NRES 798 — Lab 5

Distributions, summary statistics

Graph the PDF f(x), CDF F(x) and quantile function $F^{-1}(p)$ for the normal, uniform, log-normal, and exponential distributions (?Distributions). Hint: curve(dnorm(x, 0, 1), from=-3, to=3). See what effect the distribution parameters have. Expressions can be conveniently plotted with curve; what is the difference between curve and plot?

Generate a large sample (say 100000 values) from a standard uniform distribution: x <- runif(100000). Examine x: do a summary, head(x), plot a histogram (hist(x)). What is the (theoretical) population mean and the mean of the 100000 values? The population variance is known to be 1/12; what about var(x)?

Think now of x as a population (a Bayesian would say that any population is a sample, or *realization*, from some superpopulation). Add another standard uniform random value to each of the 100000 that you have: x <-x + runif(100000). Think about it, each x_i is now the sum of 2 (nearly) uniform random variables (independent, if the random number generator is any good). What should the (population) mean of these sums be? (expected value of a sum of 2 RVs). And the variance? Compare with your sample. Plot a histogram.

Repeat x < -x + runif(100000), so that now you have sums of 3 variables. Find the theoretical and observed means and variances. Plot histogram.

Repeat a few more times (hint: up-arrow). What is going on? What does the distribution look like?

Do the same with an exponential instead of the uniform. Then try a Bernouilli for some p, $x \le \text{sample(1:0, 100000, replace=T, prob=c(p, 1-p))}$, or $x \le \text{rbinom(100000, 1, p)}$ (why?). Amazing, huh?

Generate your sample of linphid spider tibial spine lengths: tibia <- rnorm(50, 0.253, 0.0039) (mm, parameters from Fig. 2.6). Display, sum-

marize, "histogramize". Do plot(x). Draw a box plot (??box), compare to the numbers from summary.

Compute and compare the arithmetic mean, geometric mean, harmonic mean, and median for your measurements. Do it the long way using the definitions, and using any built-in functions that you might find. For the median you can use **sort**. And indexing ([]) if you are too lazy to count.

For the GM, check that averaging logs gives the same as using products. Find the mode for histograms with various numbers of bins (breaks parameter).

Compute the two versions (denominator n and n-1) of the variance and standard deviation, the long way from the definitions, and with built-in functions. How do they, and the sample mean, compare to the population values (see above, used to generate the sample)? How can you calculate the values with denominator n from those with n-1, or vice-versa, after you lost the data? Verify.

Define an R function RMS <- function(x) ... to compute the root mean square, the square root of the mean square (= the standard deviation using denominator n). Test it.

Define a function Skewness(x) to compute the skewness. The s in the formula is the s from equation (3.7), not that of (3.9) or (3.10). You can use mean and RMS. Do the same for the kurtosis (again, s is the RMS). Try it out on tibia. The population value for a normal is 0 in both cases.

Generate another tibia lengths sample, same as before. Hint: start typing tibia <- and press Ctrl + up-arrow (or in RStudio, look up the command in the *History* tab, upper-right). Compute the skewness and kurtosis. Repeat a few times to get a feeling of the variability of these statistics.

Let's do this better, run a simulation, e.g., for the kurtosis: First, make a vector to store the results, k <- rep(x=0, times=10000). Now the loop, for(i in 1:10000) {tibia <- rnorm(50, 0.253, 0.0039); k[i] <- Skewness(tibia)}. Now k has 10000 values of skewness in samples of size 50. Plot a histogram, calculate the standard deviation (standard error), etc.

Do plot(ecdf(tibia)), and quantile(tibia, seq(0, 1, 0.1)). Look up these functions in the *Help*.

If you still have time, do some more experimenting and exploration around the above.