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parfm: Parametric Frailty Models in R

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Abstract

Frailty models are getting more and more popular to account for overdispersion and/or clustering in survival data. When the form of the baseline hazard is somehow known in advance, the parametric estimation approach can be used advantageously. Nonetheless, there is no unified widely available software that deals with the parametric frailty model. The new **parfm** package remedies that lack by providing a wide range of parametric frailty models in R. The gamma, inverse Gaussian, and positive stable frailty distributions can be specified, together with five different baseline hazards. Parameter estimation is done by maximising the marginal log-likelihood, with right-censored and possibly left-truncated data. In the multivariate setting, the inverse Gaussian may encounter numerical difficulties with a huge number of events in at least one cluster. The positive stable model shows analogous difficulties but an ad-hoc solution is implemented, whereas the gamma model is very resistant due to the simplicity of its Laplace transform.

Keywords: parametric frailty models, survival analysis, gamma, positive stable, inverse gaussian, weibull, exponential, gompertz, loglogistic, lognormal, R, **parfm**.

1. Introduction

Survival data, or time-to-event data, measure the time elapsed from a given origin to the occurrence of an event of interest. The observation of survival data is very common in the medical fields where, for instance, the clinician is interested in the time to relapse of a pathology after the therapy. However, the researcher cannot always observe the event due to censoring. Right-censoring occurs when the time of interest cannot be observed but only a lower bound is available. Particular techniques are therefore required as described by a number of textbooks, e.g., [Klein and Moeschberger \(2003\)](#).

Most commonly, survival data are handled by means of the proportional hazards regression model popularised by [Cox \(1972\)](#). But correct inference based on those proportional hazards models needs independent and identically distributed samples. Nonetheless, subjects may be

exposed to different risk levels, even after controlling for known risk factors; this is because some relevant covariates are often unavailable to the researcher or even unknown (univariate case). Also, the study population may be divided into clusters so that subjects from the same cluster behave more cohesively than subjects from different clusters (multivariate case). Lots of examples of clustered survival data arise from large-scale clinical trials in which patients are recruited at several hospital centres (Duchateau, Janssen, Lindsey, Legrand, Nguti, and Sylvester 2002; Glidden and Vittinghoff 2004). Another classical example is the analysis of lifetimes of matched human organs such as eyes or kidneys.

The frailty model, introduced in the biostatistical literature by Vaupel, Manton, and Stallard (1979), and discussed in details by Hougaard (2000), Duchateau and Janssen (2008), and by Wienke (2010), accounts for this heterogeneity in baseline. It is an extension of the proportional hazards model in which the hazard function depends upon an unobservable random quantity, the so-called frailty, that acts multiplicatively on it.

The gamma frailty model assumes a gamma distribution for the frailties. Arguably, this is the most popular frailty model due to its mathematical tractability. The lognormal frailty model is also well-liked for its strong link with generalised linear mixed models. Other frailty distributions include the positive stable and the inverse Gaussian. All of these are reviewed by Duchateau and Janssen (2008, Chapter 4)

Of particular interest in the multivariate case is the association between related event times. Indeed, different dependence structures result from different frailty distributions (Hougaard 1995). In particular, positive stable frailties typically generate very strong dependence initially while, at equal global dependence, gamma frailties lead to stronger dependence at late times, and inverse Gaussian frailties are in between the two. These three distributions therefore cover a wide range of association structures in the data.

Estimation of the frailty model can be parametric or semi-parametric. In the former case, a parametric density is assumed for the event times, resulting in a parametric baseline hazard function. Estimation is then conducted by maximising the marginal log-likelihood (see Section 2). In the second case, the baseline hazard is left unspecified and more complex techniques are available to approach that situation (Cortiñas Abrahantes, Legrand, Burzykowski, Janssen, Ducrocq, and Duchateau 2007). Even though semi-parametric estimation offers more flexibility, the parametric estimation will be more powerful if the form of the baseline hazard is somehow known in advance. Further, the estimation technique is much simpler.

Slowly but surely, a variety of estimation procedures becomes available in standard statistical software. In R (R Development Core Team 2012), the `coxph()` function from the **survival** package (Therneau 2012b) handles the semi-parametric model with gamma and lognormal frailties. Important options supported by `coxph()` and its output are described in details by Therneau and Grambsch (2000, Chapter 9). Recently, the **frailtypack** package (Rondeau, Gonzalez, Mazroui, Mauguen, Diakite, and Laurent 2012a) by Rondeau and Gonzalez (2005) and Rondeau, Mazroui, and Gonzalez (2012b) has been updated and it stands now for gamma frailty models with a semi-parametric estimation but also with a parametric approach using the Weibull baseline hazard. Other R packages include **coxme** (Therneau 2012a) and **phmm** (Donohue and Xu 2012). These two perform semi-parametric estimation in the lognormal frailty model. SAS (SAS Institute Inc. 2011) also deals with the lognormal distribution. On the one hand, `proc phreg` can now fit the semi-parametric lognormal frailty model. On the other hand, `proc nlmixed` deals with the parametric version by using Gaussian quadrature

to approach the marginal likelihood; see, e.g., Duchateau and Janssen (2008, Example 4.16). In the parametric setting, Stata (StataCorp. 2011) provides some flexibility. The `streg` command (Gutierrez 2002) is able to perform maximum likelihood estimation with various choices of baselines: exponential, Weibull, Gompertz, lognormal, loglogistic, and generalised gamma. Take notice, however, that Stata fits the accelerated failure time model. Still, with exponential or Weibull baselines, both the proportional hazards and the accelerated failure time representations are allowed. As for the frailty distribution, the gamma and the inverse Gaussian are the only two that are supported. On a side note, Bayesian analyses can be conducted in WinBUGS (Spiegelhalter, Thomas, Best, and Lunn 2003); see, e.g., Duchateau and Janssen (2008, Example 6.4). For a deeper overview of who supports what, and for a comparison of some of the aforementioned functions, see Hirsch and Wienke (2012).

