Example: Diabetes Data Set

The data in the data set **DIABETES.SAV** contain information on 68 diabetes patients falling into two clinically different categories (overt and chemical diabetics) and 76 normal controls. Measurements of plasma glucose in blood samples when fasting and in a dietary test are recorded.

The objective is to predict the test glucose levels from the fasting glucose levels in the three subgroups, and to find out if there is any significant difference between the subgroups. In this analysis, there is a single response variable, one covariate and one factor predictor:

- y : glutest, the test glucose level
- x_1 : covariate **glufast**, the fasting glucose level
- x_2 : factor predictor **group**, the diabetes group
 - GROUP 1: Overt Diabetic
 - GROUP 2: Chemical Diabetic
 - GROUP 3: Normal Patients

Tests of Between-Subjects Effects

Dependent Variable: Log(GluTest)

	Type III Sum				
Source	of Squares	df	Mean Square	F	Sig.
Corrected Model	27.187 ^a	5	5.437	569.463	.000
Intercept	.973	1	.973	101.906	.000
group	.104	2	.052	5.447	.005
loggluf	.675	1	.675	70.702	.000
group * loggluf	.155	2	.077	8.099	.000
Error	1.318	138	.010		
Total	5509.040	144			
Corrected Total	28.504	143			

a. R Squared = .954 (Adjusted R Squared = .952)

Parameter Estimates

Dependent Variable: Log(GluTest)

					95% Confide	ence Interval
Parameter	В	Std. Error	t	Sig.	Lower Bound	Upper Bound
Intercept	4.504	.559	8.060	.000	3.399	5.608
[group=1]	-2.037	.619	-3.289	.001	-3.262	813
[group=2]	-1.436	.958	-1.499	.136	-3.330	.458
[group=3]	0 ^a	-	-			
loggluf	.299	.124	2.414	.017	.054	.544
[group=1] * loggluf	.535	.134	4.001	.000	.270	.799
[group=2] * loggluf	.382	.210	1.820	.071	033	.797
[group=3] * loggluf	0 ^a	•				

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

The first ANOVA table demonstrates that there is a significant interaction between the covariate and the factor predictor (F = 8.099, p-value < 0.001). This means that there is a significantly different slope in at least two of the three subgroups.

The second table gives the slope and intercept parameters in the three groups. The SPSS parameterization is not directly in terms of the slopes and intercepts, but looks at **differences** from baseline subgroup, Group 3. For example, the Group 1 intercept and slope are, respectively,

INTERCEPT : 4.504 + (-2.037) = 2.467 SLOPE : 0.299 + 0.535 = 0.834.

Diabetes Data Set

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Create two new variables **loggluf** and **logglut** for the logged variables

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4 glutest	Numeric	11 0		Test Plasma Glucose	None	None	8	Right	Scale
5 instest	Numeric	11 0		Plasma Insulin during Test	None	None	8	Right	Scale
6 sspg	Numeric	11 0		Steady State Plasma Glucose	None	None	8	Right	Scale
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In Target Variable insert loggluf, and in Numeric Expression type

Ln(glufast), and click OK



SPSS Processor is ready

Click OK when the confirmation screen appears



The log transformed variable **loggluf** is computed.

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The same procedure computes the log transformed variable **logglut**; we log transform the glutest variable using the *Compute* pulldown

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We now perform the linear regression using the General Linear Model pulldown.⁸

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Select the *Dependent Variable* (logglut), the *Fixed Factor* (group) and the *Covariate* (loggluf).

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To specify the model, click the Model button to get the Model Dialog.

We wish to specify a *Custom* main effects plus interaction model.



We select the factor and covariate as main effects.

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The model is now built. On the General Linear Model dialog, click Options.

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E	Observed power	er	R	Residual plot		Normal	4.38	5.97															
	Parameter estin	nates		ack of fit		Normal	4.50	5.90			-					-						-	
	Contrast coefficient matrix General estimable function		Normal	4.44	5.69																		
Sig	nificance level:	.05 0	onfidence inter	rvals are 95%		Normal	4.50	5.84															
	, t		Cartin			Normal	4.50	5.93															
			Continu	ue Cancel		Normal	4.48	5.72															
23	.95	95	347	184	91	Normal	4.55	5.85															
24	.97	90	327	192	124	Normal	4.50	5.79															
25	1.12	92	365	279	235	Normal	4.52	5.96			-					-		-		-			
27	1.20	98	365	145	158	Normal	4.58	5.90			-												
28	1.13	100	352	172	140	Normal	4.61	5.86															
29	1.00	86	325	179	145	Normal	4.45	5.78															
30	.78	98	321	222	99	Normal	4.58	5.77															
31	1.00	70	360	134	90	Normal	4.25	5.89															
32	1.00	99	336	143	105	Normal	4.60	5.82															
34	.71	90	353	263	165	Normal	4.52	5.00			-					-				-		-	
35	.89	85	373	174	78	Normal	4.44	5.92															
36	.88	99	376	134	80	Normal	4.60	5.93			1												
37	1.17	100	367	182	54	Normal	4.61	5.91															
38	.85	78	335	241	175	Normal	4.36	5.81															
39	.97	106	396	128	80	Normal	4.66	5.98															
40	1.00	98	2//	165	186	Normal	4.58	5.62															
41	89	90	360	282	160	Normal	4.02	5.89			-					-						-	
43	.98	94	291	94	71	Normal	4.54	5.67															
44	.78	80	269	121	29	Normal	4.38	5.59															
45	.74	93	318	73	42	Normal	4.53	5.76															
46	.91	86	328	106	56	Normal	4.45	5.79															
47	.95	85	334	118	122	Normal	4.44	5.81															
48	.95	96	35b 291	112	122	Normal	4.55	5.8/															
45	87	87	360	292	122	Normal	4.40	5.67				1								-			
51	.87	94	313	200	233	Normal	4.54	5.75			-												
52	1.17	93	306	220	132	Normal	4.53	5.72															
53	.83	86	319	144	138	Normal	4.45	5.77															
54	.82	86	349	109	83	Normal	4.45	5.86														-	
Sec. Sec.	Bb	, 461	4471	161	104	Normal	4 56	5.81														d	

