## Assignment 2

- 1. Redo Problem 2 from Assignment 1. Improve your definition of the dpareto function for computing the Pareto density; in particular make sure that the function is properly vectorized, handles the range restriction and invalid parameter values properly, and that the optional log argument is supported. Also make sure that your graphs properly show the discontinuity of the densities.
- 2. Several mechanisms are available for calling C code from R. The simplest is the .C interface. Modify your C function for computing the Pareto density so it can be used with the .C interface, and include some examples of its use in your writeup.

The Writing R Extensions manual provides documentation for the .C interface as well as other interfaces. A simple example is provided on the class web site that you may find helpful. The files in this example show how to use the .C interface as well as the richer .Call interface.

Your submission should include the new C source file and a source file with the R code for calling your C function. Track your work using Git and include your Git log in your submission.

Be sure to follow the coding standards. Tools are available to help:

- For C code, the indent program on Linux/Mac systems; these have different control options on Linux and Mac.
- For R code, the formatR package, in particular the tidy\_source function in that package.

You should submit your assignment electronically using Icon. Your submission should include

- your writeup as a PDF file
- files with your R code for Problem 1 and Problem 2
- a file with your C code for Problem 2
- the Git log for your work

Submit your work as a single compressed tar file. If your work is in a directory mywork then you can create a compressed tar file with the command

## tar czf mywork.tar.gz mywork

## Solutions and Comments

General comments:

- Don't include things not asked for (editor temporary files like foo , .git subdirectories, executables, shared libraries, etc).
- Make sure your file name ands with .tar.gz.
- Make sure you include a Git log in your submission when asked to.
- Avoid excessively large margins.
- 1. Please follow the coding standards on use of spaces and indentation and avoiding long lines.
  - Calls to return() are only needed for early exit. Do not use them otherwise.
  - Make sure your log = TRUE code works correctly.
  - Use brief but informative error messages.
  - It is usually numerically better to compute the log density and exponentiate than the other way around.
  - One possible definition:

```
dpareto <- function(x, a, b, log = FALSE) {</pre>
   nx <- length(x)</pre>
   na <- length(a)</pre>
   nb <- length(b)</pre>
   n <- max(nx, na, nb)</pre>
   if (nx < n) x < - rep(x, length.out = n)
   if (na < n) a <- rep(a, length.out = n)
   if (nb < n) b <- rep(b, length.out = n)
   ld <- ifelse(a > 0 & b > 0,
                 ifelse(x > a,
                         log(b) + b * log(a) - (b + 1) * log(x),
                         log(0)),
                 NaN)
   if (log) ld
   else exp(ld)
}
```

This could be simpler if the **ifelse** function did not base its result size entirely on the first argument.

• These are the tests I used:

Tierney

2. • Your C code should compile without errors or warnings. You can use the PKG\_CFLAGS environment variable to set C compiler flags to enable all warnings; e.g. with

env PKG\_CFLAGS="-Wall -pedantic" R CMD SHLIB ...

- It is not a good idea to allocate vectors that might be large as local variables. These will be allocated on the process stack and the space available for the stack is not very large.
- Follow the coding guide on use of spaces, indentation, and long lines.
- Standard convention in R density functions is to return NaN for bad parameter values.
- Do not use printf or REprintf for error messages; use the C level warning/error calls. Do not terminate the process when there is an error.
- It is usually numerically better to compute the log density and exponentiate than the other way around.
- In C code you should only warn about bad parameter values once, not on every loop iteration.
- You should not use loops in R if vectorized operations can be used.
- I used the same test code as in the first problem.