Chapter 9
Paired Data

Comparison of Paired Samples

● In chapter 7 we discussed how to compare two independent samples
● In chapter 9 we discuss how to compare two samples that are paired
  - In other words the two samples are not independent, $Y_1$ and $Y_2$ are linked in some way, usually by a direct relationship
  - For example, measure the weight of subjects before and after a six month diet

Paired data

● The mean of the differences is calculated just like the one sample mean we calculated in chapter 2
  \[ \bar{d} = \frac{\sum d}{n_d} = \bar{y}_1 - \bar{y}_2 \]
  - it also happens to be equal to the difference in the sample means – this is similar to the t test
  - This sample mean differences is an estimate of the population mean difference $\mu_d = \mu_1 - \mu_2$

Paired data

● Because we are focusing on the differences, we can use the same reasoning as we did for a single sample in chapter 6 to calculate the standard error
  - aka. the standard deviation of the sampling distribution of $\bar{d}$
  - Recall: $SE = \frac{s}{\sqrt{n}}$
  - Using similar logic: $SE_d = \frac{s_d}{\sqrt{n_d}}$
    - where $s_d$ is the standard deviation of the differences and $n_d$ is the sample size of the differences
Paired data

Example: Suppose we measure the thickness of plaque (mm) in the carotid artery of 10 randomly selected patients with mild atherosclerotic disease. Two measurements are taken, thickness before treatment with Vitamin E (baseline) and after two years of taking Vitamin E daily.

<table>
<thead>
<tr>
<th>Subject</th>
<th>Before</th>
<th>After</th>
<th>Difference</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>0.66</td>
<td>0.60</td>
<td>0.06</td>
</tr>
<tr>
<td>2</td>
<td>0.72</td>
<td>0.65</td>
<td>0.07</td>
</tr>
<tr>
<td>3</td>
<td>0.65</td>
<td>0.79</td>
<td>0.14</td>
</tr>
<tr>
<td>4</td>
<td>0.62</td>
<td>0.63</td>
<td>0.01</td>
</tr>
<tr>
<td>5</td>
<td>0.59</td>
<td>0.54</td>
<td>0.05</td>
</tr>
<tr>
<td>6</td>
<td>0.63</td>
<td>0.56</td>
<td>0.07</td>
</tr>
<tr>
<td>7</td>
<td>0.64</td>
<td>0.62</td>
<td>0.02</td>
</tr>
<tr>
<td>8</td>
<td>0.70</td>
<td>0.67</td>
<td>0.03</td>
</tr>
<tr>
<td>9</td>
<td>0.73</td>
<td>0.66</td>
<td>0.07</td>
</tr>
<tr>
<td>10</td>
<td>0.68</td>
<td>0.64</td>
<td>0.04</td>
</tr>
</tbody>
</table>

What makes this paired data rather than independent data?

Why would we want to use pairing in this example?

Mean 0.682 0.637 0.045
sd 0.0742 0.0709 0.0264

Paired CI for \( \mu_d \)

Calculate the mean of the differences and the standard error for that estimate

\[
\bar{d} = 0.045 \\
sd = 0.0264 \\
SE_d = \frac{sd}{\sqrt{n_d}} = \frac{0.0264}{\sqrt{10}} = 0.00833
\]

A 100(1 - \( \alpha \))% confidence interval for \( \mu_d \)

\[
\bar{d} \pm t(df)_{0.05}(SE_d)
\]

where df = \( n_d - 1 \)

Very similar to the one sample confidence interval we learned in section 6.3, but this time we are concentrating on a difference column rather than a single sample

Paired CI for \( \mu_d \)

Example: Vitamin E (cont')

Calculate a 90% confidence interval for the true mean difference in plaque thickness before and after treatment with Vitamin E

\[
\bar{d} \pm t(df)_{0.5}(SE_d)
\]

\[
= 0.045 \pm t(9)_{0.05}(0.00833)
\]

CONCLUSION: We are highly confident, at the 0.10 level, that the true mean difference in plaque thickness before and after treatment with Vitamin E is between 0.03 mm and 0.06 mm.

Great, what does this really mean?

Does the zero rule work on this one?
Paired t test

- Of course there is also a hypothesis test for paired data
- #1 Hypotheses:
  - $H_0: \mu_d = 0$
  - $H_a: \mu_d \neq 0$ or $H_a: \mu_d < 0$ or $H_a: \mu_d > 0$
- #2 test statistic
  - Where $df = n_2 - 1$
- #3 p-value and #4 conclusion similar idea to that of the independent t test

Example: Vitamin E (cont’)

Do the data provide enough evidence to indicate that there is a difference in plaque before and after treatment with vitamin E for two years? Test using $\alpha = 0.10$

$H_0: \mu_d = 0$ (thickness in plaque is the same before and after treatment with Vitamin E)

$H_a: \mu_d \neq 0$ (thickness in plaque after treatment is different than before treatment with Vitamin E)

$df = 10 - 1 = 9$

$p < 2(0.0005) = 0.001$, so we reject $H_0$.

