

Directions: Neatly and thoroughly answer each of the following six problems in the space given, clearly showing all work. Unless otherwise noted, please use $\alpha = 5\%$. Keep four decimal places in your calculations.

1. (2 + 2 + 2 = 6 points) Pawitan (2001:154) presents data related to 40 individuals who underwent an experimental surgery, which are analyzed on p.1 of the *Appendix*. The dataset includes the age of the patient at the time of the surgery and whether ($Y = 1$) or not ($Y = 0$) the patient died within the first 30 days immediately following the surgery.
 - (a) Clearly interpret the “Point Estimate” of the odds ratio.

 - (b) Clearly give the ramifications/implications of the 95% (Wald) confidence interval for the odds ratio.

 - (b) Give the predicted probability of survival (not dying) in the first 30 days post surgery for a 65-year-old.

2. (1 + 2.5 + 1 = 4.5 points) In a genetic study of an insect population which damages crops, a random sample of four male insects were each mated with two (different) females chosen at random. The weights (in g) of three eggs produced by each of the eight mated females were measured. The data for this nested design are analyzed in Minitab on the bottom of p.1 of the *Appendix*.
 - (a) Briefly indicate why this is a nested design.

 - (b) Test whether there is a difference between the Males in terms of the average egg weights.

Null hypothesis _____

Alternative hypothesis _____

Test stat, df and p-value _____

Detailed conclusion _____

(c) Give the point estimate of the Males variance component, s_M^2 .

3. (2.5 + 2.5 = 5 points) Finney (1978:48) presents data related to the growth response of *Lactobacillus leichmannii* to one of eight different doses of vitamin B₁₂ each replicated in 6 independent test tubes. These data are analyzed on pp.2-3 of the *Appendix*, first using PROC NLIN and the resulting residual plot, and then using two PROC NLMixed's.

(a) Write out the assumed model function in terms of x = dose used in the NLIN and identify the roles of the parameters.

Model Function _____

Role of θ_1 ("th1") _____

Role of θ_2 ("th2") _____

Role of θ_3 ("th3") _____

(b) After viewing the residual plot, the two NLMixed's were then run. Using the corresponding output, perform a test of homoskedasticity clearly writing out your hypotheses, the calculated test statistic, degrees of freedom (df), p-value, and your detailed conclusion.

Null hypothesis _____

Alternative hypothesis _____

Test stat, df and p-value _____

Clear conclusion

4. (1 + 1 + 2.5 = 4.5 points) A study was conducted involving 8 subjects randomized to an experimental treatment and 8 subjects randomized to a control treatment, and in which respiration (liters of air/min/m² of body area) was measured both at rest and after 30 minutes of work. The data are plotted and analyzed on pp. 4-5 of the *Appendix*. Two separate lines (one for each treatment) are fit in the PROC REG at the top of p.5.

(a) Write down the fitted line for the experimental group

(b) Write down the fitted line for the control group

(c) Using $\alpha = 1\%$ **here**, test whether **one line can be fit to both the experimental and control groups**.

Null hypothesis _____

Alternative hypothesis _____

Test stat, df and p-value _____

Clear conclusion

5. (2.5 + 2.5 = 5 points) Schwenke & Milliken (1991) report that in meat science research, a pH level of 6.0 in the steer muscle is desired before processing begins, that at the time of slaughter pH is usually in the range of 7.0-7.2, and that pH decreases over time. In general, however, it is not practical to monitor the pH decline to determine precisely when pH of 6.0 is attained, and they conducted a study to determine the time when pH 6.0 is reached for two treatments, the **conventional treatment** (denoted '**conv**') and an **electrically simulated hot boned treatment** (denoted '**eshb**'). The study data corresponding to the postmortem pH for the *longissimus dorsi* muscle of 24 steer carcasses, 12 steers randomized to each of the treatments and measured only at 1, 2, 4, 6, 8 or 24 hours postmortem, are graphed on the bottom of p.5 of the *Appendix*. The suggested model for these data is $y = \eta(x, \theta) + \epsilon$, where x = time postmortem, and

$$h(x, q) = q_2 + (q_1 - q_2) \exp \{-q_3 x\} \quad (1.1)$$

Note that θ_1 is then the value of the expected pH at time $x = 0$ and θ_2 is the lower asymptote (at infinity).

