

A wrapper routine for MLE estimation in parametric regression models for survival analysis in R

`maxlogLreg`

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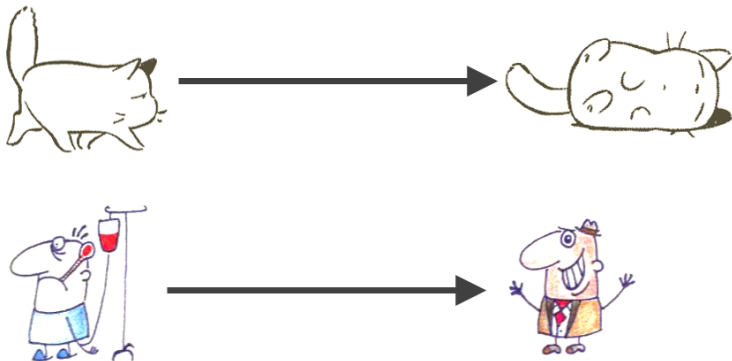
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Section 1

Survival Analysis

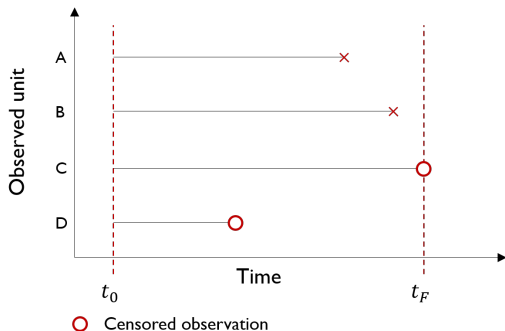
About Survival Analysis



One directional transitions, from A to B (irreversible).

Assumptions

- Observed events.
- Non observed events:
 - Right censorship (mainly).
 - Left censorship (also available).



Linear regression models

A linear regression model can be expressed as follows:

$$y_i \stackrel{\text{iid.}}{\sim} \mathcal{D}(\boldsymbol{\theta}) \quad (1)$$

$$\theta_j = \mathbf{X}_j^\top \boldsymbol{\beta}_j, \quad (2)$$

- y_i is a response (random) variable with $i = 1, 2, \dots, n$ observations.
- \mathcal{D} a distribution with k parameters.
- $\boldsymbol{\theta} = (\theta_1, \dots, \theta_k)$ is the vector of the distribution parameters.
- $\boldsymbol{\beta}_j = (\beta_1, \beta_2, \dots, \beta_k)$ are the fixed effects vector of covariates of j^{th} distribution parameter.
- \mathbf{X}_j is a known design matrix of order $n \times k$.

Section 2

Estimation

Mathematical principle behind estimation in this models:

$$\hat{\beta}_j = \arg \max_{\beta_j \in B} \left\{ \log \left[\prod_{i=1}^n f(y_i)^{a_i} S(y_i)^{r_i} F(y_i)^{l_i} \right] \right\}, \quad (3)$$

$f(\cdot)$, $F(\cdot)$ and $S(\cdot)$ are de probability density, cumulative density and survival functions.

$$a_i = \begin{cases} 1, & \text{if } y_i \text{ is not censored.} \\ 0, & \text{in other case} \end{cases} \quad (4)$$

and so on for left and right censorship with r_i and l_i respectively.

Section 3

Our routine

How to get it?

- Download and install EstimationTools package.

```
if (!require('devtools')) install.packages('devtools')
devtools::install_github('Jaimemosg/EstimationTools',
                        force = TRUE)
library(EstimationTools)
```

- Repo in GitHub: <https://github.com/Jaimemosg/EstimationTools>
- GitHub web site: <https://jaimemosg.github.io/EstimationTools/>
- CRAN repo:
<https://cran.r-project.org/web/packages/EstimationTools/index.html>

build passing build passing CRAN 1.2.1 · 12 days ago repo status Active downloads 28k/month licence GPL-3

EstimationTools

The goal of `EstimationTools` is to provide a routine for parameter estimation of probability density/mass functions in `R`.

Installation

You can install the latest version of `EstimationTools` typing the following command lines in `R` console:

```
if (!require('devtools')) install.packages('devtools')
devtools::install_github('Jaimenos/EstimationTools', force = TRUE)
library(EstimationTools)
```

Or you can install the released version from [CRAN](#) if you prefer. You can also type the following command lines in `R` console:

```
install.packages("EstimationTools")
```

You can visit the [package website](#) to explore the vignettes (articles) and functions reference.



EstimationTools **2.0.0**



Reference

Articles ▾

Reference

All functions

Fibers

`logit_link()`

`log_link()`

`maxlogi()`

`maxlogi.reg()`

`negInv_Link()`

`summary(<maxlogi>)`

Tensile strengths

Logit link function (for estimation with `maxlogi` object)

Logarithmic link function (for estimation with `maxlogi` object)

Maximum Likelihood Estimation for parametric distributions

Maximum Likelihood Estimation for parametric linear regression models

Negative inverse link function (for estimation with `maxlogi` object)

Summarize Maximum Likelihood Estimation

Main arguments

```
maxlogLreg(formulas, y_dist, data, fixed, link,  
            start, lower, upper, ...)
```

- `formulas`: a list with formula objects to specify linear predictors of each parameter.
- `y_dist`: a formula to define the response variable and its distribution.
- `fixed`: a list stating the known parameters.
- `link`: a list to define link functions application.
- `start`, `lower`, `upper`: lists to set out initial values, lower and upper bounds for search.