15

The output is generated.

Diabetes.sav [DataSet1] - SPSS Data Editor												
File Edit View Data Transform Analyze Graphs Utilities Window Help												
13:												
1 1 .81 80 356 124	<u>sspa</u> group loggut loggut var											
2 2 .95 97 289 117	76 Normal 4.57 5.67 5.67											
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12 12 .73 87 Title												
13 13 .96 78 Active Datase	set [DataSet1] C:\Work\Courses\204\SPSS\D1abetes.sav											
15 15 .74 86 Tests of Between-Sut	Alpeta Factors											
16 16 .98 80	stimates Between-Subjects Factors											
17 17 1.10 90 Coserved * P	VeluceLabel N											
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26 26 1.11 74	Dependent Verieble: Local Linear											
28 28 1.13 100												
29 29 1.00 86	Source of Squares of Mean Square F Sig.											
30 30 .78 98	Unterceted woulder 27.187* 5 54.37 569.463 J.000											
31 31 1.00 70	group .104 2 .052 5.447 .005											
33 33 .71 75	loggluf 875 1 675 70.702 000											
34 34 .76 90	group noggini .100 2 .07 0.099 .000											
35 35 .89 85	Total 5509.040 144											
37 37 1.17 100	Corrected Total 28.504 143											
38 38 .85 78	(-ce:=Densups x busice(x - ce) (-ce:=Densups x busice(x - ce) (-ce)											
39 39 .97 106												
	Parameter Estimates											
42 42 .89 90	Dependent Verlehber Log (NUTech											
43 4398 94	Dependent variable: Log(chures) 95% Confidence Interval											
44 44 .78 80	Parameter B Std. Error t Stg. Lower Bound Upper Bound											
46 46 .91 86												
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48 48 .95 96 356 112	73 Normal 4.56 5.87											
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51 51 .87 94 313 200	233 Normal 4.54 5.75											
5 2 52 1.17 93 306 220	132 Normal 4.63 5.72											
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Variable View /												
66	SPSS Processor is ready											

The ANOVA table describes the results. It can be read in the same way as an ¹⁷ ordinary ANOVA table. We note significant main effects and interaction.

Tests of Between-Subjects Effects

Dependent Variable: Log(GluTest)

	Type III Sum				
Source	of Squares	df	Mean Square	F	Sig.
Corrected Model	27.187 ^a	5	5.437	569.463	.000
Intercept	.973		.973	101.906	.000
group	.104	2	.052	5.447	.005
loggluf	.675	1	.675	70.702	.000
group * loggluf	.155	2	.077	8.099	.000
Error	1.318	138	.010		
-Total	5509.040	144			
Corrected Total	28.504	143			

a. R Squared = .954 (Adjusted R Squared = .952)

The high R squared value means that the model fit is quite good overall.

The parameter estimates/standard errors are also computed.

The SPSS parameterization of the model is used.

Parameter Estimates

Dependent Variable: Log(GluTest)

					95% Confidence Interval				
Parameter	В	Std. Error	t	Sig.	Lower Bound	Upper Bound			
Intercept	4.504	.559	8.060	.000	3.399	5.608			
[group=1]	-2.037	.619	-3.289	.001	-3.262	813			
[group=2]	-1.436	.958	-1.499	.136	-3.330	.458			
[group=3]	0 ^a								
loggluf	.299	.124	2.414	.017	.054	.544			
[group=1] * loggluf	.535	.134	4.001	.000	.270	.799			
[group=2] * loggluf	.382	.210	1.820	.071	033	.797			
[group=3] * loggluf	0 ^a								

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

In the main effects plus interaction model, there are six parameters; we are fitting three separate straight lines to the three subgroups, and there are two parameters in each straight line.



Dependent Variable: Log(GluTest)

Model: Intercept + group + loggluf + loggluf