

Risk theory features of **actuar**

Christophe Dutang
ISFA, Université Claude Bernard Lyon 1

Vincent Goulet
École d'actuariat, Université Laval

Mathieu Pigeon
École d'actuariat, Université Laval

1 Introduction

Risk theory refers to a body of techniques to model and measure the risk associated with a portfolio of insurance contracts. A first approach consists in modeling the distribution of total claims over a fixed period of time using the classical collective model of risk theory. A second input of interest to the actuary is the evolution of the surplus of the insurance company over many periods of time. In *ruin theory*, the main quantity of interest is the probability that the surplus becomes negative, in which case technical ruin of the insurance company occurs.

The interested reader can read more on these subjects in [Klugman et al. \(2004\)](#); [Gerber \(1979\)](#); [Denuit and Charpentier \(2004\)](#); [Kaas et al. \(2001\)](#), among others.

The current version of **actuar** ([Dutang et al., 2008](#)) contains four visible functions related to the above problems: two for the calculation of the aggregate claim amount distribution and two for ruin probability calculations.

2 The collective risk model

Let random variable S represent the aggregate claim amount (or total amount of claims) of a portfolio of independent risks over a fixed period of time, random variable N represent the number of claims (or frequency) in the portfolio over that period, and random variable C_j represent the amount of claim j (or severity). Then, we have the random sum

$$S = C_1 + \cdots + C_N, \tag{1}$$

where we assume that C_1, C_2, \dots are mutually independent and identically distributed random variables each independent of N . The task at hand consists in evaluating numerically the cdf of S , given by

$$\begin{aligned} F_S(x) &= \Pr[S \leq x] \\ &= \sum_{n=0}^{\infty} \Pr[S \leq x | N = n] p_n \\ &= \sum_{n=0}^{\infty} F_C^{*n}(x) p_n, \end{aligned} \quad (2)$$

where $F_C(x) = \Pr[C \leq x]$ is the common cdf of C_1, \dots, C_n , $p_n = \Pr[N = n]$ and $F_C^{*n}(x) = \Pr[C_1 + \dots + C_n \leq x]$ is the n -fold convolution of $F_C(\cdot)$. If C is discrete on $0, 1, 2, \dots$, one has

$$F_C^{*k}(x) = \begin{cases} I\{x \geq 0\}, & k = 0 \\ F_C(x), & k = 1 \\ \sum_{y=0}^x F_C^{*(k-1)}(x-y) f_C(y), & k = 2, 3, \dots \end{cases} \quad (3)$$

3 Discretization of claim amount distributions

Some numerical techniques to compute the aggregate claim amount distribution (see Section 4) require a discrete arithmetic claim amount distribution; that is, a distribution defined on $0, h, 2h, \dots$ for some step (or span, or lag) h . The package provides function `discretize` to discretize a continuous distribution. (The function can also be used to modify the support of an already discrete distribution, but this requires additional care.)

Let $F(x)$ denote the cdf of the distribution to discretize on some interval (a, b) and f_x denote the probability mass at x in the discretized distribution. Currently, `discretize` supports the following four discretization methods.

1. Upper discretization, or forward difference of $F(x)$:

$$f_x = F(x+h) - F(x) \quad (4)$$

for $x = a, a+h, \dots, b-h$. The discretized cdf is always above the true cdf.

2. Lower discretization, or backward difference of $F(x)$:

$$f_x = \begin{cases} F(a), & x = a \\ F(x) - F(x-h), & x = a+h, \dots, b. \end{cases} \quad (5)$$

The discretized cdf is always under the true cdf.

3. Rounding of the random variable, or the midpoint method:

$$f_x = \begin{cases} F(a + h/2), & x = a \\ F(x + h/2) - F(x - h/2), & x = a + h, \dots, b - h. \end{cases} \quad (6)$$

The true cdf passes exactly midway through the steps of the discretized cdf.

4. Unbiased, or local matching of the first moment method:

$$f_x = \begin{cases} \frac{E[X \wedge a] - E[X \wedge a + h]}{h} + 1 - F(a), & x = a \\ \frac{2E[X \wedge x] - E[X \wedge x - h] - E[X \wedge x + h]}{h}, & a < x < b \\ \frac{E[X \wedge b] - E[X \wedge b - h]}{h} - 1 + F(b), & x = b. \end{cases} \quad (7)$$

The discretized and the true distributions have the same total probability and expected value on (a, b) .

Figure 1 illustrates the four methods. It should be noted that although very close in this example, the rounding and unbiased methods are not identical.