- (a) Using the appropriate outputs from the first three NLIN runs on pp. 6-7 of the *Appendix*, test whether (only) the lower asymptotes are the same for the two treatments, giving your hypotheses, test statistic, p-value and conclusion.

Null hypothesis _____

Alternative hypothesis _____

Test stat, df and p-value _____

Clear conclusion

- (b) Using the appropriate outputs from the first three NLIN runs on pp. 6-7 of the *Appendix*, test whether **both** the lower and upper asymptotes are the same for the two treatments, giving your hypotheses, test stat., p-value and conclusion.

Null hypothesis _____

Alternative hypothesis _____

Test stat, df and p-value _____

Clear conclusion

The rest of the exam for G students only!!

- (c) (1.5 points) Write out the model function being fit in the fourth NLIN program and output (p.8) in the *Appendix*, and indicate the role of the new (ϕ = phi) parameter.

- (d) (1.5 points) For each treatment, how many hours postmortem should the meat scientists wait for processing in order that the pH be nearest to 6.0? Give approximate 95% confidence intervals.

For Conventional treatment _____

For ESHB treatment _____

First Problem –SAS Program and Output

AB Exam 1 Appendix

```
data onea;
  y=0; input age @@; datalines;
50 50 51 51 53 54 54 54 55 55 56 56 56 57 57 58 60 61 62 63 63 64 65 68
69 71
;
data oneb;
  y=1; input age @@; datalines;
57 57 59 61 61 62 62 62 63 64 67 67 68 70
;
data one;
  set onea oneb;
proc logistic descending;
  model y=age;
run;
```

The LOGISTIC Procedure Response Profile

Ordered Value	y	Total Frequency
1	1	14
2	0	26

Probability modeled is y=1.

Model Fit Statistics

Criterion	Intercept Only	Intercept and Covariates
AIC	53.796	49.301
SC	55.485	52.679
-2 Log L	51.796	45.301

Testing Global Null Hypothesis: BETA=0

Test	Chi-Square	DF	Pr > ChiSq
Likelihood Ratio	6.4949	1	0.0108
Score	6.1508	1	0.0131
Wald	5.3899	1	0.0203

Analysis of Maximum Likelihood Estimates

Parameter	DF	Estimate	Standard Error	Wald Chi-Square	Pr > ChiSq
Intercept	1	-10.4814	4.3042	5.9301	0.0149
age	1	0.1629	0.0702	5.3899	0.0203

Odds Ratio Estimates

Effect	Point Estimate	95% Wald Confidence Limits
age	1.177	1.026 1.351

Second Problem – Minitab Output

Minitab Output

Nested ANOVA: y versus male, female

Analysis of Variance for y

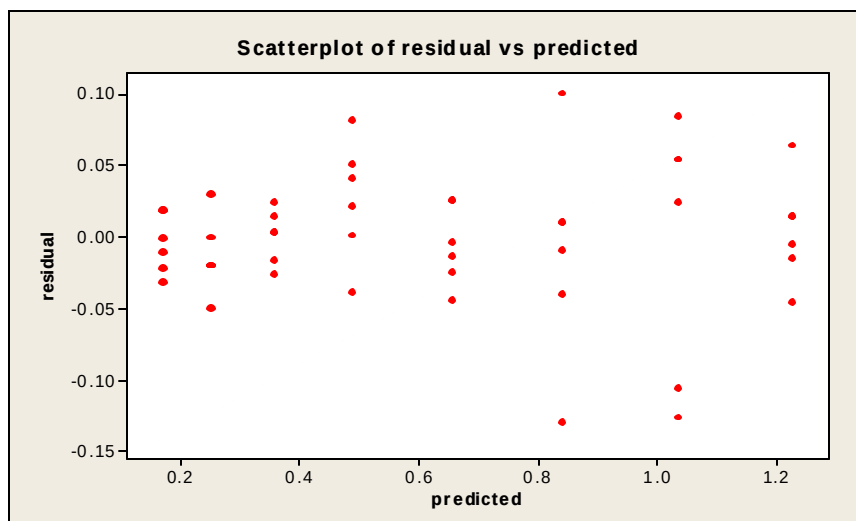
Source	DF	SS	MS	EMS
male	3	16.8913	5.6304	$s^2 + 3s_F^2 + 6s_M^2$
female	4	10.1450	2.5362	$s^2 + 3s_F^2$
Error	16	16.5933	1.0371	s^2
Total	23	43.6296		