CONCLUSION: These data show that the true mean thickness of plaque after two years of treatment with Vitamin E is statistically significantly different than before the treatment ($p < 0.001$).

In other words, vitamin E appears to be effective in changing carotid artery plaque after treatment.

May have been better to conduct this as an upper-tailed test because we would hope that vitamin E will reduce clogging.

However, researchers need to make this decision before analyzing data.

Results of Ignoring Pairing

- Suppose we accidentally analyzed the groups independently (like an independent t-test) rather than a paired test?
  - keep in mind this would be an incorrect way of analyzing the data
  - How would this change our results?

Results of Ignoring Pairing

Example Vitamin E (con’t)

Calculate the test statistic and p-value as if this were an independent t test

$df = 17$

$p < 2(0.05) < p < 2(0.1)$

Fail To Reject $H_0$
Results of Ignoring Pairing

- What happens to a CI?
  - Calculate a 90% confidence interval for \( \mu_1 - \mu_2 \)
  
\[
\bar{y}_1 - \bar{y}_2 \pm t(\frac{df}{2})(SE_{\bar{y}_1-\bar{y}_2})
\]
  
\[
= (0.682 - 0.637) \pm t(17)(0.0325)
\]
  
\[
= 0.045 \pm (1.740)(0.0325)
\]
  
\[
= (-0.0116, 0.1016)
\]

- How does the significance of this interval compare to the paired 90% CI (0.03 mm and 0.06 mm)?

- Why is this happening?

- Is there anything better about the independent CI? Is it worth it in this situation?

Paired T-Test and CI: Before, After

- Paired T for Before - After

<table>
<thead>
<tr>
<th>N</th>
<th>Mean</th>
<th>StDev</th>
<th>SE Mean</th>
</tr>
</thead>
<tbody>
<tr>
<td>10</td>
<td>0.682</td>
<td>0.074</td>
<td>0.023</td>
</tr>
<tr>
<td>After</td>
<td>0.637</td>
<td>0.070</td>
<td>0.023</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Difference</th>
<th>Mean</th>
<th>StDev</th>
<th>SE Mean</th>
</tr>
</thead>
<tbody>
<tr>
<td>10</td>
<td>0.045</td>
<td>0.026</td>
<td>0.008</td>
</tr>
</tbody>
</table>

- T-Test of mean difference = 0 (vs not = 0):
  
\[
T-Value = 5.40 \quad P-Value = 0.000
\]

Results of Ignoring Pairing

- Why would the SE be smaller for correctly paired data?
  - If we look at the within each sample at the data we notice variation from one subject to the next
  - This information gets incorporated into the SE for the independent t-test via \( s_1 \) and \( s_2 \)
  - The original reason we paired was to try to control for some of this inter-subject variation
  - This inter-subject variation has no influence on the SE for the paired test because only the differences were used in the calculation.
  - The price of pairing is smaller df.
    - However, this can be compensated with a smaller SE if we had paired correctly.

Conditions for the validity of the paired t test

- Conditions we must meet for the paired t test to be valid:
  - It must be reasonable to regard the differences as a random sample from some large population
  - The population distribution of the differences must be normally distributed.
    - The methods are approximately valid if the population is approximately normal or the sample size \( n_1 \) is large.
  - These conditions are the same as the conditions we discussed in chapter 6.

The Paired Design

- Ideally in the paired design the members of a pair are relatively similar to each other
  - Common Paired Designs
    - Randomized block experiments with two units per block
    - Observational studies with individually matched controls
    - Repeated measurements on the same individual
    - Blocking by time – formed implicitly when replicate measurements are made at different times.
  - IDEA of pairing: members of a pair are similar to each other with respect to extraneous variables
The Paired Design

Example: Vitamin E (cont’)
- Same individual measurements made at different times before and after treatment (controls for within patient variation).

Example: Growing two types of bacteria cells in a petri dish replicated on 20 different days.
- These are measurements on 2 different bacteria at the same time (controls for time variation).

Purpose of Pairing

- Pairing is used to reduce bias and increase precision
  - By matching/blocking we can control variation due to extraneous variables.
- For example, if two groups are matched on age, then a comparison between the groups is free of any bias due to a difference in age distribution
- Pairing is a strategy of design, not analysis
  - Pairing needs to be carried out before the data are observed
  - It is not correct to use the observations to make pairs after the data has been collected

Paired vs. Unpaired

- If the observed variable Y is not related to factors used in pairing, the paired analysis may not be effective
  - For example, suppose we wanted to match subjects on race/ethnicity and then we compare how much ice cream men vs. women can consume in an hour
- The choice of pairing depends on practical considerations (feasibility, cost, etc…) and on precision considerations
  - If the variability between subjects is large, then pairing is preferable
  - If the experimental units are homogenous then use the independent t