Usage of `discretize` is similar to R's `plotting function curve`. The cdf to discretize and, for the unbiased method only, the limited expected value function are passed to `discretize` as expressions in `x`. The other arguments are the upper and lower bounds of the discretization interval, the step h and the discretization method. For example, upper and unbiased discretizations of a `Gamma(2, 1)` distribution on $(0, 17)$ with a step of 0.5 are achieved with, respectively,

```
> fx <- discretize(pgamma(x, 2, 1), method = "upper",
+   from = 0, to = 17, step = 0.5)
> fx <- discretize(pgamma(x, 2, 1), method = "unbiased",
+   lev = levgamma(x, 2, 1), from = 0, to = 17,
+   step = 0.5)
```

Function `discretize` is written in a modular fashion making it simple to add other discretization methods if needed.

4 Calculation of the aggregate claim amount distribution

Function `aggregateDist` serves as a unique front end for various methods to compute or approximate the cdf of the aggregate claim amount random variable S . Currently, five methods are supported.

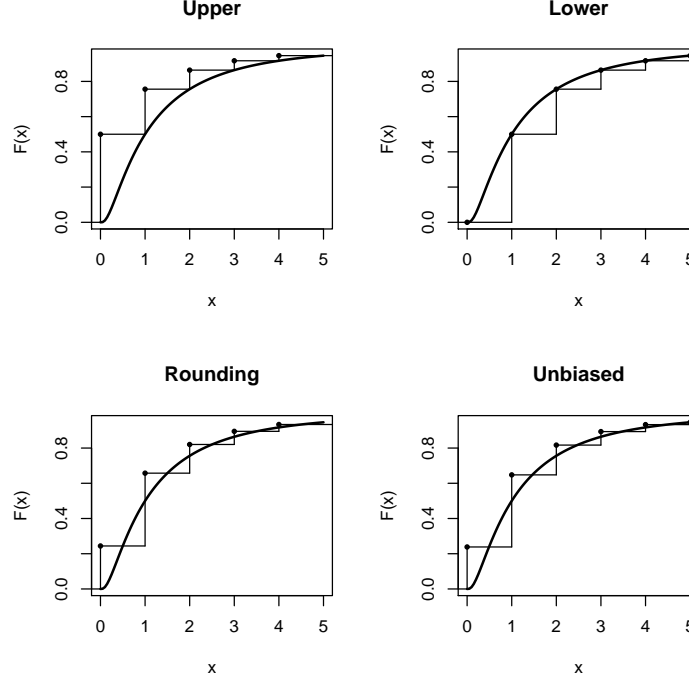


Figure 1: Comparison of four discretization methods

1. Recursive calculation using the algorithm of [Panjer \(1981\)](#). This requires the severity distribution to be discrete arithmetic on $0, 1, 2, \dots, m$ for some monetary unit and the frequency distribution to be a member of either the $(a, b, 0)$ or $(a, b, 1)$ family of distributions ([Klugman et al., 2004](#)). (These families contain the Poisson, binomial, negative binomial and logarithmic distributions and their extensions with an arbitrary mass at $x = 0$.) The general recursive formula is:

$$f_S(x) = \frac{(p_1 - (a + b)p_0)f_C(x) + \sum_{y=1}^{\min(x,m)} (a + by/x)f_C(y)f_S(x - y)}{1 - af_C(0)},$$

with starting value $f_S(0) = P_N(f_C(0))$, where $P_N(\cdot)$ is the probability generating function of N . Probabilities are computed until their sum is arbitrarily close to 1.

The recursions are done in C to dramatically increase speed. One difficulty the programmer is facing is the unknown length of the output. This was solved using a common, simple and fast technique: first allocate an arbitrary amount of memory and double this amount each time the allocated space gets full.

2. Exact calculation by numerical convolutions using (2) and (3). This also requires a discrete severity distribution. However, there is no restriction on the shape of the frequency distribution. The package merely implements the sum (2), the convolutions being computed with R's function `convolve`, which in turn uses the Fast Fourier Transform. This approach is practical for small problems only, even on today's fast computers.
3. Normal approximation of the cdf, that is

$$F_S(x) \approx \Phi\left(\frac{x - \mu_S}{\sigma_S}\right), \quad (8)$$

where $\mu_S = E[S]$ and $\sigma_S^2 = \text{Var}[S]$. For most realistic models, this approximation is rather crude in the tails of the distribution.

