STATISTICS 3N03 & 3J04 ASSIGNMENT 5 SOLUTIONS 2006-11-24 Updated 2006-12-12

Question 1

(a) You need at least 29 degrees of freedom.

```
> for (df in 25:30) print(c(df,qchisq(.995,df)/qchisq(.005,df)))
[1] 25.000000      4.460974
[1] 26.000000      4.326958
[1] 27.000000      4.204493
[1] 28.000000      4.092128
[1] 29.000000      3.988646
[1] 30.000000      3.893019
```

(b) Here, $\alpha = 0.01$, $\beta = 0.10$, $\delta = 0.5$ and $\sigma = 1.3$, so the required sample size n is given by

> ((qnorm(1-(0.01)/2) + qnorm(1-0.1))*1.3/0.5)^2
[1] 100.5847

or, using table values from the text

> ((2.576 + 1.282)*1.3/0.5)^2
[1] 100.6169

so 101 observations would be required.

The probability of a Type II error is computed by text formula (9-17); when n = 10 it gives

> pnorm(qnorm(1-(0.01)/2)-0.5*sqrt(10)/1.3)
+ pnorm(-qnorm(1-(0.01)/2)-0.5*sqrt(10)/1.3)
[1] 0.9130915

and this probability is much too high for the test to be useful.

(c) One-sided test:

```
> ((qnorm(.95)+qnorm(.95))*2/(11.5-12))^2
[1] 173.1548
> 1 - pnorm(-qnorm(.95)-(11.5-12)*sqrt(50)/2)
[1] 0.4510879
```

The number of paint samples required to reduce the Type II error rate to 5% is 174; if you could only test 50 samples, the Type II error rate would be 45% which is too high for the test to be useful.

Two-sided test:

```
> ((qnorm(.975)+qnorm(.95))*2/(11.5-12))^2
[1] 207.9154
> pnorm(qnorm(.975)-(11.5-12)*sqrt(50)/2)
- pnorm(-qnorm(.975)-(11.5-12)*sqrt(50)/2)
```

[1] 0.5761095

The number of paint samples required to reduce the Type II error rate to 5% is 208; if you could only test 50 samples, the Type II error rate would be 57.6% which is too high for the test to be useful. *(10 marks for <u>either</u> 1-sided or 2-sided answer)*

(d) Assuming that the flaws occur independently, at random, at a constant average rate over the windshield (i.e. as a Poisson process), the number of flaws per windshield will follow a Poisson distribution. The probability it was produced at Plant A given that it has 3 flaws is found by Bayes' Theorem to be 20.8%. *(11 marks)*

Let A be the event it was produced at Plant A, let X be the number of flaws.

P(A|X=3) = P(X=3|A)P(A)/[P(X=3|A)P(A) + P(X=3|B)P(B)] > dpois(3,2.1)*0.2/(dpois(3,2.1)*0.2 + dpois(3,4.3)*0.8) [1] 0.2081147

(*a*) Acceptable analyses: Paired t-test, 2-factor ANOVA without interaction, sign test, simple linear regression.

Acceptable graphs: dot, box or stem-leaf plot of differences; interaction plot; scatter plot with fitted line. (Graph 2, suitable analysis 2, correct calculation 4, assumptions 2, conclusions 2)

Paired t-test

Assumptions: Differences independent (can't test), normal (looks OK on dot plot and stem & leaf plot).

