STA 100 Homework 5 Solution

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- 1. (a) H_0 : Death is independent of treatment (Surgery, Watchful Waiting). H_A : Death is dependent of treatment (Surgery, Watchful Waiting).
 - (b) See the following table for the row, column, and grand totals.

	Surgery	WW	Total
Died	83	106	189
Alive	264	242	506
Total	347	348	695

The expected frequencies are

$$e_1 = \frac{347 \times 189}{695} = 94.364, \quad e_2 = \frac{348 \times 189}{695} = 94.636,$$

 $e_3 = \frac{347 \times 506}{695} = 252.636, \quad e_4 = \frac{348 \times 506}{695} = 253.364.$

Therefore, the test statistic is

$$T = \sum_{i=1}^{4} \frac{(o_i - e_i)^2}{e_i}$$

= $\frac{(83 - 94.364)^2}{94.364} + \frac{(106 - 94.636)^2}{94.636} + \frac{(264 - 252.636)^2}{252.636} + \frac{(242 - 253.364)^2}{253.364}$
= 3.754.

The null distribution for the test statistic is χ_1^2 . The critical value for $\alpha = 0.05$ is thus $\chi_1^2(0.05) = 3.84$.

- (c) From chi-square Table with df = 1, we find that $P(\chi_1^2 > 2.71) = 0.10$ and $P(\chi_1^2 > 3.84) = 0.05$. The range of *p*-value is thus (0.05, 0.10).
- (d) Since *p*-value > $\alpha = 0.05$, we fail to reject the null at the 0.05 level of significance.
- (e) We support the claim that death is independent of treatment at the 0.05 level of significance.
- 2. (a) H_0 : Pain is independent of treatment (Angioplasty, Bypass Surgery). H_A : Pain is dependent of treatment (Angioplasty, Bypass Surgery).
 - (b) See the following table for the row, column, and grand totals.

	Α	В	Total
Pain	111	74	185
No pain	402	441	843
Total	513	515	1028

The expected frequencies are

$$e_1 = \frac{513 \times 185}{1028} = 92.32, \quad e_2 = \frac{515 \times 185}{1028} = 92.68,$$
$$e_3 = \frac{513 \times 843}{1028} = 420.68, \quad e_4 = \frac{515 \times 843}{1028} = 422.32.$$

Therefore, the test statistic is

$$T = \sum_{i=1}^{4} \frac{(o_i - e_i)^2}{e_i}$$

= $\frac{(111 - 92.32)^2}{92.32} + \frac{(74 - 92.68)^2}{92.68} + \frac{(402 - 420.68)^2}{420.68} + \frac{(441 - 422.32)^2}{422.32}$
= 9.20.

The null distribution for the test statistic is χ_1^2 . The critical value for $\alpha = 0.01$ is thus $\chi_1^2(0.01) = 6.63$.

- (c) From chi-square Table with df = 1, we find that $P(\chi_1^2 > 6.63) = 0.01$ and $P(\chi_1^2 > 10.83) = 0.001$. The range of *p*-value is thus (0.001, 0.01).
- (d) Since *p*-value $< \alpha = 0.01$, we reject the null at the 0.01 level of significance.
- (e) We cannot support the claim that pain is independent of treatment at the 0.01 level of significance.
- 3. (a) Here we have $n_1 = 84 + 87 = 171$, $n_2 = 2916 + 4913 = 7829$. The Wilson-adjusted sample proportions are

$$\tilde{p}_1 = \frac{84+1}{171+2} = 0.4913, \quad \tilde{p}_2 = \frac{2916+1}{7829+2} = 0.3725.$$

The standard error for $\tilde{p}_1 - \tilde{p}_2$ is

$$SE_{\tilde{p}_1-\tilde{p}_2} = \sqrt{\frac{0.4913 \times (1-0.4913)}{171+2} + \frac{0.3725 \times (1-0.3725)}{7829+2}} = 0.0384$$

The 95% confidence interval for $p_1 - p_2$ is thus

$$(0.4913 - 0.3725) \pm 1.96 \times 0.0384$$

or (0.0435, 0.1941).