Hereinbelow, we illustrate **parfm** (Rotolo, Munda, and Legrand 2012), a new R package that fits the gamma, the inverse Gaussian, and the positive stable proportional hazards frailty models with either exponential, Weibull, Gompertz, lognormal, or loglogistic baseline. The main advantage of **parfm** therefore relies on the large choice of frailty distributions and parametric baseline hazards it supports. Parameter estimation is done by maximising the marginal log-likelihood.

The model and the marginal log-likelihood are shown in Section 2. There, we also outline the estimation method, while Sections 2.1–2.3 provide details for the three frailty distributions supported by **parfm**. In Section 3, we apply **parfm** to a real dataset in order to illustrate its use and its output. Section 4 concludes with remarks.

2. Model estimation

From a modelling point of view, the multivariate model includes the univariate. Because of this, we shall mainly refer to the first. However, they are used in two different contexts: in the former case, the frailty distribution variability is related to a measure of dependence between clustered subjects, whereas it is rather interpreted as a measure of overdispersion in the latter.

Model

The frailty model is defined in terms of the conditional hazard

$$h_{ij}(t \mid u_i) = h_0(t)u_i \exp(\mathbf{x}_{ij}^\top \boldsymbol{\beta}),$$

with $i \in I = \{1, \dots, G\}$ and $j \in J_i = \{1, \dots, n_i\}$, where $h_0(\cdot)$ is the baseline hazard function, u_i the frailty term in group i , \mathbf{x}_{ij} the vector of covariates for subject j in group i , and $\boldsymbol{\beta}$ the vector of regression coefficients.

If the number of subjects n_i is 1 for all groups, then the univariate frailty model is obtained (Wienke 2010, Chapter 3), otherwise the model is called the shared frailty model (Hougaard 2000, Chapter 7; Duchateau and Janssen 2008) because all subjects in the same cluster share the same frailty value u_i .

Baseline hazard

Under the parametric approach, the baseline hazard is defined as a parametric function and the vector of its parameters, say $\boldsymbol{\psi}$, is estimated together with the regression coefficients

Distribution	$h_0(t)$	$H_0(t) = \int_0^t h_0(s)ds$	Parameter space
Exponential	λ	λt	$\lambda > 0$
Weibull	$\lambda \rho t^{\rho-1}$	λt^ρ	$\rho, \lambda > 0$
Gompertz	$\lambda \exp(\gamma t)$	$\frac{\lambda}{\gamma} (\exp(\gamma t) - 1)$	$\gamma, \lambda > 0$
Lognormal	$\frac{\phi\left(\frac{\log(t)-\mu}{\sigma}\right)}{\sigma t \left[1 - \Phi\left(\frac{\log(t)-\mu}{\sigma}\right)\right]}$	$-\log\left[1 - \Phi\left(\frac{\log(t)-\mu}{\sigma}\right)\right]$	$\mu \in \mathbb{R}, \sigma > 0$
Loglogistic	$\frac{\exp(\alpha)\kappa t^{\kappa-1}}{1 + \exp(\alpha)t^\kappa}$	$\log(1 + \exp(\alpha)t^\kappa)$	$\alpha \in \mathbb{R}, \kappa > 0$

Table 1: Parametric distributions available in **parfm** for the baseline hazard. With the lognormal, $\phi(\cdot)$ and $\Phi(\cdot)$ respectively denote the probability density and the cumulative distribution functions of a standard normal random variable.

and the frailty parameter(s). A bunch of possibilities are considered in the literature; in the **parfm** package the Weibull, exponential, Gompertz, lognormal, and loglogistic distributions are available. Table 1 shows the hazard and cumulative hazard functions for each of these distributions.

Frailty distribution

The frailty u_i is an unobservable realisation of a random variable U with probability density function $f(\cdot)$ – the frailty distribution. Since u_i multiplies the hazard function, U has to be non-negative. Another constraint is further needed for identifiability reasons, similar to the zero-mean constraint of a random effect in a standard linear mixed model. More specifically, the mean of U is typically restricted to unity when possible (i.e., when $E(U)$ exists) in order to separate the baseline hazard from the overall level of the random frailties.

Various frailty distributions have been proposed in the literature (Duchateau and Janssen 2008, Chapter 4). Hereinafter, we shall focus on the gamma, the positive stable, and the inverse Gaussian frailty distributions. In all of these three, a single heterogeneity parameter (denoted either θ or ν) indexes the degree of dependence. In the following, ξ is used as a generic notation to denote either θ or ν .

Data

For right-censored clustered survival data, the observation for subject $j \in J_i = \{1, \dots, n_i\}$ from cluster $i \in I = \{1, \dots, G\}$ is the couple $\mathbf{z}_{ij} = (y_{ij}, \delta_{ij})$, where $y_{ij} = \min(t_{ij}, c_{ij})$ is the minimum between the survival time t_{ij} and the censoring time c_{ij} , and where $\delta_{ij} = I(t_{ij} \leq c_{ij})$ is the event indicator. Covariate information may also have been collected; in this case, $\mathbf{z}_{ij} = (y_{ij}, \delta_{ij}, \mathbf{x}_{ij})$, where \mathbf{x}_{ij} denotes the vector of covariates for the ij -th observation. Further, if left-truncation is also present, truncation times τ_{ij} are gathered in the vector $\boldsymbol{\tau}$.

Likelihood

In the parametric setting, estimation is based on the marginal likelihood in which the frailties have been integrated out by averaging the conditional likelihood with respect to the frailty distribution. Under assumptions of non-informative right-censoring and of independence between the censoring time and the survival time random variables, given the covariate

information, the marginal log-likelihood of the observed data $\mathbf{z} = \{\mathbf{z}_{ij}; i \in I, j \in J_i\}$ can be written as (van den Berg and Drepper 2012)

$$\begin{aligned} \ell_{\text{marg}}(\boldsymbol{\psi}, \boldsymbol{\beta}, \boldsymbol{\xi}; \mathbf{z} \mid \boldsymbol{\tau}) = \sum_{i=1}^G \left\{ \left[\sum_{j=1}^{n_i} \delta_{ij} \left(\log(h_0(y_{ij})) + \mathbf{x}_{ij}^\top \boldsymbol{\beta} \right) \right] \right. \\ \left. + \log \left[(-1)^{d_i} \mathcal{L}^{(d_i)} \left(\sum_{j=1}^{n_i} H_0(y_{ij}) \exp(\mathbf{x}_{ij}^\top \boldsymbol{\beta}) \right) \right] \right. \\ \left. - \log \left[\mathcal{L} \left(\sum_{j=1}^{n_i} H_0(\tau_{ij}) \exp(\mathbf{x}_{ij}^\top \boldsymbol{\beta}) \right) \right] \right\}, \end{aligned} \quad (1)$$

with $d_i = \sum_{j=1}^{n_i} \delta_{ij}$ the number of events in the i -th cluster, and $\mathcal{L}^{(q)}(\cdot)$ the q -th derivative of the Laplace transform of the frailty distribution defined as

$$\mathcal{L}(s) = \mathbb{E}[\exp(-Us)] = \int_0^\infty \exp(-u_i s) f(u_i) du_i, \quad s \geq 0.$$

