Basic Course on R:
Hypothesis Testing and Confidence Intervals 1
Practical

Elizabeth Ribble*

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Contents

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*emcclel3@msudenver.edu

1 Baby Data

1. Read in the data "R_data_January2015.csv" with a header and row names from the first column. Assign it to the object `babydata` and allow strings be converted to factors. Attach the data to the environment.

2. What are the dimensions of `babydata`? What is the class? Answer these questions separately with two functions and then together with one function.

3. Answer the following questions pertaining to the variable `SAH`:

   (a) What are the 20% quantiles of `SAH`?

   (b) What are the mean, median, variance and standard deviation of `SAH`?

   (c) Create a stem and leaf plot of `SAH`.

   (d) Create a histogram and a horizontal boxplot of `SAH` in one graphics window where the plot of the histogram is above the boxplot.

   (e) Utilize all 3 graphs to describe the shape of the distribution of `SAH`.

   (f) Log-transform SAH (assign it to `logSAH`).

   (g) What are the 20% quantiles of `logSAH`?

   (h) What are the mean, median, variance and standard deviation of `logSAH`?

(i) Create a stem and leaf plot of `logSAH`.

(j) Create a histogram and a horizontal boxplot of `logSAH` in one graphics window where the plot of the histogram above the boxplot.

(k) Utilize all 3 graphs to describe the shape of the distribution of `logSAH`.

(l) What did the log transformation do to the values of `SAH`?

(m) Take a random sample of size 50 from `logSAH` and make a histogram. Does this distribution have a similar shape compared to that of all `logSAH` values?

(n) Take a random sample of size 50 with replacement from `logSAH` and make a histogram. Does this distribution have a similar shape compared to that of all `logSAH` values?

4. Answer the following questions pertaining to the variable `medication`:

   (a) Use a function to create frequency table of the number of mothers taking medication and not taking medication.

   (b) Calculate the percent of the mothers who are taking medication; what is the percentage?

5. Answer the following questions pertaining to the variable `educational_level`:

   (a) Create a frequency table of the number of mothers in each education level.

(b) Create a horizontal boxplot of the `SAH` values for the different levels of education and color each box a different color. Add a rug plot of the values where the ticks for each group have the same color as their corresponding box.

(c) Are `triglycerides` normally distributed (make a plot to answer this question)? If not, log-transform them. Are the log-transformed values normal?

(d) Is the average triglyceride level for highly educated mothers different from that of mothers with a low education level? Formulate a hypothesis, test it, and make a decision about whether or not you can reject the null hypothesis. Can you use a t-test (either on the raw or log-transformed data)? Why or why not (hint: how are the data distributed)?

(e) Now re-do the test and make your decision to reject/not reject the null based on the confidence interval. Challenge: instead of just looking at the output, extract the confidence interval from the test output and use logical operators to answer the question of whether the interval contains the null value.

6. Answer the following questions pertaining to the variable `Status`:

    (a) Are the average `SAH` values for the two levels of `Status` (normal brain development or intellectual disability) different? Formulate a hypothesis, test it, and make a decision about whether or not you can reject the null hyopthesis. Can you use a t-test (either on the raw or log-transformed data)? Why or why not (hint: check distributions with plots)?

    (b) What is the fold change of log-`SAH` between the 2 groups? Calculate it two ways: use the output from the previous test and also use the data itself (function `mean` plus logical operators).

    (c) Make a boxplot of the `SAH` values of the 2 groups and calculate the fold change of `SAH` between the 2 groups. Does the difference seem clinically relevant? Why or why not?