Third Problem – SAS Program and Output

The Original NLIN Run

```
data one;
  do dose=0.23,0.35,0.53,0.79,1.19,1.78,2.67,4;
    do rep=1 to 6;
      input y @@; y=y/100; drop rep; output;
    end; end; datalines;
15 14 19 19 17 16 28 20 23 25 23 23 36 36 34 37 33 38 51 53 54 45 57 49
68 63 64 61 65 68 85 80 71 85 94 83 106 91 109 93 109 112 121 122 129 124 118 124
;
proc nlin data=one;
  parms th1=2 th2=2 th3=2;
  t=(dose/th2)**th3; den=1+t; mean=th1*t/den;
  model y=mean;
run;
```

Residual Plot



The First NLMixed

```
proc nlmixed data=one;
  parms th1=2 th2=1.5 th3=2 se=0.1 rho=1;
  t=(dose/th2)**th3; den=1+t; mean=th1*t/den; var=se*se*mean**rho;
  model y~normal(mean,var);
run;
```

The NLMIXED Procedure

Dimensions

Observations Used	48
Observations Not Used	0
Total Observations	48
Parameters	5

NOTE: GCONV convergence criterion satisfied.

Fit Statistics

-2 Log Likelihood	-174.8
AIC (smaller is better)	-164.8
AICC (smaller is better)	-163.3
BIC (smaller is better)	-155.4

Parameter Estimates

Parameter	Estimate	Standard Error	DF	t Value	Pr > t	Alpha	Lower	Upper	
Gradient									
th1	1.8021	0.1480	48	12.18	<.0001	0.05	1.5045	2.0996	-
0.00142									
th2	1.9995	0.3263	48	6.13	<.0001	0.05	1.3435	2.6556	
0.000703									
th3	1.0597	0.05145	48	20.60	<.0001	0.05	0.9562	1.1631	
0.000553									
se	0.06153	0.008886	48	6.92	<.0001	0.05	0.04366	0.07939	
0.001657									
rho	1.3608	0.3076	48	4.42	<.0001	0.05	0.7424	1.9792	-
0.00005									

The Second NLMixed

```
proc nlmixed data=one;
  parms th1=2 th2=1.5 th3=2 se=0.1; rho=0;
  t=(dose/th2)**th3; den=1+t; mean=th1*t/den; var=se*se*mean**rho;
  model y~normal(mean,var);
run;
```

The NLMIXED Procedure

Dimensions

Observations Used	48
Observations Not Used	0
Total Observations	48
Parameters	4

NOTE: GCONV convergence criterion satisfied.

Fit Statistics

-2 Log Likelihood	-159.2
AIC (smaller is better)	-151.2
AICC (smaller is better)	-150.3
BIC (smaller is better)	-143.8

Parameter Estimates

Parameter	Estimate	Standard Error	DF	t Value	Pr > t	Alpha	Lower	Upper
Gradient								
th1	1.9203	0.1854	48	10.36	<.0001	0.05	1.5474	2.2931
0.00041								
th2	2.2838	0.4627	48	4.94	<.0001	0.05	1.3535	3.2142
0.000232								
th3	1.0131	0.06874	48	14.74	<.0001	0.05	0.8749	1.1513
0.00032								
se	0.04606	0.004701	48	9.80	<.0001	0.05	0.03661	0.05552
0.01292								

Fourth Problem – Graph, SAS Program and Output



```

data one;
  do group='experim', 'control';
    egroup=(group='experim');
    do rep=1 to 8;
      input rest work @@; egress=egroup*rest; drop rep; output;
    end; end; datalines;
5.74 6.24 6.79 9.07 5.32 7.77 7.18 16.46 5.60 6.95 6.06 8.14 6.32 11.72 6.34 8.06
6.21 5.50 4.50 4.64 4.86 4.61 4.78 3.78 4.79 5.41 5.70 5.32 5.41 4.54 6.08 5.98
;
proc print; run;