Estimation

Estimates of $\boldsymbol{\psi}$, $\boldsymbol{\beta}$, and $\boldsymbol{\xi}$ are obtained by maximising the marginal log-likelihood 1; this can easily be done if one is able to compute higher order derivatives $\mathcal{L}^{(q)}(\cdot)$ of the Laplace transform up to $q = \max\{d_1, \dots, d_G\}$. Symbolic differentiation might be performed in R, but is impractical here, mainly because this is very time consuming. Therefore, explicit formulas are rather desirable. Further, they will be used in the calculation of predictions as shown below.

Prediction

Besides parameter estimates, prediction of frailties are sometimes desirable. As an aside, they are needed at each expectation step of the expectation-maximisation (EM) algorithm that fits the semi-parametric frailty model.

The frailty term u_i can be predicted by $\hat{u}_i = \mathbb{E}(U \mid \mathbf{z}_i, \boldsymbol{\tau}_i; \hat{\boldsymbol{\psi}}, \hat{\boldsymbol{\beta}}, \hat{\boldsymbol{\xi}})$, with \mathbf{z}_i and $\boldsymbol{\tau}_i$ the data and the truncation times in cluster i . This conditional expectation can be computed as

$$\mathbb{E}(U \mid \mathbf{z}_i, \boldsymbol{\tau}_i; \boldsymbol{\psi}, \boldsymbol{\beta}, \boldsymbol{\xi}) = - \frac{\mathcal{L}^{(d_i+1)} \left(\sum_{j=1}^{n_i} H_0(y_{ij}) \exp(\mathbf{x}_{ij}^\top \boldsymbol{\beta}) \right)}{\mathcal{L}^{(d_i)} \left(\sum_{j=1}^{n_i} H_0(y_{ij}) \exp(\mathbf{x}_{ij}^\top \boldsymbol{\beta}) \right)},$$

which can be seen from Appendix A.2, together with $\mathbb{E}[U^q \exp(-Us)] = (-1)^q \mathcal{L}^{(q)}(s)$.

Outline

In Sections 2.1–2.3 we illustrate the three frailty distributions which are available in the **parfm** package: the gamma, the positive stable and the inverse Gaussian. Note that the Laplace transform of a lognormal random variable does not exist in a closed form. Hence, Equation 1 requires analytical or numerical approximation in that case, which is not considered here.

2.1. Gamma frailty

A gamma frailty term is a random variable $U \sim \text{Gam}^*(\theta)$ with probability density function

$$f(u) = \frac{\theta^{-\frac{1}{\theta}} u^{\frac{1}{\theta}-1} \exp(-u/\theta)}{\Gamma(1/\theta)}, \quad \theta > 0,$$

where $\Gamma(\cdot)$ is the gamma function. It corresponds to a gamma distribution $\text{Gam}(\mu, \theta)$ where the mean μ is fixed to 1 for identifiability. Its variance is then θ .

The associated Laplace transform is given by

$$\mathcal{L}(s) = (1 + \theta s)^{-\frac{1}{\theta}}, \quad s \geq 0,$$

and it is easy to show that, for $q \geq 1$,

$$\mathcal{L}^{(q)}(s) = (-1)^q (1 + \theta s)^{-q} \left[\prod_{l=0}^{q-1} (1 + l\theta) \right] \mathcal{L}(s).$$

Therefore, in Equation 1, we have

$$\log \left((-1)^q \mathcal{L}^{(q)}(s) \right) = - \left(q + \frac{1}{\theta} \right) \log(1 + \theta s) + \sum_{l=0}^{q-1} \log(1 + l\theta). \quad (2)$$

For the gamma distribution, the Kendall's tau (Hougaard 2000, Section 4.2), which measures the association between any two event times from the same cluster in the multivariate case, can be computed as

$$\tau = \frac{\theta}{\theta + 2} \in (0, 1).$$

2.2. Positive stable frailty

Hougaard (2000, Section A.3.3) introduces the positive stable distributions as a family with two parameters: a scale $\delta > 0$ and the so-called index $\alpha < 1$. Imposing $\delta = \alpha$, the positive stable frailty distribution $\text{PS}^*(\nu)$ is obtained, with $\nu = 1 - \alpha$.

The associated probability density function is then

$$f(u) = -\frac{1}{\pi u} \sum_{k=1}^{\infty} \frac{\Gamma(k(1-\nu) + 1)}{k!} (-u^{\nu-1})^k \sin((1-\nu)k\pi), \quad \nu \in (0, 1).$$

The mean and variance are both undefined. Therefore, the heterogeneity parameter ν does not correspond to the variance of the frailty term. Because of that, we intentionally call it ν instead of θ to avoid misinterpretation.

