R functional programming approach for truncated and shifted and for shifted and truncated probability distribution functions

Introduction

R programming functional nature allows programmer to implement and manipulate functions as needed. Functional programming is a style of programming that emphasizes the evaluation of expressions rather than the execution of commands, so these functions produce results that depend only on their inputs and not on the program state. This style has a great power and includes lots of tools that combined together can allows us to produce results with the most efficient computational method. One of them is the functional factory, which is a factory for making new functions. It allows us to use closures, which are functions together with a referencing environment, to write specific functions that, in turn, can be used to generate new functions. So we can have a double layer of development: a first layer that is used to do all the complex work in common to all functions and a second layer that defines the details of each function. This concept was used to develop the computational method proposed by Andrea Spanò in "R package qdist: A functional programming approach to shifted probability distributions", and that I used in "R package qshift: A functional programming approach to shifted probability distributions", that produces functions for truncated and shifted distributions, respectively. qdist and qshift packages are available on Github:

```
devtools::install_github("veronicagiro/qtruncate")
devtools::install_github("veronicagiro/qshift")
```

In the previously cited articles, we dealt separately with shifted and truncated distributions. But in real life, we sometimes work with random variables arising from more than one transformation. So in this article, we will demonstrate that we can produce functions for truncated and shifted or for shifted and truncated distributions only combining the functions of the qdist and qshift packages. This is possible because of functional programming approach used to build them.

Distribution functions for truncated and shifted or for shifted and truncated distributions

R provides density, probability, quantile and random number generator functions in stats package. These functions share a stable naming convention both for the names of the functions and in the first argument of these functions. In particular:

- d<dist>(x) with x vector of quantiles, identifies density functions
- p<dist>(q) with q vector of quantiles, identifies probability distribution functions
- q<dist>(p) with p vector of probabilities, identifies quantile functions
- r<dist>(n) with n number of observations, identifies random number generator functions

where <dist> indicates the distribution family.

As a result we have: dweibull(x), pweibull(q), qweibull(p) and rweibull(n) for Weibull distribution and similarly for all other distributions.

Andrea Spanò proposed in qdist a functional programming method that computes truncated distribution functions for each distribution chosen without changing the well structured programming style and syntax of the previous functions.

The computational method proposed in qdist is based on two R key concepts:

- the environment of a function can be used as a placeholder for other objects (for details see Andrea Spanò article "R package qdist: A functional programming approach to truncated probability distributions")
- function formals can be manipulated

and it is made up of two steps:

1. defines the required functions suitable for computing from a specific truncated probability distribution by using the functions available within its package. These functions take as input a probability distribution as a character string, in this case the Weibull distribution, and return the equivalent truncated distribution function as object:

```
require(qdist)
dtweibull <- dtruncate("weibull") # density function
ptweibull <- ptruncate("weibull") # probability function
qtweibull <- qtruncate("weibull") # quantile function
rtweibull <- rtruncate("weibull") # random generator function</pre>
```

2. uses the newly created functions (dtweibull(), ptweibull(), qtweibull(), rtweibull()) for computing from truncated probability distributions.

Let us consider dtweibull(). It has the same formals as the original dweibull(), plus two extra parameters: L and U, respectively for lower and upper truncation thresholds set by default to -Inf and +Inf:

```
args(dweibull)
## function (x, shape, scale = 1, log = FALSE)
## NULL
args(dtweibull)
## function (x, shape, scale = 1, log = FALSE, L = -Inf, U = Inf)
## NULL
```

As dweibull() is the function that computes the density function for a Weibull distribution, conventionally named weibull, dtweibull() is the function that computes the density function for a truncated Weibull distribution, named tweibull.

Suppose we want to generate the density function for a **truncated and shifted** Weibull distribution. We need **qshift** package. **qshift** package shares the same logic and the same methodology of **qdist**. The difference is that its functions are designed to generate functions for shifted distributions. In particular, **qshift** package includes the following four functions:

- dshift(), for density function
- pshift(), for distribution function
- qshift(), for quantile function

• rshift(), for random generator function

So for computing the density function of a trucated and shifted Weibull distribution we have to set tweibull distribution as dshift() argument:

```
require(qshift)
dstweibull <- dshift("tweibull")</pre>
```

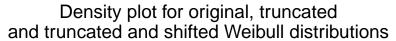
So dstweibull() is the function that computes the density function of stweibull, which is the truncated and shifted Weibull distribution.

