

# Homework 3 - P8130 Biostatistics Method I

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## Problems 1–3 use `birthwt` from `{MASS}`

We first load and lightly prepare the data referenced by the problems.

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### Problem 1 (10 points)

*In this data set, we have a variable (`smoke`) indicating the smoking status of the mothers during pregnancy. Some doctors believe that smoking status is related to weight. Using the columns `smoke` and `lwt`, test this claim. (Note: A value of 1 indicates the mother is in the “smoking” group.)*

a) Test for the equality of variances between the two groups using a 5% significance level.

We assume the weights of smoker and non-smokers are two independent samples from approximately normal populations.

```
smoke0 <- bw$lwt[bw$smoke == "No smoke"]
smoke1 <- bw$lwt[bw$smoke == "Smoke"]
f_test <- var.test(smoke1, smoke0, alternative = "two.sided") # F test for  $\sigma_1^2 = \sigma_0^2$ 
f_test

##
## F test to compare two variances
##
## data:  smoke1 and smoke0
## F = 1.4126, num df = 73, denom df = 114, p-value = 0.09744
## alternative hypothesis: true ratio of variances is not equal to 1
## 95 percent confidence interval:
##  0.9388406 2.1671700
## sample estimates:
## ratio of variances
##           1.412636
```

From the result, we can see  $p = 0.09744 > 0.05$ , so we fail to reject  $H_0$  at the 5% significance level.

b) Given your answer from part (a), what kind of hypothesis test will you perform to compare means?

Since the variances are not significantly different, we will use the pooled two-sample t test.

c) Conduct your chosen hypothesis test from part (b) at the 10% significance level. What is your decision regarding the null? Interpret this result in the context of the problem.

```
equal_var <- f_test$p.value >= 0.05
tt <- t.test(lwt ~ smoke, data = bw, var.equal = equal_var, alternative = "two.sided", conf.level = 0.10)
tt
```

```
##
## Two Sample t-test
##
## data: lwt by smoke
## t = 0.60473, df = 187, p-value = 0.5461
## alternative hypothesis: true difference in means between group No smoke and group Smoke is not equal to 0
## 90 percent confidence interval:
## -4.785414 10.306448
## sample estimates:
## mean in group No smoke mean in group Smoke
## 130.8957 128.1351
```

$H_0 : \mu_{\text{Smoke}} = \mu_{\text{No smoke}}$  vs.  $H_1 : \mu_{\text{Smoke}} \neq \mu_{\text{No smoke}}$

Since the  $p\text{-value} = 0.5461 > 10\%$ , we failed to reject  $H_0$ . There is not enough evidence that smoking has an impact on weight.

d) Construct a 95% confidence interval for the difference in means and provide a proper interpretation.

```
tt95 <- t.test(lwt ~ smoke, data = bw, var.equal = equal_var, conf.level = 0.95)
tt95
```

```
##
## Two Sample t-test
##
## data: lwt by smoke
## t = 0.60473, df = 187, p-value = 0.5461
## alternative hypothesis: true difference in means between group No smoke and group Smoke is not equal to 0
## 95 percent confidence interval:
## -6.244749 11.765783
## sample estimates:
## mean in group No smoke mean in group Smoke
## 130.8957 128.1351
```

The 95% CI for  $\mu_{\text{Smoke}} - \mu_{\text{No smoke}}$  is  $(-6.24, 11.77)$ .

## Problem 2 (10 points)

Researchers are interested in exploring whether rates of uterine irritability in the group of pregnant women who smoke vs those who do not smoke? (Use columns `ui` and `smoke`.)

a) Construct a 95% confidence interval for the rate of uterine irritability among pregnant women who smoke and interpret the interval.

```
tab <- table(bw$smoke, bw$ui)
tab
```

```
##
##           No Yes
## No smoke 100  15
## Smoke    61  13
```

```
x_smoke <- tab["Smoke", "Yes"]
n_smoke <- sum(tab["Smoke",])
p_hat   <- x_smoke / n_smoke
ci_smoke <- prop.test(x_smoke, n_smoke, conf.level = 0.95, correct = TRUE)
ci_smoke
```

```
##
## 1-sample proportions test with continuity correction
##
## data:  x_smoke out of n_smoke, null probability 0.5
## X-squared = 29.851, df = 1, p-value = 4.665e-08
## alternative hypothesis: true p is not equal to 0.5
## 95 percent confidence interval:
##  0.1004347 0.2853177
## sample estimates:
##           p
## 0.1756757
```

The 95% confidence interval for the rate of uterine irritability among women who smoke is (0.100, 0.285).

b) Construct a 95% confidence interval for the difference in the rate between those who smoke and those who do not. Interpret the interval.

```
x_nosmoke <- tab["No smoke", "Yes"]
n_nosmoke <- sum(tab["No smoke",])
diff_ci <- prop.test(x = c(x_smoke, x_nosmoke),
                     n = c(n_smoke, n_nosmoke),
                     conf.level = 0.95,
                     correct = TRUE) # gives CI for p1 - p2
diff_ci
```

```
##
## 2-sample test for equality of proportions with continuity correction
##
## data:  c(x_smoke, x_nosmoke) out of c(n_smoke, n_nosmoke)
## X-squared = 0.41576, df = 1, p-value = 0.5191
## alternative hypothesis: two.sided
## 95 percent confidence interval:
## -0.07219452  0.16267631
## sample estimates:
##      prop 1      prop 2
## 0.1756757 0.1304348
```

The 95% confidence interval for the differences in the rate of uterine irritability between women who smoke and those who do not is (-0.072, 0.163).