In contrast to the probability density function, the associated Laplace transform takes a very simple form,

$$\mathcal{L}(s) = \exp(-s^{1-\nu}), \quad s \geq 0,$$

and Wang, Klein, and Moeschberger (1995) found that, for $q \geq 1$,

$$\mathcal{L}^{(q)}(s) = (-1)^q ((1 - \nu)s^{-\nu})^q \left[\sum_{m=0}^{q-1} \Omega_{q,m} s^{-m(1-\nu)} \right] \mathcal{L}(s),$$

where the $\Omega_{q,m}$'s are polynomials of degree m , given recursively by

$$\begin{aligned} \Omega_{q,0} &= 1, \\ \Omega_{q,m} &= \Omega_{q-1,m} + \Omega_{q-1,m-1} \left\{ \frac{q-1}{1-\nu} - (q-m) \right\}, \quad m = 1, \dots, q-2, \\ \Omega_{q,q-1} &= (1-\nu)^{1-q} \frac{\Gamma(q - (1-\nu))}{\Gamma(\nu)}. \end{aligned} \quad (3)$$

It follows that

$$\begin{aligned} \log \left((-1)^q \mathcal{L}^{(q)}(s) \right) &= q (\log(1 - \nu) - \nu \log(s)) \\ &\quad + \log \left[\sum_{m=0}^{q-1} \Omega_{q,m} s^{-m(1-\nu)} \right] - s^{1-\nu}. \end{aligned} \quad (4)$$

With clustered data, the Kendall's tau for positive stable distributed frailties is

$$\tau = \nu \in (0, 1).$$

2.3. Inverse Gaussian frailty

The inverse Gaussian frailty distribution $\text{IG}^*(\theta)$ has density

$$f(u) = \frac{1}{\sqrt{2\pi\theta}} u^{-\frac{3}{2}} \exp \left(-\frac{(u-1)^2}{2\theta u} \right), \quad \theta > 0.$$

The mean and the variance are 1 and θ , respectively. For the Laplace transform, one has

$$\mathcal{L}(s) = \exp \left(\frac{1}{\theta} \left(1 - \sqrt{1 + 2\theta s} \right) \right), \quad s \geq 0,$$

and, for $q \geq 1$,

$$\mathcal{L}^{(q)}(s) = (-1)^q (2\theta s + 1)^{-\frac{q}{2}} \frac{K_{q-(1/2)} \left(\sqrt{2\theta^{-1}(s + \frac{1}{2\theta})} \right)}{K_{1/2} \left(\sqrt{2\theta^{-1}(s + \frac{1}{2\theta})} \right)} \mathcal{L}(s), \quad (5)$$

where K is the modified Bessel function of the second kind (Hougaard 2000, Section A.4.2)

$$K_\gamma(\omega) = \frac{1}{2} \int_0^\infty t^{\gamma-1} \exp \left\{ -\frac{\omega}{2} \left(t + \frac{1}{t} \right) \right\} dt, \quad \gamma \in \mathbb{R}, \omega > 0.$$

The proof of this result, given in Appendix A.1, sketches a general constructive method to obtain the derivatives of the Laplace transform for any distribution for which the moments of $U \mid \mathbf{z}_i, \boldsymbol{\tau}_i; \boldsymbol{\psi}, \boldsymbol{\beta}, \xi$, the conditional frailty given the data, are known.

Noting that $K_{1/2}(\omega) = \sqrt{\frac{\pi}{2\omega}} \exp(-\omega)$, we have

$$\log\left((-1)^q \mathcal{L}^{(q)}(s)\right) = -\frac{q}{2} \log(2\theta s + 1) + \log\left(K_{q-(1/2)}(z)\right) - \left[\frac{1}{2} \left(\log\left(\frac{\pi}{2z}\right)\right) - z\right] + \frac{1}{\theta} \left(1 - \sqrt{1 + 2\theta s}\right), \quad (6)$$

with $z = \sqrt{2\theta^{-1}(s + \frac{1}{2\theta})}$.

With multivariate data, an inverse Gaussian distributed frailty yields a Kendall's tau given by

$$\tau = \frac{1}{2} - \frac{1}{\theta} + 2 \frac{\exp(2/\theta)}{\theta^2} \int_{2/\theta}^{\infty} \frac{\exp(-u)}{u} du \in (0, 1/2).$$

3. Case study

We illustrate the **parfm** package with the very well-known kidney dataset that contains the recurrence times to kidney infection for 38 patients using portable dialysis equipment (McGilchrist and Aisbett 1991).

```
R> R.Version()[["version.string"]]
```

```
[1] "R version 2.15.1 (2012-06-22)"
```

```
R> library("parfm")
```

```
R> packageDescription("parfm", fields = "Version")
```

```
[1] "2.5.2"
```

The dataset is available in **parfm** via the command `data("kidney")` and it looks like the following:

```
R> head(kidney)
```

	id	time	status	age	sex	disease	frail
1	1	8	1	28	1	Other	2.3
2	1	16	1	28	1	Other	2.3
3	2	23	1	48	2	GN	1.9
4	2	13	0	48	2	GN	1.9
5	3	22	1	32	1	Other	1.2
6	3	28	1	32	1	Other	1.2

Each observation corresponds to a kidney, the variable `id` being the patient's code. The time from insertion of the catheter to infection or censoring is stored in `time` while `status` is 1 when infection has occurred and 0 for censored observations (catheters may be removed for reasons other than infection). Three covariates are available: `age`, the age of the patient in years, `sex`, being 1 for males and 2 for females, and `disease`, the disease type (GN, AN,

PKD or Other). Finally `frail` is the frailty prediction from the original paper which fits a semi-parametric lognormal frailty model.

First and foremost, `sex` is recoded as a 0/1 indicator for ease of interpretation:

```
R> kidney$sex <- kidney$sex - 1
```

The hazard of infection will be modelled as a function of the patient's age and sex. Clearly, kidneys from the same patient cannot be considered independent. Therefore, the use of a shared frailty model is advisable, with clusters of size 2 corresponding to patients.

The `parfm()` function must have the following inputs. `formula`: a formula with an object of class `Surv` on the left-hand side; `cluster`: the cluster variable's name; `data`: the dataset; `dist`: the baseline hazard, either `exponential`, `weibull`, `gompertz`, `lognormal` or `loglogistic`; `frailty`: the frailty distribution, either `none`, `gamma`, `possta` or `ingau`.