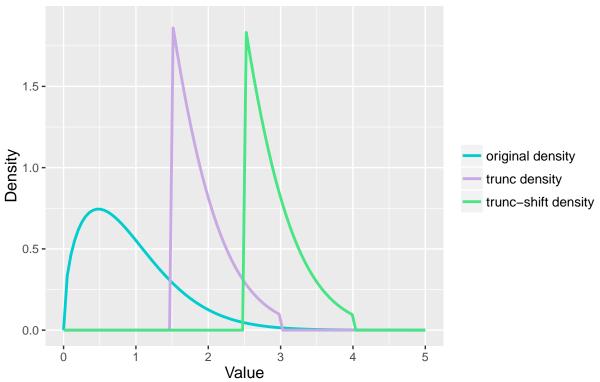
```
## function (x, shape, scale = 1, log = FALSE, L = -Inf, U = Inf,
## shift = 0)
## NULL
```

Looking at the function arguments, we see that the syntax style is that of dweibull() function plus L and U arguments for truncated component of the distribution and shift argument for shifted component of the distribution.

Let us graphically compare the results of dweibull(), dtweibull() and dstweibull() functions:

```
x \leftarrow seq(0, 5, len = 100)
orig density \leftarrow dweibull(x = x, shape = 1.5, scale = 1)
trunc_density <- dtweibull(x = x, shape = 1.5, scale = 1, L = 1.5, U = 3)
trunc_shift_density <- dstweibull(x = x, shape = 1.5, scale = 1, L = 1.5, U = 3, shift = 1)
df <- data.frame(x=x,</pre>
                 type = factor(c(rep("original density", times=length(x)),
                    rep("trunc density", times=length(x)),
                    rep("trunc-shift density", times=length(x))),
                    levels = c("original density", "trunc density", "trunc-shift density")),
                  value = c(orig_density, trunc_density, trunc_shift_density))
require(ggplot2)
pl <- ggplot(data = df, mapping = aes(x=x, y=value, col=type)) +</pre>
  geom_line(size=1) +
  scale_color_manual(values = c("original density" = "cyan3",
                                 "trunc density" = "#c8a9e3",
                                 "trunc-shift density" = "#47e583")) +
  xlab("Value") + ylab("Density") + ylim(c(0,1.9)) +
  ggtitle(paste("Density plot for original, truncated",
                "and truncated and shifted Weibull distributions", sep = "\n")) +
  theme(legend.title=element_blank(), legend.text= element_text(size=10),
  plot.title = element_text(size=14), axis.title=element_text(size=12))
print(pl)
```





The inverse procedure generates the density function for a shifted and truncated Weibull distribution.

Starting from dshift() function of qshift package, we define dsweibull(), the density function of sweibull, which is the shifted Weibull distribution:

```
dsweibull <- dshift("weibull") # density function</pre>
```

We have to define also the others functions for shifted distribution, as they are implicitly necessary for further step:

```
psweibull <- pshift("weibull") # probability function
qsweibull <- qshift("weibull") # quantile function
rsweibull <- rshift("weibull") # random generator function</pre>
```

The density function of a shifted and truncated Weibull distribution is generated setting sweibull distribution as ptruncate() argument:

```
dtsweibull <- dtruncate("sweibull")</pre>
```

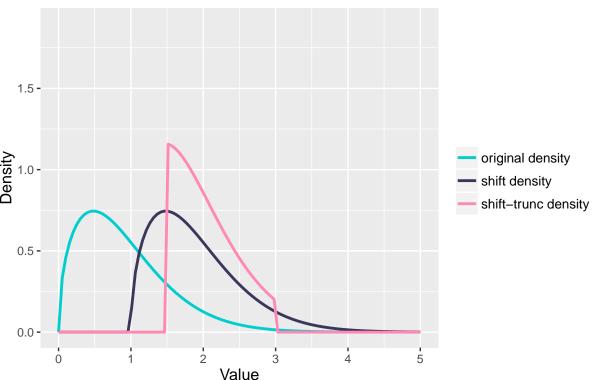
So dtsweibull() is the function that computes the density function of tsweibull, which is the shifted and truncated Weibull distribution.