- (b) We are 95% confident that the difference in the proportion of people who develop CHD between smokers and nonsmokers falls within the range of 0.0435 to 0.1941, with smokers having a higher proportion.
- (c) Yes, it suggests a dependence on CHD and smoking since the confidence interval does not contain 0.
- (d) No, it does not support the claim since 0.20 is not included in the interval.
- 4. (a) Here we have $I = 4, n = \sum_{i=1}^{4} n_i = 40$, and

$$\bar{Y} = \frac{\sum_{i=1}^{4} n_i \bar{Y}_i}{n}$$

= $\frac{10 \times 3.22 + 10 \times 3.57 + 10 \times 2.87 + 10 \times 2.98}{40}$
= 3.16.

It follows that

$$SSB = \sum_{i=1}^{4} n_i (\bar{Y}_i - \bar{Y})^2$$

= 10 × (3.22 - 3.16)² + 10 × (3.57 - 3.16)² + 10 × (2.87 - 3.16)² + 10 × (2.98 - 3.16)²
= 2.882.

and

SSW =
$$\sum_{i=1}^{4} (n_i - 1)s_i^2$$

= $(10 - 1) \times 0.54^2 + (10 - 1) \times 0.35^2 + (10 - 1) \times 0.21^2 + (10 - 1) \times 0.23^2$
= 4.600.

Therefore,

$$SSTO = SSB + SSW = 2.882 + 4.600 = 7.482$$

If follows that

$$MSB = \frac{SSB}{4-1} = 0.9607, \quad MSW = \frac{SSW}{40-4} = 0.1278.$$

The ANOVA table is as follows.

Source	df	\mathbf{SS}	\mathbf{MS}
Between groups	3	2.882	0.9607
Within groups	36	4.600	0.1278
Total	39	7.482	

- (b) H_0 : $\mu_1 = \mu_2 = \mu_3 = \mu_4$ v.s. H_A : The μ_i 's are not all equal.
- (c) The test statistic is

$$T = \frac{\text{MSB}}{\text{MSW}} = \frac{0.9607}{0.1278} = 7.5172.$$

The null distribution of the test statistic is $F_{3,36}$. The critical value for $\alpha = 0.01$ is thus $F_{3,30}(0.01) = 4.51$.

- (d) From F Table with numerator df = 3 and denominator df = 30, we find that $P(F_{3,30} > 7.05) = 0.001$ and $P(F_{3,30} > 9.99) = 0.0001$. The range of *p*-value is thus (0.0001, 0.001).
- (e) Since the *p*-value $< \alpha = 0.01$, we reject the null at the 0.01 level of significance.
- (f) We conclude at the 0.01 level of significance that at least two of the average GPA's of the four sororities are different.
- (g) We could falsely reject the null and thus possibly made a Type I error.
- (h) To construct family-wise 99% confidence intervals for $\mu_2 \mu_1, \mu_2 \mu_3$, and $\mu_2 \mu_4$. The individual coverage probability for each confidence interval is $1 \alpha/3$ where $\alpha = 0.01$. The $1 \alpha/3$ confidence interval for $\mu_i \mu_j$ is given by

$$(\bar{Y}_i - \bar{Y}_j) \pm t_{n-I}(\alpha/(2 \times 3)) \times \operatorname{SE}_{\bar{Y}_i - \bar{Y}_j}$$

Running qt (p = 1 - 0.01 / (2 * 3), df = 40 - 4) in R, we know that $t_{36}(0.01/6) = 3.143858$. The family-wise 99% confidence intervals for $\mu_2 - \mu_1, \mu_2 - \mu_3$, and $\mu_2 - \mu_4$ are thus as follows.

$$(\bar{Y}_2 - \bar{Y}_1) \pm t_{36}(0.01/6) \times \sqrt{\text{MSW} \times \left(\frac{1}{n_2} + \frac{1}{n_1}\right)}$$

= (3.57 - 3.22) ± 3.143858 × $\sqrt{0.1278 \times \left(\frac{1}{10} + \frac{1}{10}\right)}$
= (-0.1526, 0.8526),

$$(\bar{Y}_2 - \bar{Y}_3) \pm t_{36}(0.01/6) \times \sqrt{\text{MSW} \times \left(\frac{1}{n_2} + \frac{1}{n_3}\right)}$$