Model estimation

The model with exponential baseline hazard and gamma frailty distribution is first fitted.

```
R> mod <- parfm(Surv(time, status) ~ sex + age, cluster = "id",
+ data = kidney, dist = "exponential", frailty = "gamma")
R> mod
```

Execution time: 1.15 second(s)

Frailty distribution: Gamma
 Basline hazard distribution: Exponential
 Loglikelihood: -333.248

	ESTIMATE	SE	p-val
theta	0.301	0.157	
lambda	0.025	0.015	
sex	-1.485	0.398	0.000 ***
age	0.005	0.011	0.663

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Kendall's Tau: 0.131

Standard errors are computed as the square roots of the diagonal elements of the observed information matrix. According to this model, `sex` has a significant impact on the hazard of infection while it is not affected by `age`. Conditional on the patient's frailty and on the age, the hazard of infection for a female at any time t is estimated to be $\exp(-1.485) \approx 0.227$ times that of a male, with Wald confidence interval

```
R> ci.parfm(mod, level = 0.05)["sex", ]
```

	low	up
	0.104	0.495

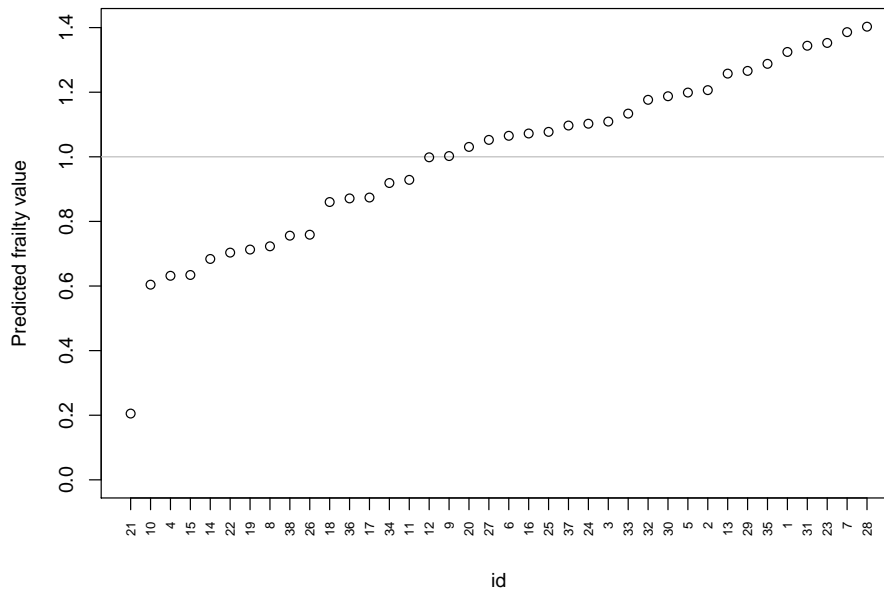


Figure 1: Prediction of frailties for the kidney dataset as given by the parametric gamma-exponential frailty model.

As for the heterogeneity parameter, it is estimated to be 0.301 which corresponds to a Kendall's tau equal to 0.131.

Frailty prediction

Prediction of frailties can be obtained via the `predict()` function, with the parametric frailty model object as unique argument. For instance, the predictions for the gamma-exponential model, `mod`, are obtained via the command

```
R> u <- predict(mod)
```

which returns an object of class `predict.parfm`. These predictions can easily be plotted (Figure 1) with the command `plot(u, sort = "i")`.

Comparison of different models

In some circumstances, it might be useful to easily obtain AIC and BIC values for a series of candidate models. This can be done using the `select.parfm()` function. Its use is similar to that of the `parfm()` function, but the `dist` and `frailty` values are vectors that contain all the alternatives to try.

```
R> kidney.parfm <- select.parfm(Surv(time, status) ~ sex + age,
+   cluster = "id", data = kidney, dist = c("exponential", "weibull",
+     "gompertz", "loglogistic", "lognormal"),
+   frailty = c("gamma", "ingau", "possta"))
R> kidney.parfm
```

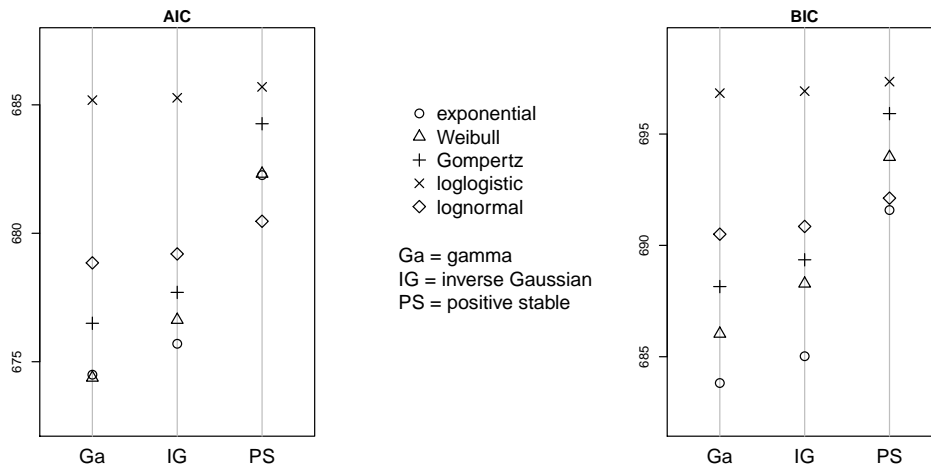


Figure 2: AIC and BIC values of **parfm** models for the kidney dataset.

AIC:

	gamma	ingau	possta
exponential	674.496	675.699	682.264
weibull	674.376	676.627	682.315
gompertz	676.496	677.699	684.264
loglogistic	685.184	685.274	685.699
lognormal	678.849	679.196	680.467

BIC:

	gamma	ingau	possta
exponential	683.819	685.022	691.587
weibull	686.029	688.281	693.969
gompertz	688.150	689.353	695.918
loglogistic	696.837	696.927	697.353
lognormal	690.502	690.850	692.121

The results can be plotted (Figure 2) via the command `plot(kidney.parfm)`. In this particular example, the exponential baseline seems to be a good candidate.