c) Conduct a hypothesis test for a difference in proportions at the  $\alpha = 0.01$  level. What can we conclude?

```
# Two-sample test for equality of proportions (with continuity correction)
prop_test <- prop.test(x = c(x_smoke, x_nosmoke),
                      n = c(n_smoke, n_nosmoke),
                      alternative = "two.sided",
                      conf.level = 0.99,
                      correct = TRUE)

prop_test
```

```
##
## 2-sample test for equality of proportions with continuity correction
##
## data:  c(x_smoke, x_nosmoke) out of c(n_smoke, n_nosmoke)
## X-squared = 0.41576, df = 1, p-value = 0.5191
## alternative hypothesis: two.sided
## 99 percent confidence interval:
## -0.1056061  0.1960879
## sample estimates:
##      prop 1      prop 2
## 0.1756757 0.1304348
```

$H_0 : p_{\text{Smoke}} = p_{\text{No smoke}}$  vs.  $H_1 : p_{\text{Smoke}} \neq p_{\text{No smoke}}$

Since the p-value = 0.5191 > 0.01, there is not enough evidence to say smoking has an impact on the rate of uterine irritability.

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### Problem 3 (10 points)

*Is race related to birth weight? (Use columns race and bwt.)*

a) What test would be most appropriate to answer this question?

Since the dataset *bwt* across 3 groups of race, the standard choice is a **one-way ANOVA** to compare group means.

b) What assumptions are we making by using this test? Are all assumptions met?

ANOVA assumes - (i) the 3 set of data are independent to each other - (ii) data within each group are normally distributed - (iii) the variances of each group are roughly the same

c) Conduct the test at the 5% significance level and interpret your results. Be sure to write the hypotheses you are testing.

```
fit_aov <- aov(bwt ~ race, data = bw)
fit_aov
```

```
## Call:
## aov(formula = bwt ~ race, data = bw)
##
## Terms:
##              race Residuals
## Sum of Squares  5015725  94953931
## Deg. of Freedom      2      186
##
## Residual standard error: 714.4963
## Estimated effects may be unbalanced
```

```
anova_res <- summary(fit_aov)
anova_res
```

```
##              Df    Sum Sq Mean Sq F value    Pr(>F)
## race           2   5015725  2507863    4.913 0.00834 **
## Residuals    186  94953931   510505
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

$H_0 : \mu_{\text{White}} = \mu_{\text{Black}} = \mu_{\text{Other}}$  vs.  $H_1 : \text{at least one group mean differs.}$

From the result, the p-value is  $0.00834 < 5\%$ , we reject  $H_0$ . So, at least one race has a different mean of birth weight.

d) Perform multiple comparisons - which races can you claim are different from which? Interpret your results.

Use Tukey's HSD to control the family-wise error rate for all pairwise mean differences.

```
tuk <- TukeyHSD(fit_aov)
tuk
```

```
## Tukey multiple comparisons of means
## 95% family-wise confidence level
##
## Fit: aov(formula = bwt ~ race, data = bw)
##
## $race
##              diff          lwr          upr          p adj
## Black-White -383.02644 -756.2363  -9.816581 0.0428037
## Other-White -297.43517 -566.1652 -28.705095 0.0260124
## Other-Black   85.59127 -304.4521 475.634630 0.8624372
```

From the result, since both adjusted p-values for Black-White and Other-White are  $< 5\%$ , people in the ‘White’ race group are more likely to have a different birth weight than other people.

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## Problem 4 (10 points)

A public-health team is evaluating a new community program intended to reduce average systolic blood pressure (SBP) among participants. The current population guideline level is  $\mu = 130$  mmHg. Past studies indicate that the population standard deviation of SBP is  $\sigma = 15$  mmHg.

- a) The investigators will use a one-sample Z test and a 5% significance level to test  $H_0: \mu = 130$  vs.  $H_a: \mu < 130$ .
- b) If the true mean after the program is 125 mmHg, calculate the power of the test for  $n = 30$ . What is the power if the true mean is 122?

$$Power = P(Z < z_\alpha + \frac{\mu_0 - \mu}{\sigma/\sqrt{n}}) = P(Z < z_\alpha + \frac{130 - 125}{15/\sqrt{30}})$$

```
power_mu <- function(mu1, mu0 = 130, sigma = 15, n = 30, z_alpha = qnorm(0.05)) {
  pnorm(z_alpha + (mu0 - mu1)/(sigma/sqrt(n)))
}
```

```
power_mu(125)
```

```
## [1] 0.5717723
```

```
power_mu(122)
```

```
## [1] 0.8990811
```

The power is 57.1% if the true mean is 125 mmHg or 89.9% if the true mean is 122 mmHg.

- c) What should the sample size  $n$  be to ensure at least 80% power if the true mean after the program is 124? What would it need to be if we needed 90% power?

$$n = \left( \frac{\sigma}{\mu_0 - \mu_1} (z_{1-\beta} - z_\alpha) \right)^2 = \left( \frac{15}{130 - 124} (z_{1-\beta} - z_\alpha) \right)^2$$

```
min_n <- function(power, sigma = 15, mu0 = 130, mu1 = 124, alpha = 0.05) {
  (sigma / (mu0 - mu1) * (qnorm(power) - qnorm(alpha)))^2
}
```

```
min_n(0.8)
```

```
## [1] 38.64098
```

```
min_n(0.9)
```

```
## [1] 53.52405
```

The required sample size n is at least 39 to have a power of 80% or 54 to have a power of 90%.