Conclusions: There is no evidence (P = 0.37) from these data of a difference in mean time between the two processors.

```
> stem(procspeed$diff)
 The decimal point is 1 digit(s) to the right of the |
  -0 | 8
  -0 | 310
  0 | 03
> procspeed
 code procA procB diff
1
   1 27.2 24.1 3.1
    2 18.1 19.3 -1.2
2
3
    3 27.2 26.8 0.4
4
    4 19.7 20.1 -0.4
5
    5 24.5 27.6 -3.1
    6 22.1 29.8 -7.7
6
> t.test(procspeed$procA, procspeed$procB, pair=T)
       Paired t-test
data: procspeed$procA and procspeed$procB
t = -0.9921, df = 5, p-value = 0.3667
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
-5.326857 2.360190
sample estimates:
mean of the differences
             -1.483333
```

2-factor ANOVA

>	procspeed2					
	code	time	process			
1	1	27.2	A			
2	2	18.1	A			
3	3	27.2	A			
4	4	19.7	A			
5	5	24.5	A			
6	6	22.1	A			
7	1	24.1	В			
8	2	19.3	В			
9	3	26.8	В			
1() 4	20.1	В			
11	L 5	27.6	В			

```
12
      6 29.8
                   В
> anova(lm(time~code+process, procspeed2))
Analysis of Variance Table
Response: time
          Df Sum Sq Mean Sq F value Pr(>F)
code
          5 129.068 25.814 3.8488 0.0827 .
process
           1
               6.601
                       6.601
                              0.9842 0.3667
Residuals 5
              33.534
                       6.707
Signif. codes:
               0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Sign test

Assumptions: Differences are independent (can't test).

Conclusions: There are 2 positive differences out of 6 non-zero differences, so P = 0.69 and there is no evidence from these data of a difference in the median time between the two processors.

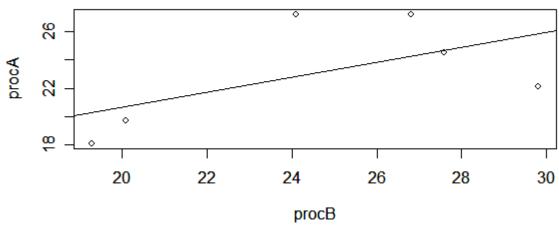
```
> 2*pbinom(2,6,.5)
[1] 0.6875
```

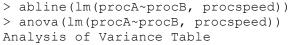
Simple Linear Regression

Assumptions: The time with Processor A is linearly related to the time with Processor B, data are normal and homoscedastic. We can't test any of these because the sample is small and there are no repeated x-values.

Conclusions: There is no evidence from these data of a linear relationship between Processor A and Processor B times.

```
> plot(procA~procB, procspeed)
```



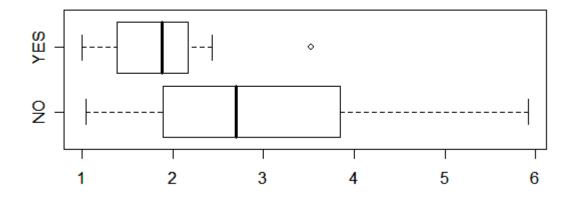


(b) Acceptable analyses: Two-sample t-test or 1-factor ANOVA. Acceptable graphs: <u>Comparative</u> dot, box or stem-leaf plots. (*Graph 2, suitable analysis 2, correct calculation 3, F-test of variances 2, assumptions 2, conclusions 2*)

Assumptions: Independence (can't test), normality (OK by plot), homoscedasticity (graph looks heteroscedastic; F-test gives no evidence (P = 0.14) that the variances are not equal).

Conclusions: There is no evidence from these data that either the mean or the variance of resilient modulus differs between rutted and non-rutted pavement.

>	ruts				
	resmod	rutted			
1	1.48	YES			
2	1.88	YES			
3	1.90	YES			
4	1.29	YES			
5	3.53	YES			
6	2.43	YES			
7	1.00	YES			
8	3.06	NO			
9	2.58	NO			
1	0 1.70	NO			
1	1 5.76	NO			
1:	2 2.44	NO			
1	3 2.03	NO			
1	4 1.76	NO			
1	5 4.63	NO			
1	6 2.86	NO			
1	7 2.82	NO			
1	8 1.04	NO			
1	9 5.92	NO			
>	boxplot	(resmod~	rutted,	ruts,	horizontal=T)



> t.test(resmod~rutted, ruts, var.eq=T)

Two Sample t-test

The only acceptable analysis is a 2-factor ANOVA with a test for interaction.

Assumptions: Normality, independence, homoscedasticity.

Conclusions: There is no evidence (P = 0.49) from these data of an interaction between metal type and sintering time. The interaction plots confirm this as the lines are parallel. There is strong evidence that the mean compressive strength is different for the different sintering times (P = 0.0003) and for the different metals (P = 0.0006).

The 95% confidence interval for the residual variance is (0.522, 4.199).

(Correct calculation with P-values 9, assumptions 2, conclusions 3, either interaction plot 3, CI for residual variance 3)