As a comparison, the model with inverse Gaussian distributed frailties is fitted by changing the frailty argument into "ingau".

```
R> parfm(Surv(time, status) ~ sex + age, cluster = "id",
+ data = kidney, dist = "exponential", frailty = "ingau")
```

Execution time: 1.15 second(s)

```
Frailty distribution: Inverse Gaussian
Baseline hazard distribution: Exponential
Loglikelihood: -333.85
```

```

      ESTIMATE SE    p-val
theta  0.375  0.259
lambda 0.022  0.013
sex    -1.310  0.373 0.000 ***
age     0.004  0.011 0.694
---
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Kendall's Tau: 0.125

```

In this case, the conclusions drawn from the previous two models are essentially analogous. Consider now the model with the positive stable frailty distribution. In this example, it converges to a solution which is not valid ($\nu = 0$) with the default settings.

```
R> parfm(Surv(time, status) ~ sex + age, cluster = "id",
+ data = kidney, dist = "exponential", frailty = "possta")
```

Execution time: 1.16 second(s)

```

Frailty distribution: Positive Stable
Baseline hazard distribution: Exponential
Loglikelihood: -337.132

```

```

      ESTIMATE SE p-val
nu      0.000
lambda  0.012
sex     -0.885
age      0.004
---
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

Kendall's Tau: 0

Warning message:

```

In parfm(Surv(time, status) ~ sex + age, cluster = "id", data = kidney, :
  Error in solve.default(res$hessian) :
  Lapack routine dgesv: system is exactly singular

```

The default initial value for ν is $1/2$ in the case of positive stable frailties; it can be changed by means of the `iniFpar` option in `parfm()`. Let us try with $\nu = 0.25$.

```
R> parfm(Surv(time, status) ~ sex + age, cluster = "id",
+ data = kidney, dist = "exponential", frailty = "possta", iniFpar = 0.25)
```

Execution time: 1.71 second(s)

```

Frailty distribution: Positive Stable
Baseline hazard distribution: Exponential

```

Loglikelihood: -336.182

	ESTIMATE	SE	p-val
nu	0.112	0.084	
lambda	0.014	0.008	
sex	-0.951	0.348	0.006 **
age	0.004	0.011	0.698

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Kendall's Tau: 0.112

The problem might also be fixed by changing the optimisation method (see `optim()`). By default it is set to "BFGS", but it can be changed through the `method` option.

```
R> parfm(Surv(time, status) ~ sex + age, cluster = "id",
+ data = kidney, dist = "exponential", frailty = "possta",
+ method = "Nelder-Mead")
```

Execution time: 1.51 second(s)

Frailty distribution: Positive Stable
 Basline hazard distribution: Exponential
 Loglikelihood: -336.182

	ESTIMATE	SE	p-val
nu	0.112	0.084	
lambda	0.014	0.008	
sex	-0.951	0.348	0.006 **
age	0.004	0.011	0.694

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Kendall's Tau: 0.112

In this example the results obtained by changing the optimisation method are the same as those obtained by changing the initial value of ν . When convergence problems occur, using different starting values and/or different optimisation methods is generally sufficient to find the global maximum of the marginal likelihood function.

Finally we provide a comparison with the semi-parametric model. As an example, we fit the semi-parametric model with gamma frailties via the `coxph()` function.

```
R> coxph(Surv(time, status) ~ sex + age +
+ frailty(id, distribution = "gamma", eps = 1e-11),
+ outer.max = 50, data = kidney)
```

	coef	se(coef)	se2	Chisq	DF	p
sex	-1.58323	0.4594	0.3515	11.88	1.0	0.00057

```
age                0.00522 0.0119   0.0088  0.19  1.0 0.66000
frailty(id, distribution          22.96 12.9 0.04100
```