= $(3.57 - 2.87) \pm 3.143858 \times \sqrt{0.1278 \times \left(\frac{1}{10} + \frac{1}{10}\right)}$
= $(-0.1974, 1.2026),$

$$(\bar{Y}_2 - \bar{Y}_4) \pm t_{36}(0.01/6) \times \sqrt{\text{MSW} \times \left(\frac{1}{n_2} + \frac{1}{n_4}\right)}$$

= $(3.57 - 2.98) \pm 3.143858 \times \sqrt{0.1278 \times \left(\frac{1}{10} + \frac{1}{10}\right)}$
= $(0.0874, 1.0926).$

- (i) The confidence interval for $\mu_2 \mu_4$ suggest a significant difference in the means as it does not include zero.
- 5. (a) Here we have $I = 3, n = \sum_{i=1}^{3} n_i = 21$, and

$$\bar{Y} = \frac{\sum_{i=1}^{3} n_i \bar{Y}_i}{n} \\ = \frac{7 \times 2.57 + 7 \times 3.71 + 7 \times 4.29}{21} \\ = 3.5233.$$

It follows that

$$SSB = \sum_{i=1}^{3} n_i (\bar{Y}_i - \bar{Y})^2$$

= 7 × (2.57 - 3.5233)² + 7 × (3.71 - 3.5233)² + 7 × (4.29 - 3.5233)²
= 10.72027

and

SSW =
$$\sum_{i=1}^{3} (n_i - 1)s_i^2$$

= $(7 - 1) \times (0.98^2 + 1.11^2 + 1.38^2)$
= 24.5814.

Therefore,

$$SSTO = SSB + SSW = 10.7203 + 24.5814 = 35.3017.$$

If follows that

$$MSB = \frac{SSB}{3-1} = 5.3601, \quad MSW = \frac{SSW}{21-3} = 1.3656.$$

The ANOVA table is as follows.

Source	df	\mathbf{SS}	\mathbf{MS}
Between groups	2	10.7202	5.3601
Within groups	18	24.5814	1.3656
Total	20	35.3017	

(b) $H_0: \mu_1 = \mu_2 = \mu_3$ v.s. $H_A:$ The μ_i 's are not all equal.

(c) The test statistic is

$$T = \frac{\text{MSB}}{\text{MSW}} = \frac{5.3601}{1.3656} = 3.9251.$$

The null distribution of the test statistic is $F_{2,18}$. The critical value for $\alpha = 0.01$ is thus $F_{2,18}(0.01) = 6.01$.

- (d) From F Table with numerator df = 2 and denominator df = 18, we find that $P(F_{2,18} > 3.55) = 0.05$ and $P(F_{2,18} > 4.90) = 0.02$. The range of *p*-value is thus (0.02, 0.05).
- (e) Since the *p*-value > $\alpha = 0.01$, we fail to reject the null at the 0.01 level of significance.
- (f) There is insufficient evidence to conclude that there is any difference among the average number of calls in the morning, afternoon and night shifts, at the 0.01 level of significance.
- (g) We could falsely fail to reject the null and thus possibly made a Type II error.
- (h) To construct family-wise 99% confidence intervals for $\mu_1 \mu_2$, $\mu_1 \mu_3$, and $\mu_2 \mu_3$. The individual coverage probability for each confidence interval is $1 \alpha/3$ where $\alpha = 0.01$. The $1 \alpha/3$ confidence interval for $\mu_i \mu_j$ is given by

$$(\bar{Y}_i - \bar{Y}_j) \pm t_{n-I}(\alpha/(2 \times 3)) \times \operatorname{SE}_{\bar{Y}_i - \bar{Y}_j}$$

Running qt (p = 1 - 0.01 / (2 * 3), df = 21 - 3) in R, we know that $t_{18}(0.01/6) = 3.380362$. The family-wise 99% confidence intervals for $\mu_1 - \mu_2, \mu_1 - \mu_3$, and $\mu_2 - \mu_3$ are thus as follows.

$$(\bar{Y}_1 - \bar{Y}_2) \pm t_{18}(0.01/6) \times \sqrt{\text{MSW} \times \left(\frac{1}{n_1} + \frac{1}{n_2}\right)}$$