```

Obs	group	egroup	rest	work	egrest
1	experim	1	5.74	6.24	5.74
2	experim	1	6.79	9.07	6.79
3	experim	1	5.32	7.77	5.32
4	experim	1	7.18	16.46	7.18
5	experim	1	5.60	6.95	5.60
6	experim	1	6.06	8.14	6.06
7	experim	1	6.32	11.72	6.32
8	experim	1	6.34	8.06	6.34
9	control	0	6.21	5.50	0.00
10	control	0	4.50	4.64	0.00
11	control	0	4.86	4.61	0.00
12	control	0	4.78	3.78	0.00
13	control	0	4.79	5.41	0.00
14	control	0	5.70	5.32	0.00
15	control	0	5.41	4.54	0.00
16	control	0	6.08	5.98	0.00

```

proc reg; model work=rest egroup egress; run;

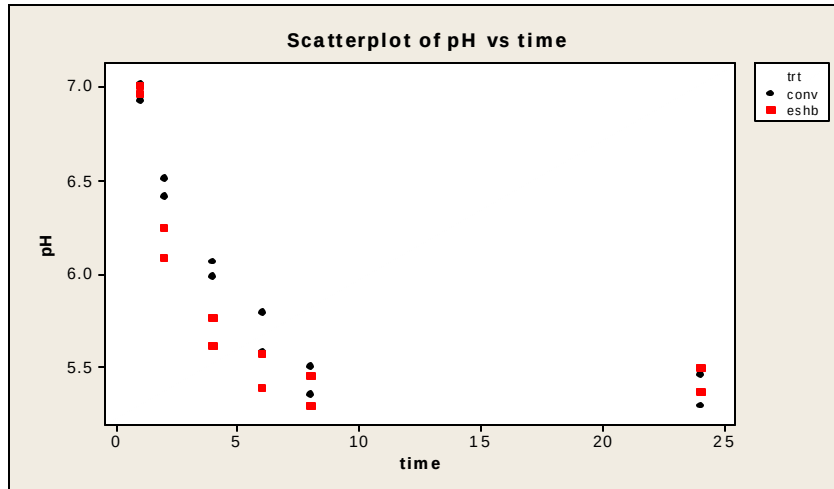
```


The REG Procedure						
Dependent Variable: work						
Analysis of Variance						
	Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
0.0002	Model	3	124.04476	41.34825	15.68	
	Error	12	31.64918	2.63743		
	Corrected Total	15	155.69394			
	Root MSE		1.62402	R-Square	0.7967	
	Dependent Mean		7.13688	Adj R-Sq	0.7459	
	Coeff Var		22.75530			
Parameter Estimates						
	Variable	DF	Parameter Estimate	Standard Error	t Value	Pr > t
	Intercept	1	1.09182	5.01742	0.22	0.8314
	rest	1	0.73342	0.94202	0.78	0.4513
	egroup	1	-17.66140	7.91656	-2.23	0.0455
	egrest	1	3.46044	1.36533	2.53	0.0262

```
proc reg; model work=rest; run;
```

The REG Procedure						
Dependent Variable: work						
Analysis of Variance						
	Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
0.0004	Model	1	94.09904	94.09904	21.39	
	Error	14	61.59491	4.39964		
	Corrected Total	15	155.69394			
	Root MSE		2.09753	R-Square	0.6044	
	Dependent Mean		7.13688	Adj R-Sq	0.5761	
	Coeff Var		29.39005			
Parameter Estimates						
	Variable	DF	Parameter Estimate	Standard Error	t Value	Pr > t
	Intercept	1	-11.65126	4.09626	-2.84	0.0130
	rest	1	3.27891	0.70900	4.62	0.0004

Fifth Problem – Graph, SAS Program and Output



The First NLIN

```
data one;
  do time=1,2,4,6,8,24;
    do trt='conv', 'eshb';
      trtc=(trt='conv'); trte=(trt='eshb');
      do rep=1,2;
        input ph @@; drop rep; output;
      end; end; end; datalines;
7.02 6.93 7.01 6.96 6.42 6.51 6.25 6.09 6.07 5.99 5.77 5.62
5.59 5.80 5.57 5.39 5.51 5.36 5.46 5.30 5.30 5.47 5.37 5.50
;
proc nlin data=one;
  parms th1c=7 th1e=7 th2c=5 th2e=5 th3c=0.5 th3e=0.5;
  th1=th1c*trtc+th1e*trte; th2=th2c*trtc+th2e*trte; th3=th3c*trtc+th3e*trte;
  model ph=th2+(th1-th2)*exp(-th3*time);
run;
```

The NLIN Procedure
 Dependent Variable ph
 NOTE: Convergence criterion met.