4. Normal Power II approximation of the cdf, that is

$$F_S(x) \approx \Phi\left(-\frac{3}{\gamma_S} + \sqrt{\frac{9}{\gamma_S^2} + 1} + \frac{6}{\gamma_S} \frac{x - \mu_S}{\sigma_S}\right), \quad (9)$$

where $\gamma_S = E[(S - \mu_S)^3]/\sigma_S^3$. The approximation is valid for $x > \mu_S$ only and performs reasonably well when $\gamma_S < 1$. See [Daykin et al. \(1994\)](#) for details.

5. Simulation of a random sample from S and approximation of $F_S(x)$ by the empirical cdf (??). The simulation itself is done with function `simul` (see the "simulation" vignette). This function admits very general hierarchical models for both the frequency and the severity components.

Here also, adding other methods to `aggregateDist` is simple due to its modular conception.

The arguments of `aggregateDist` differ depending on the calculation method; see the help page for details. One interesting argument to note is `x.scale` to specify the monetary unit of the severity distribution. This way, one does not have to mentally do the conversion between the support of $0, 1, 2, \dots$ assumed by the recursive and convolution methods and the true support of S .

The function returns an object of class "aggregateDist" inheriting from the "function" class. Thus, one can use the object as a function to compute the value of $F_S(x)$ in any x .

For illustration purposes, consider the following model: the distribution of S is a compound Poisson with parameter $\lambda = 10$ and severity distribution $\text{Gamma}(2, 1)$. To obtain an approximation of the cdf of S we first discretize the gamma distribution on $(0, 22)$ with the unbiased method and a step of 0.5, and then use the recursive method in `aggregateDist`:

```

> fx <- discretize(pgamma(x, 2, 1), from = 0, to = 22,
+   step = 0.5, method = "unbiased", lev = levgamma(x,
+   2, 1))
> Fs <- aggregateDist("recursive", model.freq = "poisson",
+   model.sev = fx, lambda = 10, x.scale = 0.5)
> summary(Fs)

```

Aggregate Claim Amount Empirical CDF:

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
0.0	14.0	19.0	20.0	24.5	71.0

Hence, object Fs contains an empirical cdf with support

```

> knots(Fs)

[1] 0.0 0.5 1.0 1.5 2.0 2.5 3.0 3.5 4.0 4.5 5.0
[12] 5.5 6.0 6.5 7.0 7.5 8.0 8.5 9.0 9.5 10.0 10.5
[23] 11.0 11.5 12.0 12.5 13.0 13.5 14.0 14.5 15.0 15.5 16.0
[34] 16.5 17.0 17.5 18.0 18.5 19.0 19.5 20.0 20.5 21.0 21.5
[45] 22.0 22.5 23.0 23.5 24.0 24.5 25.0 25.5 26.0 26.5 27.0
[56] 27.5 28.0 28.5 29.0 29.5 30.0 30.5 31.0 31.5 32.0 32.5
[67] 33.0 33.5 34.0 34.5 35.0 35.5 36.0 36.5 37.0 37.5 38.0
[78] 38.5 39.0 39.5 40.0 40.5 41.0 41.5 42.0 42.5 43.0 43.5
[89] 44.0 44.5 45.0 45.5 46.0 46.5 47.0 47.5 48.0 48.5 49.0
[100] 49.5 50.0 50.5 51.0 51.5 52.0 52.5 53.0 53.5 54.0 54.5
[111] 55.0 55.5 56.0 56.5 57.0 57.5 58.0 58.5 59.0 59.5 60.0
[122] 60.5 61.0 61.5 62.0 62.5 63.0 63.5 64.0 64.5 65.0 65.5
[133] 66.0 66.5 67.0 67.5 68.0 68.5 69.0 69.5 70.0 70.5 71.0

```

A nice graph of this function is obtained with a method of plot (see Figure 2):

```

> plot(Fs, do.points = FALSE, verticals = TRUE, xlim = c(0,
+   60))

```

The package defines a few summary methods to extract information from "aggregateDist" objects. First, there are methods of mean and quantile to easily compute the mean and obtain the quantiles of the approximate distribution:

```

> mean(Fs)

[1] 20

> quantile(Fs)

 25%  50%  75%  90%  95% 97.5%  99% 99.5%
14.0 19.0 24.5 30.0 33.5 36.5 40.5 43.0

> quantile(Fs, 0.999)

99.9%
 49