The Sign Test

- The sign test is a non-parametric version of the paired t test
- We use the sign test when pairing is appropriate, but we can’t meet the normality assumption required for the t test
- The sign test is not very sophisticated and therefore quite easy to understand
- Sign test is also based on differences
  \[ d = Y_1 - Y_2 \]
  The information used by the sign test from this difference is the sign of \( d \) (+ or -)

The Sign Test - Method

- #2 Test Statistic \( B_n \):
  1. Find the sign of the differences
  2. Calculate \( N_+ \) and \( N_- \)
  3. If \( H_0 \) is non-directional, \( B_n \) is the larger of \( N_+ \) and \( N_- \)
     If \( H_0 \) is directional, \( B_n \) is the N that jives with the direction of \( H_a \):
       - If \( H_a: Y_1 < Y_2 \) then we expect a larger \( N_- \)
       - If \( H_a: Y_1 > Y_2 \) then we expect a larger \( N_+ \)

NOTE: If we have a difference of zero it is not included in \( N_+ \) or \( N_- \), therefore \( n_y \) needs to be adjusted
The Sign Test

- #3 p-value:
  Table 7 p.684
  Similar to the WMW
  Use the number of pairs with "quality information"

- #4 Conclusion:
  Similar to the Wilcoxon-Mann-Whitney Test
  Do NOT mention any parameters!

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The Sign Test (cont')

Do the data provide sufficient evidence to indicate that the first born of a set of twins is more aggressive than the second? Test using $\alpha = 0.05$.

$H_0$: The aggressiveness is the same for 1st born and 2nd born twins
$H_a$: The aggressiveness of the 1st born twin tends to be more than the 2nd born.

NOTE: Directional $H_a$ (we're expecting higher scores for the 1st born twin), this means we predict that most of the differences will be positive

$N_+ = \text{number of positive} = 7$
$N_- = \text{number of negative} = 4$
$n_d = \text{number of pairs with useful info} = 11$

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The Sign Test

$B_s = N_+ = 7$ (because of directional alternative)
$P > 0.10$, Fail to reject $H_0$

CONCLUSION: These data show that the aggressiveness of 1st born twins is not significantly greater than the 2nd born twins ($P > 0.10$).

$X \sim B(11, 0.5)$
$P(X \geq 7) = 0.2744140625$
$http://socr.stat.ucla.edu/Applets.dir/Normal_T_Chi2_F_Tables.htm$

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The Sign Test

- Hold on did we actually need to carry out a sign test? What should we have checked first?

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Practice

- Suppose $H_a$: one-tailed, $n_d = 11$
- And $B_s = 10$
- Find the appropriate p-value
  $0.005 < p < 0.01$
  Pick the smallest p-value for $B_s = 10$ and bracket
- NOTE: Distribution for the sign test is discrete, so probabilities are somewhat smaller (similar to Wilcoxon-Mann-Whitney)
Applicability of the Sign Test

- Valid in any situation where d’s are independent of each other
- Distribution-free, doesn’t depend on population distribution of the d’s
  - although if d’s are normal the t-test is more powerful
- Can be used quickly and can be applied on data that do not permit a t-test

Example: 10 randomly selected rats were chosen to see if they could be trained to escape a maze. The rats were released and timed (sec.) before and after 2 weeks of training. Do the data provide evidence to suggest that the escape time of rats is different after 2 weeks of training?

Test using $\alpha = 0.05$.

<table>
<thead>
<tr>
<th>Rat</th>
<th>Before</th>
<th>After</th>
<th>Sign of d</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>100</td>
<td>50</td>
<td>+</td>
</tr>
<tr>
<td>2</td>
<td>38</td>
<td>12</td>
<td>+</td>
</tr>
<tr>
<td>3</td>
<td>N</td>
<td>45</td>
<td>+</td>
</tr>
<tr>
<td>4</td>
<td>122</td>
<td>62</td>
<td>+</td>
</tr>
<tr>
<td>5</td>
<td>95</td>
<td>90</td>
<td>+</td>
</tr>
<tr>
<td>6</td>
<td>116</td>
<td>100</td>
<td>+</td>
</tr>
<tr>
<td>7</td>
<td>56</td>
<td>75</td>
<td>-</td>
</tr>
<tr>
<td>8</td>
<td>135</td>
<td>53</td>
<td>+</td>
</tr>
<tr>
<td>9</td>
<td>104</td>
<td>44</td>
<td>+</td>
</tr>
<tr>
<td>10</td>
<td>N</td>
<td>50</td>
<td>+</td>
</tr>
</tbody>
</table>

N denotes a rat that could not escape the maze.

H$_0$: The escape times (sec.) of rats are the same before and after training.
H$_a$: The escape times (sec.) of rats are different before and after training.

$N_+ = 9; N_- = 1; n_d = 10$

$B_0 = $larger of $N_+ or N_- = 9$

$0.01 < p < 0.05$, reject $H_0$

CONCLUSION: These data show that the escape times (sec.) of rats before training are different from the escape times after training ($0.01 < p < 0.05$).