Source	DF	Sum of Squares	Mean Square	F Value	Approx Pr > F
Model	5	7.8707	1.5741	163.09	<.0001
Error	18	0.1737	0.00965		
Corrected Total	23	8.0445			

Parameter	Estimate	Approx Std Error	Approximate 95% Confidence Limits	
th1c	7.5714	0.1394	7.2785	7.8643
th1e	8.4361	0.3304	7.7421	9.1302
th2c	5.3530	0.0635	5.2196	5.4865
th2e	5.4196	0.0453	5.3244	5.5148
th3c	0.3229	0.0408	0.2371	0.4087
th3e	0.6658	0.0916	0.4734	0.8581

The Second NLIN

```
proc nlin data=one;
  parms th1c=7 th1e=7 th2=5 th3c=0.5 th3e=0.5;
  th1=th1c*trtc+th1e*trte; th3=th3c*trtc+th3e*trte;
  model ph=th2+(th1-th2)*exp(-th3*time);
run;
```

The NLIN Procedure
 Dependent Variable ph
 NOTE: Convergence criterion met.

Source	DF	Sum of Squares	Mean Square	F Value	Approx Pr > F
Model	4	7.8636	1.9659	206.51	<.0001
Error	19	0.1809	0.00952		
Corrected Total	23	8.0445			

Parameter	Estimate	Approx Std Error	Approximate 95% Confidence Limits	
th1c	7.6072	0.1386	7.3171	7.8974
th1e	8.3537	0.2958	7.7345	8.9729
th2	5.3950	0.0369	5.3178	5.4723
th3c	0.3432	0.0359	0.2680	0.4184
th3e	0.6339	0.0794	0.4678	0.8000

The Third NLIN

```
proc nlin data=one;
  parms th1=7 th2=5 th3c=0.5 th3e=0.5;
  th3=th3c*trtc+th3e*trte;
  model ph=th2+(th1-th2)*exp(-th3*time);
run;
```

The NLIN Procedure

Dependent Variable ph

NOTE: Convergence criterion met.

Source	DF	Sum of Squares	Mean Square	F Value	Approx Pr > F
Model	3	7.7979	2.5993	210.87	<.0001
Error	20	0.2465	0.0123		
Corrected Total	23	8.0445			

Parameter	Estimate	Std Error	Approximate 95% Confidence Limits
th1	7.7938	0.1478	7.4856 8.1021
th2	5.3806	0.0442	5.2884 5.4729
th3c	0.3728	0.0415	0.2862 0.4594
th3e	0.5019	0.0560	0.3851 0.6188

The Fourth NLIN

```
proc nlin data=one;
  parms th2=5 th3c=0.5 th3e=0.5 phic=3 phie=3;
  th3=th3c*trtc+th3e*trte; phi=phic*trtc+phie*trte;
  model ph=th2+(6-th2)*exp(-th3*(time-phi));
run;
```

The NLIN Procedure

Dependent Variable ph

NOTE: Convergence criterion met.

Source	DF	Sum of Squares	Mean Square	F Value	Approx Pr > F
Model	4	7.8636	1.9659	206.51	<.0001
Error	19	0.1809	0.00952		
Corrected Total	23	8.0445			

Parameter	Estimate	Std Error	Approximate 95% Confidence Limits
th2	5.3950	0.0369	5.3178 5.4723
th3c	0.3432	0.0359	0.2680 0.4184
th3e	0.6339	0.0794	0.4678 0.8000
phic	3.7779	0.2141	3.3298 4.2260
phie	2.5039	0.1474	2.1954 2.8125