```

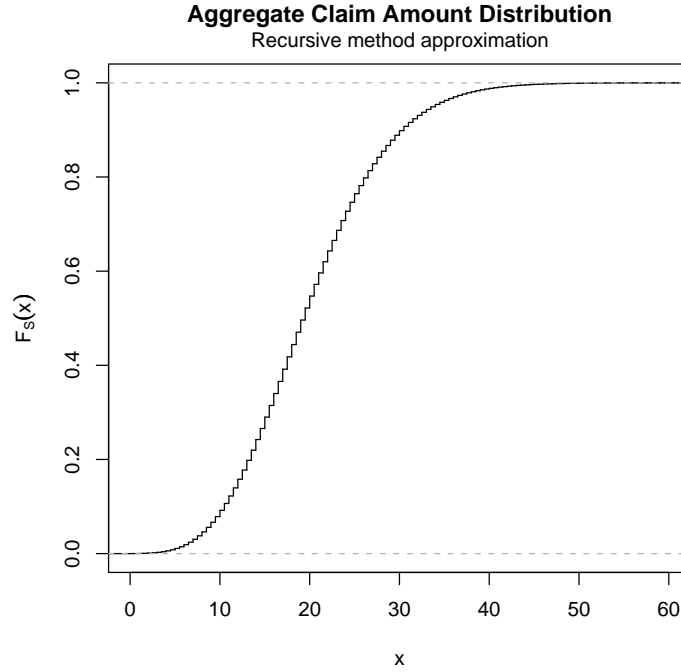


Figure 2: Graphic of the empirical cdf of S obtained with the recursive method

Second, the package introduces the generic functions VaR and CTE with methods for objects of class "aggregateDist". The former computes the value-at-risk VaR_α such that

$$\Pr[S \leq \text{VaR}_\alpha] = \alpha, \quad (10)$$

where α is the confidence level. Thus, the value-at-risk is nothing else than a quantile. As for the method of CTE, it computes the conditional tail expectation

$$\text{CTE}_\alpha = E[S | S > \text{VaR}_\alpha]. \quad (11)$$

Here are examples using object F_S obtained above:

```
> VaR(Fs)
 90%  95%  99%
30.0 33.5 40.5

> CTE(Fs)
 90%  95%  99%
34.98 38.10 44.54
```

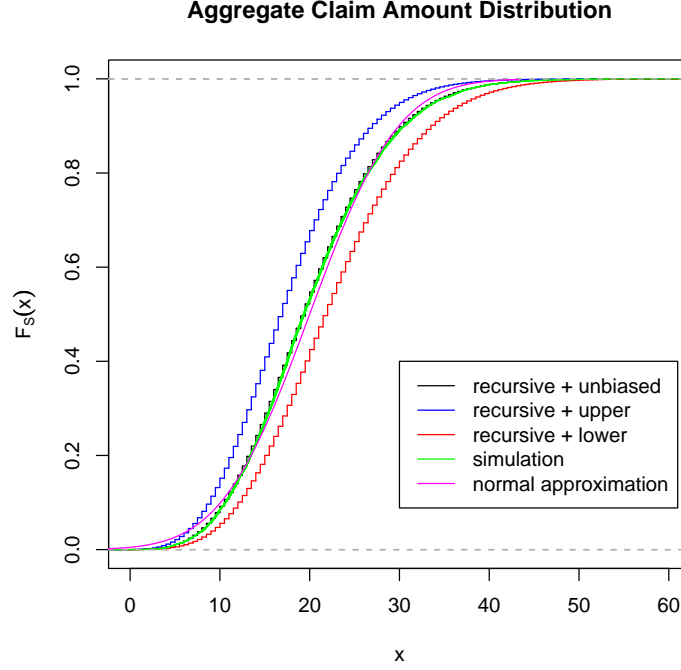


Figure 3: Comparison between the empirical or approximate cdf of S obtained with five different methods

To conclude on the subject, Figure 3 shows the cdf of S using five of the many combinations of discretization and calculation method supported by **actuar**.

5 The continuous time ruin model

We now turn to the multi-period ruin problem. Let $U(t)$ denote the surplus of an insurance company at time t , $c(t)$ denote premiums collected through time t , and $S(t)$ denote aggregate claims paid through time t . If u is the initial surplus at time $t = 0$, then a mathematically convenient definition of $U(t)$ is

$$U(t) = u + c(t) - S(t). \quad (12)$$

As mentioned previously, technical ruin of the insurance company occurs when the surplus becomes negative. Therefore, the definition of the infinite time probability of ruin is

$$\psi(u) = \Pr[U(t) < 0 \text{ for some } t \geq 0]. \quad (13)$$

We define some other quantities needed in the sequel. Let $N(t)$ denote the number of claims up to time $t \geq 0$ and C_j denote the amount of claim j . Then the definition of $S(t)$ is analogous to (1):

$$S(t) = C_1 + \dots + C_{N(t)}, \quad (14)$$

assuming $N(0) = 0$ and $S(t) = 0$ as long as $N(t) = 0$. Furthermore, let T_j denote the time when claim j occurs, such that $T_1 < T_2 < T_3 < \dots$. Then the random variable of the interarrival (or wait) time between claim $j - 1$ and claim j is defined as $W_1 = T_1$ and

$$W_j = T_j - T_{j-1}, \quad j \geq 2. \quad (15)$$

For the rest of this discussion, we make the following assumptions:

1. premiums are collected at a constant rate c , hence $c(t) = ct$;
2. the sequence $\{T_j\}_{j \geq 1}$ forms an ordinary renewal process, with the consequence that random variables W_1, W_2, \dots are independent and identically distributed;
3. claim amounts C_1, C_2, \dots are independent and identically distributed.

6 Adjustment coefficient

The quantity known as the adjustment coefficient ρ hardly has any physical interpretation, but it is useful as an approximation to the probability of ruin since we have the inequality

$$\psi(u) \leq e^{-\rho u}, \quad u \geq 0.$$

The adjustment coefficient is defined as the smallest strictly positive solution (if it exists) of the Lundberg equation

$$h(t) = E[e^{tC - tcW}] = 1, \quad (16)$$

where the premium rate c satisfies the positive safety loading constraint $E[C - cW] < 0$. If C and W are independent, as in the most common models, then the equation can be rewritten as

$$h(t) = M_C(t)M_W(-tc) = 1. \quad (17)$$

Function `adjCoef` of **actuar** computes the adjustment coefficient ρ from the following arguments: either the two moment generating functions $M_C(t)$ and $M_W(t)$ (thereby assuming independence) or else function $h(t)$; the premium rate c ; the upper bound of the support of $M_C(t)$ or any other upper bound for ρ .

For example, if W and C are independent, $W \sim \text{Exponential}(2)$, $C \sim \text{Exponential}(1)$ and the premium rate is $c = 2.4$ (for a safety loading of 20% using the expected value premium principle), then the adjustment coefficient is

```
> adjCoef(mgf.claim = mgfexp(x), mgf.wait = mgfexp(x,
+      2), premium.rate = 2.4, upper = 1)

[1] 0.1667
```

The function also supports models with proportional or excess-of-loss reinsurance (Centeno, 2002). Under the first type of treaty, an insurer pays a proportion α of every loss and the rest is paid by the reinsurer. Then, for fixed α the adjustment coefficient is the solution of

$$h(t) = E[e^{t\alpha C - tc(\alpha)W}] = 1. \quad (18)$$

Under an excess-of-loss treaty, the primary insurer pays each claim up to a limit L . Again, for fixed L , the adjustment coefficient is the solution of

$$h(t) = E[e^{t\min(C,L) - tc(L)W}] = 1. \quad (19)$$

For models with reinsurance, `adjCoef` returns an object of class "adjCoef" inheriting from the "function" class. One can then use the object to compute the adjustment coefficient for any retention rate α or retention limit L . The package also defines a method of `plot` for these objects.

For example, using the same assumptions as above with proportional reinsurance and a 30% safety loading for the reinsurer, the adjustment coefficient as a function of $\alpha \in [0, 1]$ is (see Figure 4 for the graph):

```
> mgfx <- function(x, y) mgfexp(x * y)
> p <- function(x) 2.6 * x - 0.2
> rho <- adjCoef(mgfx, mgfexp(x, 2), premium = p,
+      upper = 1, reins = "prop", from = 0, to = 1)
> rho(c(0.75, 0.8, 0.9, 1))

[1] 0.1905 0.1862 0.1765 0.1667

> plot(rho)
```

7 Probability of ruin

In this subsection, we always assume that interarrival times and claim amounts are independent.

The main difficulty with the calculation of the infinite time probability of ruin lies in the lack of explicit formulas except for the most simple models. If interarrival times are $\text{Exponential}(\lambda)$ distributed (Poisson claim number process) and claim amounts are $\text{Exponential}(\beta)$ distributed, then

