## STA 410/2102, Fall 2014 — Assignment #2

Due at the start of class on November 13. Please hand it in on 8 1/2 by 11 inch paper, stapled in the upper left, with no other packaging.

This assignment is to be done by each student individually. You may discuss it in general terms with other students, but the work you hand in should be your own. In particular, you should not leave any discussion with someone else with any written notes (either paper or electronic).

In this assignment, you will use R's integrate function for univariate numerical quadrature, applying it to maximum likelihood estimation for rounded data and to Bayesian inference for a model of bivariate normal data with known mean but unknown variance and correlation.

The integrate function takes as its first argument an R function defining what to integrate. This function must accept a *vector* of values, and return a vector of results of applying the function to those values. Many functions will naturally operate in this way, but if necessary, the function will need to loop over elements in its argument in order to compute the integrand at each element value. The second and third arguments to integrate are the low and high bounds for integration. To get the result of the integration, extract the element value from the result of integrate.

First problem. For this problem, we observe n i.i.d. values from a normal distribution with unknown mean,  $\mu$ , and unknown standard deviation,  $\sigma$ , and wish to find maximum likelihood estimates for  $\mu$  and  $\sigma$ . However, our data is rounded to d decimal places (eg, we might have data like 2.3, 1.2, 3.0, -1.7 if d=1). We wish to find maximum likelihood estimates that account for this, by maximizing the probability of obtaining n data points that round to the data that is recorded. This probability is found by integrating the normal probability density function over the range of real values that round to what was recorded.

You should write an R function called  $rnd\_norm\_mle1$  that uses nlm to find the maximum likelihood estimates for  $\mu$  and  $\log(\sigma)$  (use the log of  $\sigma$  so that there is no constraint on valid values). The function should take as arguments a vector of rounded data values and the number of decimal places to which these values were rounded. Inside your  $rnd\_norm\_mle1$  function, the function you give to nlm should compute the log likelihood using integrate. (You should not try to compute the gradient or Hessian of the log likelihood.)

You should also write an R function called rnd\_norm\_mle2 that does the same thing as rnd\_norm\_mle1 except that rather than using integrate, it finds the integrals using R's pnorm function for computing the cumulative distribution function for a normal distribution. (You'll need to call pnorm twice, for each end of the integration range, and take the difference.)

You should test your functions on three datasets generated as follows:

```
set.seed(1); d1 <- 1; x1 <- round(rnorm(25,2.24,1),d1)
set.seed(2); d2 <- 1; x2 <- round(rnorm(50,2.24,0.1),d2)
set.seed(3); d3 <- 0; x3 <- round(rnorm(4000,2.24,0.8),d3)</pre>
```

Comment on how the maximum likelihood estimates of  $\mu$  and  $\sigma$  compare to simply using the sample mean and sample standard deviation of the rounded data points. Also comment on the speed using integrate versus using pnorm.

**Second problem.** For this problem, we observe n pairs of values (to high precision, so rounding is not an issue), which we will denote by  $(X_{11}, X_{12}), (X_{21}, X_{22}), \ldots, (X_{n1}, X_{n2})$ . The n pairs are i.i.d. but the two values within each pair may be dependent, having a bivariate normal distribution with means of zero, with both elements having the same unknown standard deviation,  $\sigma$ , and with an unknown correlation,  $\rho$ , between the elements in the pair. The density function for this bivariate normal distribution for a pair  $(X_{i1}, X_{i2})$  is

$$f(x_{i1}, x_{i2}) = \frac{1}{2\pi\sigma^2\sqrt{1-\rho^2}} \exp\left(-\left(x_{i1}^2 + x_{i2}^2 - 2\rho x_{i1}x_{i2}\right) / \left(2(1-\rho^2)\sigma^2\right)\right)$$

We wish to do Bayesian inference for this model, using a prior distribution for  $\sigma$  and  $\rho$  with the following density function over  $(0, \infty) \times (-1, +1)$ :

$$f(\sigma, \rho) = e^{-\sigma} / \left(4|\rho|^{1/2}\right)$$

You should write a function bvn\_likelihood that takes as arguments a  $2 \times n$  matrix of data values, X, and parameter values sigma and rho, and returns the likelihood for these parameter values given the data. You should also write a function bvn\_prior that takes as arguments parameters values sigma and rho, and returns the prior density for these parameter values.

You should then write a function bvn\_normalize to compute the normalizing constant for the posterior distribution given data X — that is, the integral of the product of the prior density and the likelihood over the whole parameter space. You should use R's integrate function to do this. Note that integrate will accept Inf (infinity) as an upper bound. It also will handle singularities in the integrand at the end-points of the interval, but not in the interior of the integral (so you may need to break up an integral into parts to handle such singularities). Since the normalizing constant is a double integral, you will need to use an inner call of integrate inside the function integrated by an outer call of integrate.

You should also write a function bvn\_posterior\_rho with arguments X and rho that computes the marginl posterior density (with correct normalization) of rho given data X. You should then write a function for plotting both the prior density and the marginal posterior density for  $\rho$ , over its range (-1, +1), given a dataset X. Since these densities are singular at zero, you should put an upper limit of 5 on the density plotted (using an option of ylim=c(0,5) for plot).

You should try out your functions on two datasets Xa and Xb generated as follows:

```
set.seed(1)
n <- 40
x1 <- rnorm(n); x2 <- rnorm(n); z <- rnorm(n)
Xa <- cbind (x1, x2); Xb <- cbind (x1+0.5*z, x2+0.5*z)</pre>
```

Discuss how the posterior distributions you see for these datasets compare to the sample correlations and to the true values of the correlation.

For graduate students in STA 2012 (undergrads may do this for bonus marks): Investigate for what size of n the functions you wrote start to fail because of floating point overflow or underflow, and suggest a way of alleviating this problem.