= (2.57 - 3.71) ± 3.380362 × $\sqrt{1.3656 \times \left(\frac{1}{7} + \frac{1}{7}\right)}$
= (-3.2515, 0.9715),

$$(\bar{Y}_1 - \bar{Y}_3) \pm t_{18}(0.01/6) \times \sqrt{\text{MSW} \times \left(\frac{1}{n_1} + \frac{1}{n_3}\right)}$$
$$= (2.57 - 4.29) \pm 3.380362 \times \sqrt{1.3656 \times \left(\frac{1}{7} + \frac{1}{7}\right)}$$
$$= (-3.8315, 0.3915),$$

$$(\bar{Y}_2 - \bar{Y}_3) \pm t_{18}(0.01/6) \times \sqrt{\text{MSW} \times \left(\frac{1}{n_2} + \frac{1}{n_3}\right)}$$

= $(3.71 - 4.29) \pm 3.380362 \times \sqrt{1.3656 \times \left(\frac{1}{7} + \frac{1}{7}\right)}$
= $(-2.6915, 1.5315).$

- (i) Yes, the confidence intervals are consistent with the conclusion in (f) since all of them include zero.
- 6. (a) The response variable is the peak flow, and the explanatory variable is the height.
 - (b) The slope is

$$b_1 = r \frac{s_Y}{s_X} = 0.32725 \times \frac{117.9952}{8.5591} = 4.5114.$$

The intercept is

$$b_0 = \bar{Y} - b_1 \bar{X} = 660 - 4.5114 \times 180.4118 = -153.9098$$

(c) Slope: When height of male increases by 1 cm, we expect peak flow to increase by 4.5114 liters/min, on average.

Intercept: No practical meaning because height can never equal to 0.

(d) The prediction based on the fitted regression line is

$$\hat{Y} = -153.9098 + 4.5114 \times 174 = 631.0738.$$

(e) Their average difference in peak flow would be

$$10 \times b_1 = 45.114.$$

- 7. (a) $H_0: \rho = 0$ v.s. $H_A: \rho \neq 0$ or $H_0: \beta_1 = 0$ v.s. $H_A: \beta_1 \neq 0$.
 - (b) The test statistic is

$$T = r\sqrt{\frac{n-2}{1-r^2}} = 0.32725 \times \sqrt{\frac{17-2}{1-0.32725^2}} = 1.3413.$$

The null distribution for the test statistic is t_{15} . The critical value for $\alpha = 0.05$ is thus $t_{15}(0.05/2) = 2.131$.

- (c) From t Table with df = 15, we find that $P(t_{15} > 1.341) = 0.10$ and $P(t_{15} > 1.753) = 0.05$. The range of p-value is thus (0.10, 0.20).
- (d) Since the *p*-value > $\alpha = 0.05$, we fail to reject the null at the 0.05 level of significance.
- (e) There is insufficient evidence to conclude that that peak flow is linearly related with height.
- (f) The residual standard deviation is

$$s_e = \sqrt{\frac{\text{SSE}}{n-2}} = \sqrt{\frac{198909.3}{17-2}} = 115.1548$$

The standard error for b_1 is

$$SE_{b_1} = \frac{s_e}{s_X \sqrt{n-1}} = \frac{115.1548}{8.5591 \times \sqrt{17-1}} = 3.3635.$$

The 95% confidence interval for the slope is thus

$$4.5114 \pm 2.131 \times 3.3635$$

or (-2.6562, 11.6790). The confidence interval is consistent with the conclusion in (e) since it includes zero.

(g) We are 95% confident that the when the height increases by 1 cm, we expect the peak flow to increase by between -2.6562 liters/min and 11.6790 liters/min on average.

Problem 8

```
blood = read.csv("blood.csv", header = T)
head(blood)
##
     Type Disease
## 1
        0
              yes
## 2
        0
              yes
## 3
        0
              yes
## 4
        0
              yes
## 5
        0
              yes
## 6
        0
              yes
 (a) - (b)
# create the observaion table, group by Type and Disease
0 = xtabs(~ Type + Disease, data = blood)
test_result = chisq.test(0)
## Warning in chisq.test(0): Chi-squared approximation may be incorrect
test_result
##
##
    Pearson's Chi-squared test
##
## data: 0
## X-squared = 10.654, df = 3, p-value = 0.01375
```

According to the R result, the test-statistic is T = 10.654, and *p*-value is 0.01375.