$$\psi(u) = \frac{\lambda}{c\beta} e^{-(\beta - \lambda/c)u}. \quad (20)$$

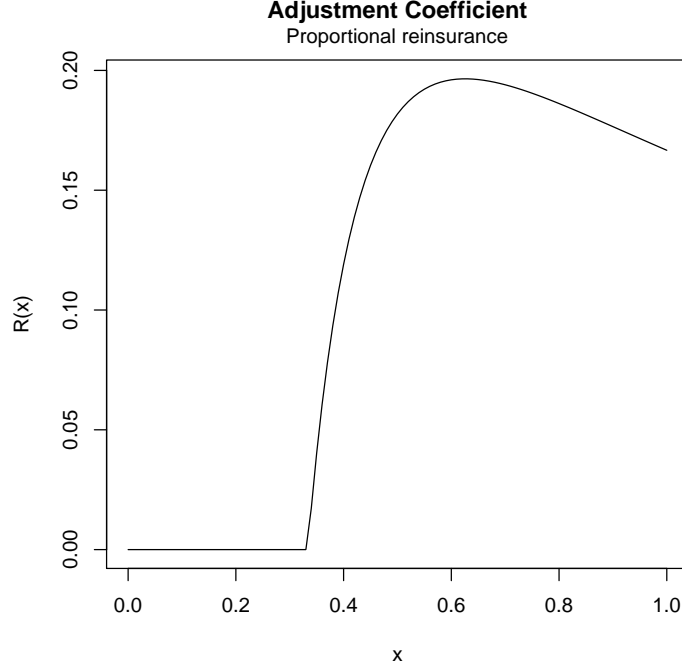


Figure 4: Adjustment coefficient as a function of the retention rate

If the frequency assumption of this model is defensible, the severity assumption can hardly be used beyond illustration purposes.

Fortunately, phase-type distributions have come to the rescue since the early 1990s. [Asmussen and Rolski \(1991\)](#) first show that in the classical Cramér-Lundberg model where interarrival times are $\text{Exponential}(\lambda)$ distributed, if claim amounts are $\text{Phase-type}(\boldsymbol{\pi}, \mathbf{T})$ distributed, then $\psi(u) = 1 - F(u)$, where F is $\text{Phase-type}(\boldsymbol{\pi}_+, \mathbf{Q})$ with

$$\begin{aligned}\boldsymbol{\pi}_+ &= -\frac{\lambda}{c} \boldsymbol{\pi} \mathbf{T}^{-1} \\ \mathbf{Q} &= \mathbf{T} + \mathbf{t} \boldsymbol{\pi}_+, \end{aligned} \tag{21}$$

and $\mathbf{t} = -\mathbf{T}\mathbf{e}$, \mathbf{e} is a column vector with all components equal to 1; see the "lossdist" vignette for details.

In the more general Sparre Andersen model where interarrival times can have any $\text{Phase-type}(\nu, \mathbf{S})$ distribution, [Asmussen and Rolski \(1991\)](#) also show that using the same claim severity assumption as above, one still has $\psi(u) = 1 - F(u)$ where F is $\text{Phase-type}(\boldsymbol{\pi}_+, \mathbf{Q})$, but with parameters

$$\boldsymbol{\pi}_+ = \frac{\mathbf{e}'(\mathbf{Q} - \mathbf{T})}{c\mathbf{e}'\mathbf{t}} \tag{22}$$

and \mathbf{Q} solution of

$$\begin{aligned}\mathbf{Q} &= \Psi(\mathbf{Q}) \\ &= \mathbf{T} - \mathbf{t}\boldsymbol{\pi} \left[(\mathbf{I}_n \otimes \boldsymbol{\nu})(\mathbf{Q} \oplus \mathbf{S})^{-1}(\mathbf{I}_n \otimes \mathbf{s}) \right].\end{aligned}\tag{23}$$

In the above, $\mathbf{s} = -\mathbf{S}\mathbf{e}$, \mathbf{I}_n is the $n \times n$ identity matrix, \otimes denotes the usual Kronecker product between two matrices and \oplus is the Kronecker sum defined as

$$\mathbf{A}_{m \times m} \oplus \mathbf{B}_{n \times n} = \mathbf{A} \otimes \mathbf{I}_n + \mathbf{B} \otimes \mathbf{I}_m.\tag{24}$$

Function `ruin` of **actuar** returns a function object of class "ruin" to compute the probability of ruin for any initial surplus u . In all cases except the exponential/exponential model where (20) is used, the output object calls function `pphtype` to compute the ruin probabilities.

Some thought went into the interface of `ruin`. Obviously, all models can be specified using phase-type distributions, but the authors wanted users to have easy access to the most common models involving exponential and Erlang distributions. Hence, one first states the claim amount and inter-arrival times models with any combination of "exponential", "Erlang" and "phase-type". Then, one passes the parameters of each model using lists with components named after the corresponding parameters of `dexp`, `dgamma` and `dphtype`. If a component "weights" is found in a list, the model is a mixture of exponential or Erlang (mixtures of phase-type are not supported). Every component of the parameter lists is recycled as needed.