```
> sinter
    strength metal time
1
          17.1 1 100

      1
      100

      16.5
      1

      14.9
      1

      12.3
      2

      13.8
      2

      10.8
      2

2
3
4
5
6

      19.4
      1
      200

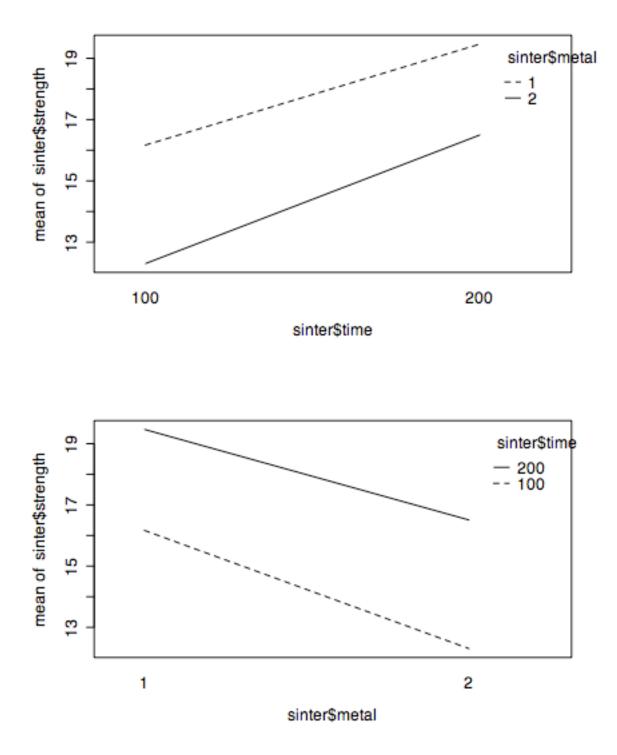
      18.9
      1
      200

      20.1
      1
      200

      15.6
      2
      200

7
8
9
10
          17.2220016.72200
11
12
> anova(lm(strength~metal*time, sinter))
Analysis of Variance Table
Response: strength
          Df Sum Sq Mean Sq F value
                                                           Pr(>F)
metal 1 35.021 35.021 30.608 0.0005522 ***
time 1 42.187 42.187 36.872 0.0002985 ***
metal
metal:time 1 0.608 0.608 0.531 0.4869859
Residuals 8 9.153 1.144
___
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> mse <- anova(lm(strength~metal*time, sinter))["Residuals","Mean Sq"]
> mse
[1] 1.144167
> mse/(qchisq(c(.975,.025),8)/8)
 [1] 0.5220171 4.1992954
> interaction.plot(sinter$time,sinter$metal,sinter$strength)
```

> interaction.plot(sinter\$metal,sinter\$time,sinter\$strength)

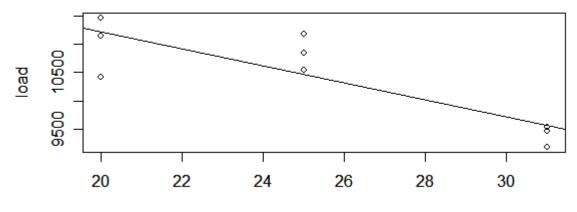


Assumptions: Independence, normality, homoscedasticity. The test of the slope also assumes linearity. Linearity is tested by the lack of fit F-test. Normality and Homoscedasticity can't be tested in such a small sample but look OK on the graph.

Conclusions: There is some evidence (P = 0.07) from these data that the relationship is non-linear. However, if we choose to ignore that and test the slope, there is strong evidence (P = 0.002) that the slope is not zero.

(Graph with line 4, regression analysis 7, lack of fit test 5, assumptions 3, conclusions 3, 99% confidence interval for MSE <u>or</u> MSPE 3)

> concrete load age 1 11450 20 2 10420 20 3 11142 20 4 10840 25 5 11170 25 6 10540 25 7 9470 31 9190 8 31 9540 9 31 > plot(load~age, concrete) > abline(lm(load~age, concrete))



