

## CPMA 573 — Homework #3

Before you begin: precede your R script with `pdf(file='myfile.pdf')`, and include as the last line `dev.off()`. This will funnel every plot you make into a single pdf to submit along with your code. While you are working in Rstudio, I recommend commenting out these two lines so that you can see the graphs as you go. When you are done, uncomment them and re-run your whole script to generate the file with the figures. Make sure all of your figures have informative titles (using the option `main='mytitle'` within most plotting commands e.g. `hist()` and `curve()`).

**Q1** : Recall the beta density function  $f(x|\alpha, \beta)$  given by

$$f(x|\alpha, \beta) = \frac{\Gamma(\alpha + \beta)}{\Gamma(\alpha)\Gamma(\beta)} x^{\alpha-1} (1-x)^{\beta-1},$$

for  $x \in (0, 1)$  and positive  $\alpha$  and  $\beta$ . Let  $X \sim \text{beta}(2, 5)$ . Find  $P(X > \frac{1}{2})$  by:

- Monte Carlo (MC) integration.
- Simulating 25,000  $\text{beta}(2, 5)$  realizations. Use rejection sampling and uniform proposal density  $q(x)$ .
- Using the following command to find the theoretical solution: `pbeta(0.5, shape1=2, shape2=5, lower.tail=FALSE)`. Note: without the `lower.tail=FALSE` argument, this would return  $P(X < \frac{1}{2})$

**Q2** : Let  $X$  and  $Y$  be independent standard normal random variables. Use the polar method to simulate 25,000 pairs  $(x, y)$  of independent, standard normal realizations.

- Use these realizations to estimate  $P(X > Y)$
- Use these realizations to find the value of  $k$  that satisfies  $P(\sqrt{X^2 + Y^2} < k) = \frac{1}{2}$
- Estimate  $P(-1.96 < X < 1.96)$  for  $X \sim \text{Normal}(0, 1)$  using:
  - your  $X$  draws
  - MC integration
  - the command `pnorm(1.96)-pnorm(-1.96)` (this will give you the theoretical answer!)

**Q3** : Let  $X$  and  $Y$  have jointly normal density  $f(x, y)$  given by:

$$f(x, y) = \frac{1}{2\pi\sqrt{1-\rho^2}} \cdot \exp\left\{-\frac{(x^2 + y^2 - 2\rho xy)}{(2(1-\rho^2))}\right\} .$$

Write a function that uses Gibbs sampling to produce realizations from this joint density, where  $\rho = 0.75$ . Your program should be flexible enough to accommodate a choice of

- Correlation  $\rho$
- Initial values  $x_0$ .
- Number of desired  $(x_i, y_i)$  realizations.
- Number of iterations to skip (lag) in order to avoid serial correlation (a lag of 1 should be the same as no lag)
- Number of burn-in iterations to skip

Note that the number of desired realizations, together with the burn-in and lag, should determine the number of required iterations. Use this program to complete the following:

- a) Generate 500 realizations with  $x_0 = 80$ , no burn-in and no lag. Plot these realizations and connect the points in order using the following commands:

```
plot(xvec[1:500],yvec[1:500])
arrows(xvec[1:499],yvec[1:499],xvec[2:500],yvec[2:500])
```

- b) Use `ts.plot()` to draw time series plots for both x and y. Use these two plots to choose a reasonable burn-in.
- c) Use `acf()` to look at autocorrelation plots for both your x and y draws. Use these two plots to choose a reasonable lag for generating independent realizations of  $(x_i, y_i)$ .
- d) Using the lag and burn-in from parts b) and c) draw 10,000 independent realizations of  $X$ . Plot a histogram of x, and overlay the theoretical standard normal curve to confirm that the marginal density is correct.

**Q4** : Use your Gibbs sampling function from above to generate 15,000 independent realizations each for  $\rho = \{-0.9, -0.8, \dots, 0.8, 0.9\}$  (note: the function `seq` may be useful here). For each value of  $\rho$ , approximate  $P(X > 0 \text{ and } Y > 0)$ , and plot these probabilities as a function of  $\rho$ , setting your vertical axis limits to (0,1) using `ylim`.