```
Variance of random effect= 0.408   I-likelihood = -181.6
```

Estimates of regression parameters are quite similar to those of the exponential–gamma model, while the frailty variance is sensibly different, arguably because of the difference in how the baseline hazard is treated.

4. Discussion

To the best of our knowledge, parametric frailty models are currently especially handled in *Stata* by means of the `streg` command. With **parfm**, they are now readily fitted in *R*. Further, **parfm** provides the positive stable frailty distribution which is presently unavailable in *Stata*. Actually, except for a *SAS* macro, `ps_frail`, developed by [Shu and Klein \(1999\)](#) in the semi-parametric setting, we are not aware of another package that provides the positive stable frailty distribution.

The **parfm** package is flexible and easy to use. It provides five distributions for the baseline hazard and three frailty distributions. Parameter estimation is done by maximising the marginal log-likelihood given in Equation 1. The `optim()` function is employed, and its `method` option is passed to `parfm()` (with `method = "BFGS"` by default). If not specified in the `inip` option, initial values for all but the heterogeneity parameter are obtained by fitting an unadjusted (i.e., without frailty) parametric proportional hazards model using the `phreg()` function from the **eha** package ([Broström 2012](#)). The initial heterogeneity parameter can also be specified by the user via the `iniFpar` option; otherwise it is set to 1 when frailties follow a gamma or an inverse Gaussian distribution, or to 1/2 when they follow the positive stable distribution.

Additionally, when `frailty = "none"`, `parfm()` fits the unadjusted parametric proportional hazards model, similar to `survreg()` (from the **survival** package) or to `phreg()`. However, `survreg()` returns the parameter estimates in the log-linear model and `phreg()` uses yet another parametrisation (see the documentation). Often, the user has then to transform back the parameters and to employ the delta method in order to get estimates for the standard errors. The `parfm()` function directly uses the proportional hazards representation.

Nonetheless, **parfm** might reach its limits when at least one d_i , the number of events in cluster i , $i \in \{1, \dots, G\}$, is very large. First, consider the positive stable distribution and observe that, for a fixed value of $m \in \{1, \dots, q - 1\}$, $\Omega_{q,m}$ rapidly grows as q increases; see Equations 3. At the extreme, some of them might exceed the largest representable number in *R*. These are then stored as `Inf`. This, in turn, prevents the marginal log-likelihood in Equation 1 to be evaluated and hence maximised. On a side note, also the *SAS* macro `ps_frail` that implements the EM algorithm to fit the semi-parametric positive stable frailty model has analogous difficulties when the number of events is large (or even moderate). The following ad-hoc solution is implemented in **parfm**: in order to keep the polynomials $\Omega_{q,m}$'s reasonably small, they are divided by some factor 10^K which does not change the marginal log-likelihood except for an additive constant (which equals $-G \times K \times \log(10)$). The value of K is specified via the `correct` option (default is `correct=0`, i.e., no correction) and `parfm()` returns the re-adjusted log-likelihood value. That solution serves the purpose for moderately large values

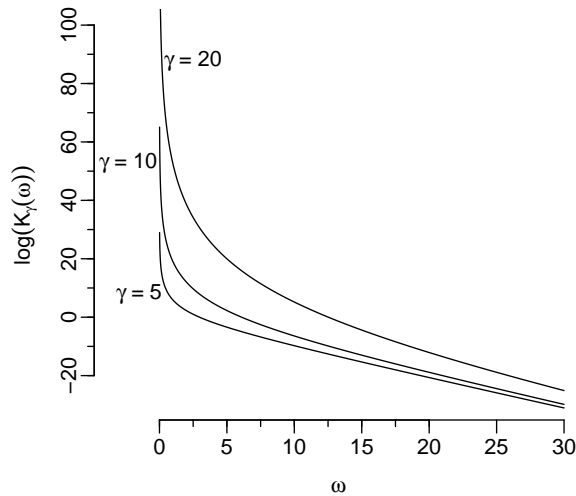


Figure 3: The logarithm of the Bessel function, $\log(K_\gamma(\omega))$, versus ω for different values of γ .

of d_i (say up to about 200 events per cluster according to our experience, but it depends on the data, on the other parameters, and on the hardware characteristics). With the inverse Gaussian distribution, the Bessel function $K_{q-1/2}(z)$ in Equation 6 raises the same problem. Indeed, it explodes when z is small relative to q ; see Figure 3. Currently, that distribution should, therefore, preferably be avoided when there are very large values of d_i (say above 200 events per cluster according to our experience, but, again, it depends on the data, on the other parameters and on the hardware characteristics). Moreover, $K_{q-1/2}(z)$ rapidly goes to zero as z increases. So, in case of very small apparent heterogeneity, $\theta \rightarrow 0$ which implies $z \rightarrow \infty$, $K_{q-1/2}(z)$ might be stored as 0 in R and hence $\log(K_{q-1/2}(z))$ cannot be computed. However, as this problem occurs in the case of very small heterogeneity, this would rather suggest to fit the model with `frailty = "none"`. When frailties are gamma distributed, which is by far the most popular assumption in common practice, the quantities involved in Equation 2 do not raise any worry. In practice, even when dealing with datasets with huge numbers of events per cluster, there is no real risk of exceeding the range of floating-point numbers.

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A. Proofs

A.1. Derivatives of the Laplace transform of the inverse Gaussian frailty

On the one hand, for any frailty distribution $f(u_i; \xi)$, the α -th moment of U ($\alpha \in \mathbb{N}$), conditional on the data from the i -th cluster and on the parameters, can be written in the form

$$\mathbb{E}(U^\alpha \mid \mathbf{z}_i, \boldsymbol{\tau}_i; \boldsymbol{\psi}, \boldsymbol{\beta}, \xi) = \frac{\mathbb{E}(U^{d_i+\alpha} \exp(-UH_{i,c}(\mathbf{y}_i)))}{\mathbb{E}(U^{d_i} \exp(-UH_{i,c}(\mathbf{y}_i)))}, \quad (7)$$

