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- 1) a) matched pairs
  - b) independent samples
  - c) independent samples
  - d) matched pairs

2)

a) There is not enough evidence to conclude that the thickness of eggs from owls exposed to the chemical is less than owl whos eggs have not been exposed to the chemical.

Welch Two Sample t-test

- b) Probability of a Type II error
- $[1] \ \ 0.0500000000 \ \ 0.0207474914 \ \ 0.0074988945 \ \ 0.0023529776 \ \ 0.0006392649$ 
  - c) n = 11, so m = 33

One-sample t test power calculation

```
n = 10.66853
d = 0.8207294
sig.level = 0.05
power = 0.8
alternative = greater
```

d) There is not sufficient evidence to suggest that owl eggs from owls exposed to the chemical are less than owl eggs from owls who have not been exposed to the chemical.

Wilcoxon rank sum test with continuity correction

```
data: dt$exposed and dt$not_exposed
W = 40, p-value = 0.2347
alternative hypothesis: true location shift is less than 0
```

e) There is not significant evidence to suggest that the variability of exposed eggs is greater than the variability of unexposed eggs.

p-value: 0.8794254

f) data collected from the non exposed group does not bass the shapiro test of normality so wilcox will be a better test.

shapiro.test(dt\$exposed)

Shapiro-Wilk normality test

data: dt\$exposed
W = 0.96279, p-value = 0.8171

shapiro.test(dt\$not\_exposed)

Shapiro-Wilk normality test

data: dt\$not\_exposed
W = 0.81803, p-value = 0.02399

3) With a significance level of .05, both tests fail to reject the null hypothesis that the vitaminB group is larger than the placebo group. The t-test is more reliable in this case because both samples pass the shapiro test of normality.

Paired t-test

Wilcoxon signed rank test with continuity correction

data: dtvitaminB and dtplacebo V = 47, p-value = 0.1144 alternative hypothesis: true location shift is greater than 0

4)

- a)  $H_o: p_1 = p_2$ ,  $H_1: p_1 \neq p_2$ , where  $p_1$  and  $p_2$  are the probabilities of normal and diabetic patients having low excretions.
- b) There is substantial evidience to suggest that the two probabilities are not the same so we reject  $H_o$

```
cat('p-value:', phyper(4, 12, 12, 14))
```

p-value: 0.01803742

5)

- a)  $H_o: p_1 = p_2, H_1: p_1 \neq p_2$ , where  $p_1$  and  $p_2$  are the probabilities of Drug 1 and Drug 2 successfully anesthetizing membranes.
- b) There is sufficient evidence to conclude that the probabilities of the 2 drugs are different.

```
cat('p-value:', phyper(18, 45, 45, 46))
```

p-value: 0.02856317

6) There is not substantial evidence to conclude that the phenotypes follow the proportion 9:3:3:1.

```
Oi Ei TS
1 926 906.1875 0.4331721
```

2 293 302.0625 0.2718938

3 288 302.0625 0.6546788

4 104 100.6875 0.1089773

p-value: 0.3104921

7)

a) Based on the Shapiro Wilks test of normality the data from Lab2 appears normal, while the data from Lab1 does not.

Shapiro-Wilk normality test

```
data: dt$Lab.1
W = 0.85222, p-value = 0.005809
```

Shapiro-Wilk normality test

```
data: dt$Lab.2
W = 0.9357, p-value = 0.1987
```

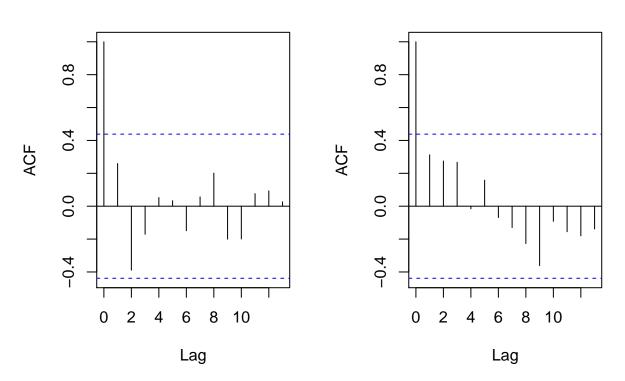
b) Based on the F test, the two variances appear to be different

F test to compare two variances

c) There is no significant correlation between day to tests, all correlation are within the acceptable "noise" range.

## **Autocorrelation Plot Lab 1**

## **Autocorrelation Plot Lab 2**



d) Based on wilcox test, the two averages appear to be different. The wilcox test is a better approximation than the t test because one of the labs data appears not to be from a normal distribution.

Wilcoxon rank sum test with continuity correction

data: dt\$Lab.1 and dt\$Lab.2 W = 331.5, p-value = 0.0003827

alternative hypothesis: true location shift is not equal to 0

e)

Lab.1 95% Confidence Interval: 3.891581 4.172419

Lab.1 95% Confidence Interval: 3.929172 4.134828

8) There is substantial evidence that the relationship between carrier status and size are not independent. Futhermore, measures of assocation show that they have very little association.

Oi

	Carrier	NonCarrier
Norm	19	497
Large	29	560
V. Large	24	269

Εi

	Carrier	NonCarrier
Norm	26.57511	489.4249
Large	30.33476	558.6652
V. Large	15 09013	277 9099

TS

Carrier NonCarrier
Norm 2.15924812 0.11724424
Large 0.05873112 0.00318902
V. Large 5.26077722 0.28565306

p-value: 0.01940118

Association
Phi 0.07510052
Contingency 0.07488962
Cramer's V 0.07507367

9)

a) Although the p-value is very close to .05, there is not substantial evidence to conclude that the proportions of white and black death penally proportions are different.

Oi

DP No.DP White 30 184 Black 6 106

Εi

DP No.DP White 23.6319 190.3681 Black 12.3681 99.6319

TS

DP No.DP White 1.716014 0.2130224 Black 3.278812 0.4070250

p-value: 0.0603595

Association
Phi 0.1312385
Contingency 0.1301227
Cramer's V 0.1310377

b) There is not enough evidence to conclude that there is a difference between white and black death sentence proportions.

0i

DP No.DP White 19 141 Black 17 149

Εi

DP No.DP White 17.66871 142.3313 Black 18.33129 147.6687

TS

DP No.DP White 0.10030888 0.01245214 Black 0.09668326 0.01200206

p-value: 0.8951865

Association
Phi 0.02606306
Contingency 0.02605421
Cramer's V 0.02602318

c) There is not evidence to support that the ratio of dealth penalty sentences for whites and blacks are any different.

White Victim Black Victim log OR -0.3850145 -Inf Weight 0.1630864 1.498052

OR Stat df pvalue 0.6352968 0.1818143 1.0000000 0.6698187

d) All 3 tests were unable to support the evidence that the proportions and ratios between white and black death penalty sentences were any different, although when testing victims race only it was very close to meeting the alpha conditions and with more data it might prove to be significant.