The following examples should make the matter clearer. (All examples use $c = 1$, the default value in `ruin`.) First, for the exponential/exponential model, one has

```
> psi <- ruin(claims = "e", par.claims = list(rate = 5),
+   wait = "e", par.wait = list(rate = 3))
> psi
```

```
function (u, survival = FALSE, lower.tail = !survival)
{
  res <- 0.6 * exp(-(2) * u)
  if (lower.tail)
    res
  else 0.5 - res + 0.5
}
<environment: 0x2be8d50>
attr(,"class")
[1] "ruin"      "function"
```

```
> psi(0:10)

[1] 6.000e-01 8.120e-02 1.099e-02 1.487e-03 2.013e-04
[6] 2.724e-05 3.687e-06 4.989e-07 6.752e-08 9.138e-09
[11] 1.237e-09
```

Second, for a mixture of two exponentials claim amount model and exponential interarrival times, the simplest call to `ruin` is

```
> ruin(claims = "e", par.claims = list(rate = c(3,
+   7), weights = 0.5), wait = "e", par.wait = list(rate = 3))

function (u, survival = FALSE, lower.tail = !survival)
pphype(u, c(0.5, 0.214285714285714), c(-1.5, 3.5, 0.642857142857143,
-5.5), lower.tail = !lower.tail)
<environment: 0x2c5b454>
attr("class")
[1] "ruin"      "function"
```

Finally, one will obtain a function to compute ruin probabilities in a model with phase-type claim amounts and mixture of exponentials interarrival times with

```
> prob <- c(0.5614, 0.4386)
> rates <- matrix(c(-8.64, 0.101, 1.997, -1.095),
+   2, 2)
> ruin(claims = "p", par.claims = list(prob = prob,
+   rates = rates), wait = "e", par.wait = list(rate = c(5,
+   1), weights = c(0.4, 0.6)))

function (u, survival = FALSE, lower.tail = !survival)
pphype(u, c(0.146595513877824, 0.761505562273639), c(-7.66616600130962,
0.246715940794557, 7.05568145018378, -0.338063471100003), lower.tail = !lower.tail)
<environment: 0x1aa63c8>
attr("class")
[1] "ruin"      "function"
```

To ease plotting of the probability of ruin function, the package provides a method of `plot` for objects returned by `ruin` that is a simple wrapper for `curve` (see Figure 5):

```
> psi <- ruin(claims = "p", par.claims = list(prob = prob,
+   rates = rates), wait = "e", par.wait = list(rate = c(5,
+   1), weights = c(0.4, 0.6)))
> plot(psi, from = 0, to = 50)
```

8 Approximation to the probability of ruin

When the model for the aggregate claim process (14) does not fit nicely into the framework of the previous section, one can compute ruin probabilities using the so-called Beekman's convolution formula (Beekman, 1968; Kass, 2004).

Let the surplus process and the aggregate claim amount process be defined as in (12) and (14), respectively, and let $\{N(t)\}$ be a Poisson process

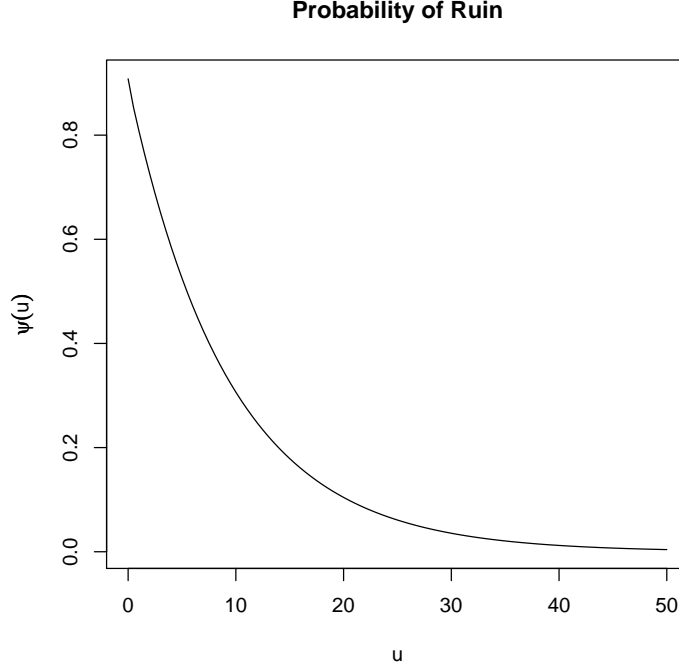


Figure 5: Graphic of the probability of ruin as a function of the initial surplus u