with $H_{i,c}(\mathbf{y}_i) = \sum_{j=1}^{n_i} H_0(y_{ij}) \exp(\mathbf{x}_{ij}^\top \boldsymbol{\beta})$. This is a generalisation of a result found by Wang *et al.* (1995) which follows from Bayes's formula applied to $f(u_i \mid \mathbf{z}_i, \boldsymbol{\tau}_i; \boldsymbol{\psi}, \boldsymbol{\beta}, \xi)$ in

$$\mathbb{E}(U^\alpha \mid \mathbf{z}_i, \boldsymbol{\tau}_i; \boldsymbol{\psi}, \boldsymbol{\beta}, \xi) = \int_0^\infty u_i^\alpha f(u_i \mid \mathbf{z}_i, \boldsymbol{\tau}_i; \boldsymbol{\psi}, \boldsymbol{\beta}, \xi) du_i$$

(see Appendix A.2 for more details). Now, since the expected values in the right-hand side of Equation 7 can be written in terms of derivatives of the Laplace transform

$$\mathbb{E}(U^q \exp(-sU)) = (-1)^q \mathcal{L}^{(q)}(s), \quad q, s \geq 0,$$

we have that

$$\mathcal{L}^{(d_i+\alpha)}(H_{i,c}(\mathbf{y}_i)) = (-1)^\alpha \mathbb{E}(U^\alpha \mid \mathbf{z}_i, \boldsymbol{\tau}_i; \boldsymbol{\psi}, \boldsymbol{\beta}, \xi) \mathcal{L}^{(d_i)}(H_{i,c}(\mathbf{y}_i)). \quad (8)$$

On the other hand, if $U \sim \text{IG}^*(\theta)$, then it is easy to show that the conditional distribution of U given the data and the parameters is a generalised inverse Gaussian distribution (Appendix A.3):

$$U \mid \mathbf{z}_i, \boldsymbol{\tau}_i; \boldsymbol{\psi}, \boldsymbol{\beta}, \theta \sim \text{GIG}(\gamma_{\text{GIG}}, \delta_{\text{GIG}}, \theta_{\text{GIG}})$$

with

$$\gamma_{\text{GIG}} = d_i - \frac{1}{2}, \quad (9)$$

$$\theta_{\text{GIG}} = \frac{1}{2\theta} + H_{i,c}(\mathbf{y}_i), \quad (10)$$

$$\delta_{\text{GIG}} = \frac{1}{\sqrt{2\theta}}. \quad (11)$$

Hence (Hougaard 2000, Section A.3.6)

$$\mathbb{E}(U^\alpha \mid \mathbf{z}_i, \boldsymbol{\tau}_i; \boldsymbol{\psi}, \boldsymbol{\beta}, \xi) = \left(\frac{\theta_{\text{GIG}}^{1/2}}{\delta_{\text{GIG}}} \right)^{-\alpha} \frac{K_{\gamma_{\text{GIG}}+\alpha}(2\delta_{\text{GIG}} \theta_{\text{GIG}}^{1/2})}{K_{\gamma_{\text{GIG}}}(2\delta_{\text{GIG}} \theta_{\text{GIG}}^{1/2})}. \quad (12)$$

Combining (8) and (12), Equation 5 is deduced.

A.2. Conditional expectation of frailty terms

For ease of notation, let $H_{i,c}(\mathbf{y}_i)$ denote $\sum_{j=1}^{n_i} H_0(y_{ij}) \exp(\mathbf{x}_{ij}^\top \boldsymbol{\beta})$.

For any frailty distribution $f(u_i; \xi)$ and for any $\alpha \in \mathbb{N}$, we have

$$\begin{aligned} \mathbb{E}(U^\alpha \mid \mathbf{z}_i, \boldsymbol{\tau}_i; \boldsymbol{\psi}, \boldsymbol{\beta}, \xi) &= \int_0^\infty u_i^\alpha f(u_i \mid \mathbf{z}_i, \boldsymbol{\tau}_i; \boldsymbol{\psi}, \boldsymbol{\beta}, \xi) du_i \\ &= \int_0^\infty u_i^\alpha \frac{L_{\text{cond}}(\boldsymbol{\psi}, \boldsymbol{\beta}; \mathbf{z}_i \mid \boldsymbol{\tau}_i, u_i) f(u_i \mid \boldsymbol{\tau}_i; \xi)}{L_{\text{marg}}(\boldsymbol{\psi}, \boldsymbol{\beta}, \xi; \mathbf{z}_i \mid \boldsymbol{\tau}_i)} du_i, \end{aligned}$$

with

$$\begin{aligned} L_{\text{cond}}(\boldsymbol{\psi}, \boldsymbol{\beta}; \mathbf{z}_i \mid \boldsymbol{\tau}_i, u_i) &= \left[\prod_{j=1}^{n_i} \left(h_0(y_{ij}) u_i \exp(\mathbf{x}_{ij}^\top \boldsymbol{\beta}) \right)^{\delta_{ij}} \right] \\ &\quad \times \exp(-u_i H_{i,c}(\mathbf{y}_i)) \exp(u_i H_{i,c}(\boldsymbol{\tau}_i)), \\ f(u_i \mid \boldsymbol{\tau}_i; \xi) &= \frac{\exp(-u_i H_{i,c}(\boldsymbol{\tau}_i)) f(u_i; \xi)}{\mathcal{L}(H_{i,c}(\boldsymbol{\tau}_i))}, \\ L_{\text{marg}}(\boldsymbol{\psi}, \boldsymbol{\beta}, \xi; \mathbf{z}_i \mid \boldsymbol{\tau}_i) &= \int_0^\infty L_{\text{cond}}(\boldsymbol{\psi}, \boldsymbol{\beta} \mid \boldsymbol{\tau}_i, u_i; \mathbf{z}_i) f(u_i \mid \boldsymbol{\tau}_i; \xi) du_i. \end{aligned}$$

Thus,

$$\begin{aligned} \mathbb{E}(U^\alpha \mid \mathbf{z}_i, \boldsymbol{\tau}_i; \boldsymbol{\psi}, \boldsymbol{\beta}, \xi) &= \frac{\int_0^\infty u_i^{d_i + \alpha} \exp(-u_i H_{i,c}(\mathbf{y}_i)) f(u_i; \xi) du_i}{\int_0^\infty u_i^{d_i} \exp(-u_i H_{i,c}(\mathbf{y}_i)) f(u_i; \xi) du_i} \\ &= \frac{\mathbb{E}\left[U^{d_i + \alpha} \exp(-U H_{i,c}(\mathbf{y}_i))\right]}{\mathbb{E}\left[U^{d_i} \exp(-U H_{i,c}(\mathbf{y}_i))\right]}. \end{aligned}$$

A.3. Conditional distribution of inverse Gaussian frailty

Let $H_{i,c}(\mathbf{y}_i) = \sum_{j=1}^{n_i} H_0(y_{ij}) \exp(\mathbf{x}_{ij}^\top \boldsymbol{\beta})$.

If $U \sim \text{IG}^*(\theta)$, then

$$\begin{aligned} f(u_i \mid \mathbf{z}_i, \boldsymbol{\tau}_i; \boldsymbol{\psi}, \boldsymbol{\beta}, \theta) &= \frac{L_{\text{cond}}(\boldsymbol{\psi}, \boldsymbol{\beta}; \mathbf{z}_i \mid \boldsymbol{\tau}_i, u_i) f(u_i \mid \boldsymbol{\tau}_i; \theta)}{L_{\text{marg}}(\boldsymbol{\psi}, \boldsymbol{\beta}, \theta; \mathbf{z}_i \mid \boldsymbol{\tau}_i)} \\ &\propto L_{\text{cond}}(\boldsymbol{\psi}, \boldsymbol{\beta}; \mathbf{z}_i \mid \boldsymbol{\tau}_i, u_i) f(u_i \mid \boldsymbol{\tau}_i; \theta) \\ &\propto u_i^{d_i} \exp(-u_i H_{i,c}(\mathbf{y}_i)) \\ &\quad \times \sqrt{\frac{1}{2\pi\theta}} u_i^{-\frac{3}{2}} \exp\left(-\frac{1}{2\theta u_i} (u_i - 1)^2\right) \\ &\propto u_i^{d_i - \frac{3}{2}} \exp\left(-u_i H_{i,c}(\mathbf{y}_i) - \frac{1}{2\theta} u_i - \frac{1}{2\theta} \frac{1}{u_i}\right) \\ &= u_i^{d_i - \frac{3}{2}} \exp\left(-\left(\frac{1}{2\theta} + H_{i,c}(\mathbf{y}_i)\right) u_i - \frac{1}{2\theta} \frac{1}{u_i}\right), \end{aligned}$$

which is proportional to the density of a generalised inverse Gaussian distribution (Hougaard 2000, Section A.3.6) with parameters given by Equations 9–11.

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