```
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> mse <- anova(lm(load~age, concrete))["Residuals", "Mean Sq"]</pre>
> mse
[1] 212265.4
> mse/(qchisq(c(.995,.005),7)/7)
[1] 73275.32 1501995.83
> anova(lm(load~age+as.factor(age), concrete))
Analysis of Variance Table
Response: load
              Df Sum Sq Mean Sq F value Pr(>F)
               1 4039390 4039390 29.3341 0.001639 **
age
as.factor(age) 1 659642 659642 4.7903 0.071204 .
Residuals
              6 826216 137703
___
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> mspe <- anova(lm(load~age+as.factor(age), concrete))["Residuals", "Mean Sq"]</pre>
> mspe
[1] 137702.7
> mspe/(qchisq(c(.995,.005),6)/6)
[1] 44545.75 1222707.21
```

The analyses in original units and on a log scale give very similar results and lead to the same conclusion: the interaction is significant at the 5% level (or, better to say, $P \ll 0.001$ so there is very strong evidence of an interaction between frequency and environment). That means that both frequency and environment affect the crack growth rate, but the effect of the environment is different at different frequencies; the higher the frequency, the less difference the environment makes. Because the interaction is significant, we do not test the main effects.

The residual plots show that the residuals from the log-scale analysis follow a normal distribution more closely than residuals from the original-scale analysis. In the original scale, the 23rd observation is an outlier with a large negative residual.

>	cracks		
	growth	environ	freq
1	2.29	Air	10
2	2.47	Air	10
3	2.48	Air	10
4	2.12	Air	10
5	2.65	Air	1
6	2.68	Air	1
7	2.06	Air	1
8	2.38	Air	1
9	2.24	Air	0.1
10		Air	0.1
11		Air	0.1
12		Air	0.1
13		Water	10
14	2.05	Water	10
15	5 2.23	Water	10
16	5 2.03	Water	10
17	3.20	Water	1
18	3.18	Water	1
19	3.96	Water	1
20	3.64	Water	1
21	11.00	Water	0.1
22	2 11.00	Water	0.1
23		Water	0.1
24		Water	0.1
25	5 1.90	Saltwater	10
26		Saltwater	10
27		Saltwater	10
28		Saltwater	10
29		Saltwater	1
30		Saltwater	1
31		Saltwater	1
32		Saltwater	1
33	9.96	Saltwater	0.1
34	10.01	Saltwater	0.1
35	9.36	Saltwater	0.1
36	5 10.40	Saltwater	0.1

> anova(lm(growth~environ*freq, cracks)) Analysis of Variance Table Response: growth Df Sum Sq Mean Sq F value Pr(>F) 2 64.252 32.126 159.92 1.076e-15 *** environ 2 209.893 104.946 522.40 < 2.2e-16 *** freq environ:freq 4 101.966 25.491 126.89 < 2.2e-16 *** Residuals 27 5.424 0.201 ___ Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1 > plot(lm(growth~environ*freq, cracks))

Residuals vs Fitted 240 0<mark>31</mark> 0.5 0 o 00 **0** 0 0.0 8 000 Residuals 0 o œ ò0 -1.5 -1.0 -0.5 o o 23⁰ 2 6 4 8 10 Fitted values Im(growth ~ environ * freq)

12