(c) Since *p*-value > $\alpha = 0.01$, we fail to reject the null at the 0.01 level of significance. We conclude that blood type and whether to develop a disease are independent.

test_result\$observed

Disease ## Type no yes ## А 12 15 ## AB 7 2 В 8 17 ## 9 30 ## 0

test_result\$expected

Disease ## Type no yes ## 9.72 17.28 А 3.24 5.76 ## AB 9.00 16.00 ## В 14.04 24.96 ## 0

- (d) The observed frequency for blood type A is 15, while the expected frequency is 17.28. Blood type A is thus less likely to have the disease than what we expected if the null was true.
- (e) The observed frequency for blood type A is 30, while the expected frequency is 24.96. Blood type A is thus more likely to have the disease than what we expected if the null was true.

(f)

(test_result\$observed - test_result\$expected)^2/test_result\$expected

 ##
 Disease

 ##
 Type
 no
 yes

 ##
 A
 0.5348148
 0.3008333

 ##
 AB
 4.3634568
 2.4544444

 ##
 B
 0.1111111
 0.0625000

 ##
 0
 1.8092308
 1.0176923

The group blood type "AB" and no disease contributes most to the test statistic.

Problem 9

```
IQ = read.csv("IQ.csv")
head(IQ)
     group iq
##
## 1
          A 44
## 2
          A 40
         A 44
## 3
## 4
         A 39
## 5
         A 25
         A 37
## 6
 (a)
anova = aov(iq ~ group, data = IQ)
summary(anova)
##
                Df Sum Sq Mean Sq F value Pr(>F)
                      1529
                             764.7
## group
                 2
                                      20.02 7.84e-07 ***
## Residuals
                42
                      1604
                               38.2
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
 (b) The test statistic is 20.02 and the p-value is 7.84 \times 10^{-7}.
 (c) Since p-value < \alpha = 0.05, we fail to reject the null at the 0.05 level of significance.
 (d) There is significant difference for the mean IQ of students among the three majors.
 (e)
library(ggplot2)
ggplot(IQ,
       aes(sample = iq)) +
  stat_qq() +
  stat_qq_line() +
  labs(x = "Theoretical Quantiles",
       y = "Sample Quantiles",
       title = "Normal Quantile Plot") +
  theme(plot.title = element_text(hjust = 0.5))
```



This data do not appear to be approximately normally distributed.

(f) library(asbio)

```
## Loading required package: tcltk
bonfCI(y = IQ$iq, x = factor(IQ$group), conf.level = 0.95)
##
## 95% Bonferroni confidence intervals
##
##
                Diff
                         Lower
                                  Upper Decision Adj. p-value
           -0.06667 -5.69471 5.56138
                                           FTR HO
## muA-muB
                                                             1
## muA-muC
               -12.4 -18.02805 -6.77195 Reject HO
                                                         6e-06
## muB-muC -12.33333 -17.96138 -6.70529 Reject H0
                                                         7e-06
```

(g) The confidence intervals for $\mu_A - \mu_C$ and $\mu_B - \mu_C$ suggest a significant difference in the means.

Problem 10

```
fitness = read.csv("fitness.csv")
head(fitness)
```

Tread Run
1 7.5 43.5
2 7.8 45.2
3 7.9 44.9

```
## 4
      8.1 41.1
## 5
      8.3 43.8
## 6
      8.7 44.4
 (a)
reg = lm(Run ~ Tread, data = fitness)
summary(reg)
##
## Call:
## lm(formula = Run ~ Tread, data = fitness)
##
## Residuals:
##
      Min
                1Q Median
                                ЗQ
                                       Max
## -2.9440 -1.5788 0.1860 0.7863 4.5603
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 59.9211
                            3.1166 19.226 1.90e-13 ***
## Tread
                -1.9601
                            0.3164 -6.194 7.59e-06 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.921 on 18 degrees of freedom
## Multiple R-squared: 0.6807, Adjusted R-squared: 0.6629
## F-statistic: 38.37 on 1 and 18 DF, p-value: 7.589e-06
```

The slope and intercept of the fitted regression line are -1.9601 and 59.9211.

(b) confint(reg, 'Tread', level = 0.95) ## 2.5 % 97.5 % ## Tread -2.624957 -1.295313

(c) From the summary table, we find that $s_e = 1.921$.

(d) From the summary table, we find that $r^2 = 0.6807$.

(e) Yes, the interval suggests a significant linear relationship.