Let us graphically represent the output of the three functions:

```
orig_density <- dweibull(x = x, shape = 1.5, scale = 1) shift_density <- dsweibull(x = x, shape = 1.5, scale = 1, shift = 1) shift_trunc_density <- dtsweibull(x = x, shape = 1.5, scale = 1, L = 1.5, U = 3, shift = 1)
```

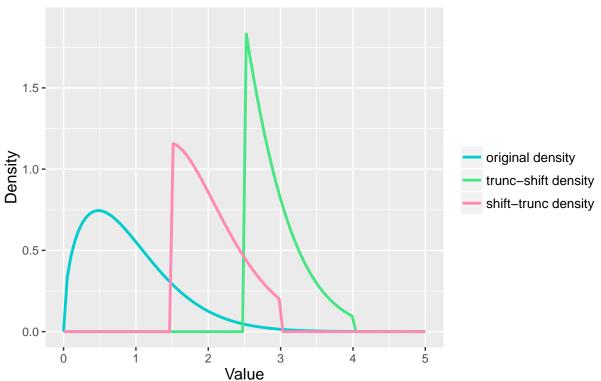
```
df <- data.frame(x=x,</pre>
                 type = factor(c(rep("original density", times=length(x)),
                    rep("shift density", times=length(x)),
                    rep("shift-trunc density", times=length(x))),
                    levels = c("original density", "shift density", "shift-trunc density")),
                  value = c(orig_density, shift_density, shift_trunc_density))
pl <- ggplot(data = df, mapping = aes(x=x, y=value, col=type)) +
  geom line(size=1) +
  scale color manual(values = c("original density" = "cyan3",
                                "shift density" = "\#39375b",
                                 "shift-trunc density" = "#FF89B4")) +
  xlab("Value") + ylab("Density") + ylim(c(0,1.9)) +
  ggtitle(paste("Density plot for original, shifted and",
                "shifted and truncated Normal distributions", sep = "\n")) +
  theme(legend.title=element_blank(), legend.text= element_text(size=10),
  plot.title = element_text(size=14), axis.title=element_text(size=12))
print(pl)
```

Density plot for original, shifted and shifted and truncated Normal distributions



Let us see graphically the differences between the density function of original Weibull distribution, of truncated and shifted Weibull distribution and of shifted and truncated Weibull distribution. We refers to values computed in the previous examples.

Density plot for original, truncated-shifted and shifted-truncated Weibull distribution



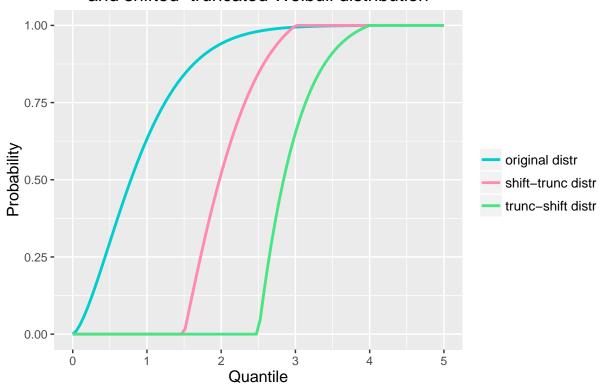
Looking at the figure we can see that a truncated and shifted distribution is different from a shifted and truncated one.

Let us analyze the other distribution functions:

• probability distribution function of a truncated and shifted Weibull distribution and of a shifted and truncated Weibull distribution:

```
# probability distribution function of a truncated and shifted Weibull distribution
pstweibull <- pshift("tweibull")</pre>
# probability distribution function of a shifted and truncated Weibull distribution
ptsweibull <- ptruncate("sweibull")</pre>
orig_distr <- pweibull(q = x, shape = 1.5, scale = 1)</pre>
trunc_shift_distr <- pstweibull(q = x, shape = 1.5, scale = 1, L = 1.5, U = 3, shift = 1)</pre>
shift_trunc_distr <- ptsweibull(q = x, shape = 1.5, scale = 1, L = 1.5, U = 3, shift = 1)</pre>
df <- data.frame(x=x,</pre>
                 type = factor(c(rep("original distr", times=length(x)),
                    rep("shift-trunc distr", times=length(x)),
                    rep("trunc-shift distr", times=length(x))),
                    levels = c("original distr", "shift-trunc distr", "trunc-shift distr")),
                  value = c(orig_distr, shift_trunc_distr, trunc_shift_distr))
pl <- ggplot(data = df, mapping = aes(x=x, y=value, col=type)) +</pre>
  geom line(size=1) +
  scale_color_manual(values = c("original distr" = "cyan3",
                                 "shift-trunc distr" = "#FF89B4",
                                 "trunc-shift distr" = "#47e583")) +
  xlab("Quantile") + ylab("Probability") +
  ggtitle(paste("Distribution plot for original, truncated-shifted",
                "and shifted-truncated Weibull distribution", sep="\n")) +
  theme(legend.title=element_blank(), legend.text= element_text(size=10),
  plot.title = element_text(size=14), axis.title=element_text(size=12))
print(pl)
```

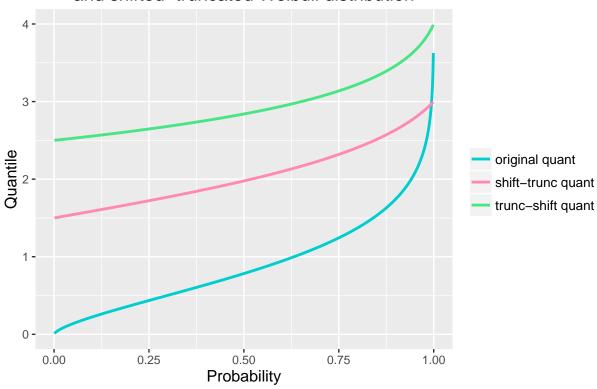
Distribution plot for original, truncated—shifted and shifted—truncated Weibull distribution



• quantile function of a truncated and shifted Weibull distribution and of a shifted and truncated Weibull distribution:

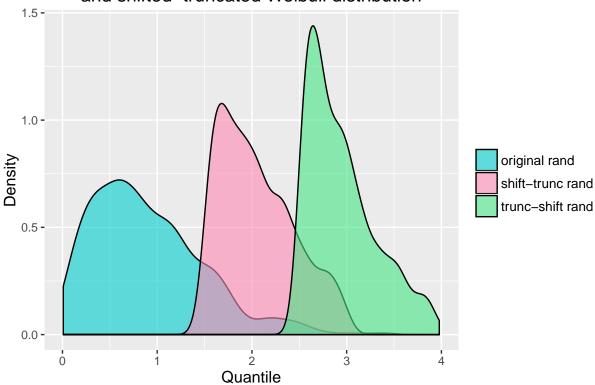
```
# quantile function of a truncated and shifted Weibull distribution
qstweibull <- qshift("tweibull")</pre>
# quantile function of a shifted and truncated Weibull distribution
qtsweibull <- qtruncate("sweibull")</pre>
x <- 1:999/1000
orig_quant <- qweibull(p = x, shape = 1.5, scale = 1)</pre>
trunc_shift_quant <- qstweibull(p = x, shape = 1.5, scale = 1, L = 1.5, U = 3, shift = 1)
shift_trunc_quant <- qtsweibull(p = x, shape = 1.5, scale = 1, L = 1.5, U = 3, shift = 1)</pre>
df <- data.frame(x=x.</pre>
                  type = factor(c(rep("original quant", times=length(x)),
                     rep("shift-trunc quant", times=length(x)),
                     rep("trunc-shift quant", times=length(x))),
                     levels = c("original quant", "shift-trunc quant", "trunc-shift quant")),
                   value = c(orig_quant, shift_trunc_quant, trunc_shift_quant))
pl <- ggplot(data = df, mapping = aes(x=x, y=value, col=type)) +</pre>
  geom_line(size=1) +
  scale_color_manual(values = c("original quant" = "cyan3",
                                 "shift-trunc quant" = "#FF89B4",
                                 "trunc-shift quant" = "#47e583")) +
  xlab("Probability") + ylab("Quantile") +
```

Quantile plot for original, truncated—shifted and shifted—truncated Weibull distribution



• random generation function of a truncated and shifted Weibull distribution and of a shifted and truncated Weibull distribution:

Density plot for original, truncated-shifted and shifted-truncated Weibull distribution



Extending the computation

Once we demonstrated that we can generate distribution functions for truncated and shifted and for shifted and truncated distributions, we can use them for further implementations. Suppose we want to define a function for maximum likelihood estimate for truncated and shifted distributions and one for shifted and truncated distributions. In the following examples, we will continue to consider the Weibull distribution.