with mean λ . As before, claim amounts C_1, C_2, \dots are independent and identically distributed with cdf $P(\cdot)$ and mean $\mu = E[C_1]$. Then the infinite time probability of ruin is given by

$$\psi(u) = 1 - F(u), \quad (25)$$

where $F(\cdot)$ is Compound Geometric(p, H) with

$$p = 1 - \frac{\lambda\mu}{c} \quad (26)$$

and

$$H(x) = \int_0^x \frac{1 - P(y)}{\mu} dy. \quad (27)$$

In other words, we have (compare with (2))

$$\psi(u) = 1 - \sum_{n=0}^{\infty} H^{*n}(u)p(1-p)^n. \quad (28)$$

In most practical situations, numerical evaluation of (28) is done using Panjer's recursive formula. This usually requires discretization of $H(\cdot)$. In

such circumstances, Beekman's formula yields approximate ruin probabilities.

For example, let claim amounts have a Pareto(5, 4) distribution, that is

$$P(x) = 1 - \left(\frac{4}{4+x} \right)^5$$

and $\mu = 1$. Then

$$\begin{aligned} H(x) &= \int_0^x \left(\frac{4}{4+y} \right)^5 dy \\ &= 1 - \left(\frac{4}{4+x} \right)^4, \end{aligned}$$

or else H is Pareto(4, 4). Furthermore, we determine the premium rate c with the expected value premium principle and a safety loading of 20%, that is $c = 1.2\lambda\mu$. Thus, $p = 0.2/1.2 = 1/6$.

One can get functions to compute lower bounds and upper bounds for $F(u)$ with functions `discretize` and `aggregateDist` as follows:

```
> f.L <- discretize(ppareto(x, 4, 4), from = 0, to = 200,
+   step = 1, method = "lower")
> f.U <- discretize(ppareto(x, 4, 4), from = 0, to = 200,
+   step = 1, method = "upper")
> F.L <- aggregateDist(method = "recursive", model.freq = "geometric",
+   model.sev = f.L, prob = 1/6)
> F.U <- aggregateDist(method = "recursive", model.freq = "geometric",
+   model.sev = f.U, prob = 1/6)
```

Corresponding functions for the probability of ruin $\psi(u)$ lower and upper bounds are (see figure 6 for the graphic):

```
> psi.L <- function(u) 1 - F.U(u)
> psi.U <- function(u) 1 - F.L(u)
> u <- seq(0, 50, by = 5)
> cbind(lower = psi.L(u), upper = psi.U(u))
```

	lower	upper
[1,]	0.6719160	0.83333
[2,]	0.2892792	0.51572
[3,]	0.1361541	0.32938
[4,]	0.0662486	0.21200
[5,]	0.0329848	0.13700
[6,]	0.0167551	0.08877
[7,]	0.0086802	0.05764
[8,]	0.0045911	0.03749
[9,]	0.0024843	0.02443
[10,]	0.0013790	0.01595
[11,]	0.0007877	0.01043

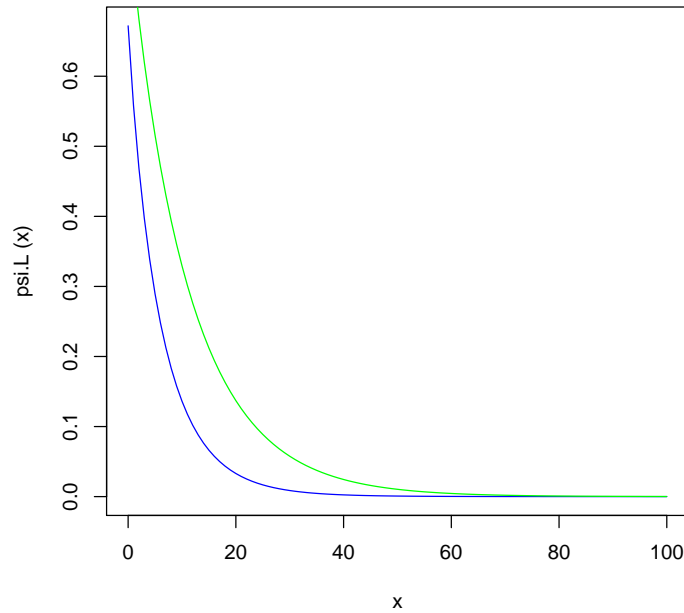


Figure 6: Lower and upper bounds for the probability of ruin as determined using Beekman's convolution formula.

```
> curve(psi.L, from = 0, to = 100, col = "blue")
> curve(psi.U, add = TRUE, col = "green")
```

One can make the bounds as close as one wishes by reducing the discretization step.

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