Let us start from the function for maximum likelihood estimate for truncated and shifted Weibull distribution. We consider the previously defined density function, dstweibull(), for a truncated and shifted Weibull distribution and we use it inside the estimator function:

```
ltweibull <- function(x, shift = 0 ,L = -Inf, U = Inf){
  x1 \leftarrow x - shift
  # starting values for parameters (scale, shape)
  shape <- (sd(x1)/mean(x1))^{(-1.086)}
  scale <- mean(x1)/gamma(1+1/shape)</pre>
  # parameters vector definition
  theta <- c(shape, scale)
  # likelihood function
  11 <- function(theta, x, shift = 0, L = -Inf, U = Inf){</pre>
    shape <- theta[1]</pre>
    scale <- theta[2]</pre>
    ld <- dstweibull(x = x, shape = shape, scale = scale, shift = shift, L = L, U = U, log=TRUE)</pre>
    -sum(1d)
  # maximum likelihood estimation
  optim(par = theta, fn = ll, x = x, shift=shift, L = L, U = U, method = "Nelder-Mead")[["par"]]
}
```

Let's test it to see how it works.

We consider the previously defined random generator function rstweibull() for a truncated and shifted Weibull distribution to create the sample for testing the maximum likelihood estimate function:

```
x \leftarrow rstweibull(n = 100000, shape = 1.6, scale = 1, shift = 1, L = 1, U = 2.5)
# maximum likelihood estimate for the shifted and truncated weibull distribution
ltweibull(x = x, shift=1, L = 1, U = 2.5)
```

```
## [1] 1.593260 0.997412
```

The maximum likelihood estimate function produces a good estimate of the scale and shape parameters of truncated and shifted Weibull distribution.

Let us continue with the function for maximum likelihood estimate for shifted and truncated Weibull distribution.

We will consider the previously defined density function, dtsweibull(), for a shifted and truncated Weibull distribution and we use it inside the estimator function:

```
ltweibull <- function(x, shift = 0, L = -Inf, U = Inf){</pre>
  x1 \leftarrow x - shift
  # starting values for parameters (scale, shape)
  shape <- (sd(x1)/mean(x1))^{(-1.086)}
  scale <- mean(x1)/gamma(1+1/shape)</pre>
  # parameters vector definition
  theta <- c(shape, scale)
  # likelihood function
  11 \leftarrow function(theta, x, L = -Inf, U = Inf){
    shape <- theta[1]</pre>
    scale <- theta[2]</pre>
    ld <- dtsweibull(x = x, shape = shape, scale = scale, shift = shift, L = L, U = U, log=T)</pre>
    -sum(1d)
  }
  # maximum likelihood estimation
  optim(par = theta, fn = 11, x = x, L = L, U = U, method = "Nelder-Mead")[["par"]]
}
```

Let's test it to see how it works.

We consider the previously defined random generator function rtsweibull() for a shifted and truncated Weibull distribution to create the sample for testing the maximum likelihood estimate function:

```
x <- rtsweibull(n = 100000, shape = 1.6, scale = 1, shift = 1, L = 0.5, U = 3)
# maximum likelihood estimate for the shifted weibull distribution
ltweibull(x = x, shift = 1, L = 0.5, U = 3)</pre>
```

```
## [1] 1.6035625 0.9992417
```

Also in this case, the maximum likelihood estimate function produces a good estimate of the scale and shape parameters of shifted and truncated Weibull distribution.

Bibliography

Giro V. . R package qshift: A functional programming approach to shifted probability ditributions. url Spanò A. . R package qdist: A functional programming approach to truncated probability ditributions. url Spanò A. . Ramarro. R for Developers. http://www.quantide.com/ramarro-r-for-developers/. Accessed: 2016-09-20

Wickham H. . Advanced r. http://adv-r.had.co.nz/. Accessed: 2016-09-20.