Section 4

Motivation: usual approach vs. `maxlogLreg`

Application example # 1

Lets assume a failure test of 500 hours of 20 electronic devices. Only 10 of these failures have been observed (those with *Status* = 1).

Table 1: Failure times in the test.

Time	Status	Time	Status
55	1	500	0
187	1	500	0
216	1	500	0
240	1	500	0
244	1	500	0
335	1	500	0
361	1	500	0
373	1	500	0
375	1	500	0
386	1	500	0

Application example # 1

Selected distribution:

$$f(y|\alpha, k) = \frac{\alpha}{k} \left(\frac{y}{k}\right)^{\alpha-1} \exp \left[- \left(\frac{y}{k}\right)^{\alpha} \right]$$

The model:

$$y_i \stackrel{\text{iid.}}{\sim} WEI(\alpha, k),$$

$$\alpha = c_1,$$

$$k = c_2,$$

c_1 and c_2 are unknown parameters.

Application example # 1

Code implementation:

```
logL <- function(theta){  
  beta <- theta[1]  
  eta <- theta[2]  
  logf <- dweibull(x = y[,1], shape = beta, scale = eta,  
                  log = TRUE)  
  logS <- log(pweibull(q = y[,1], shape = beta, scale = eta,  
                     lower.tail = FALSE))  
  l <- sum( logf*y[,2] + logS*(1 - y[,2]) )  
  return(-l)  
}  
fit <- nlminb(start = c(1,100), objective = logL, lower = c(0,0))  
fit$par
```

```
## [1] 1.725626 606.004873
```

With our routine

```
formulas <- list(scale.fo = ~ 1, shape.fo = ~ 1)

start <- list(
  scale = list(Intercept = 100),
  shape = list(Intercept = 10)
)

lower <- list(
  scale = list(Intercept = 0),
  shape = list(Intercept = 0)
)

mod_weibull <- maxlogLreg(formulas,
  y_dist = Surv(times, status) ~ dweibull,
  start = start,
  lower = lower)
```

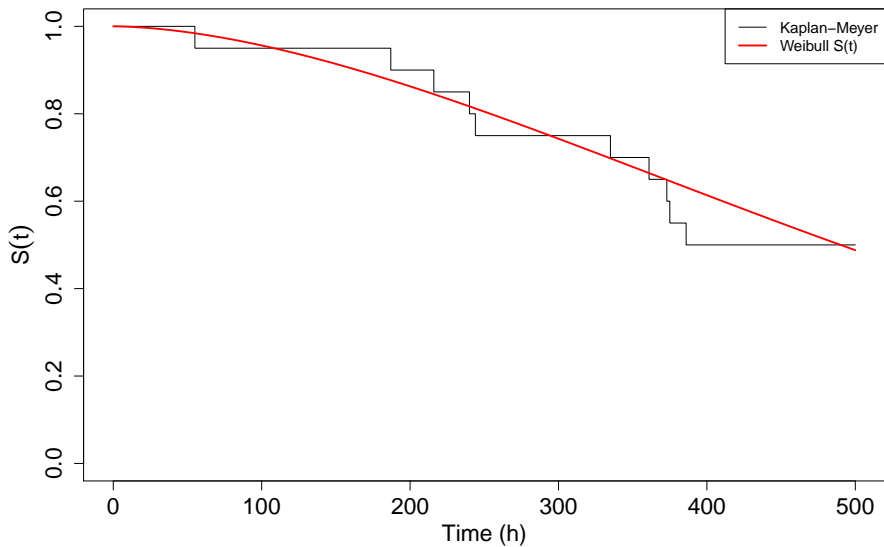

With our routine

The user does not have to write loglikelihood again and again.

```
summary(mod_weibull)
```

```
## -----  
## Optimization routine: nlminb  
## Standard Error calculation: Hessian from optim  
## -----  
##           AIC       BIC  
##    154.2437 156.2352  
## -----  
## Fixed effects for g(shape)  
## -----  
##           Estimate Std. Error Z value Pr(>|z|)  
## (Intercept)   1.7256    0.5034  3.4279 0.0006083 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
## -----  
## Fixed effects for g(scale)  
## -----  
##           Estimate Std. Error Z value Pr(>|z|)  
## (Intercept)   606.00    124.43  4.8703 1.114e-06 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
## -----  
## Note: p-values valid under asymptotic normality of estimators  
## ---
```

The fit



Section 5

Implementation of a new regression model

Application example # 2

Neck cancer data

The data corresponds to a randomized clinical trial to compare two therapies for head and neck cancer (Khan 2018; Efron 1988):

- i. 51 patients treated with radiation only.
- ii. 45 patients treated with radiation plus chemotherapy.
- iii. Response: survival times in days.

We fitted the Weibull model (usually used) and Exponentiated Weibull (EW) model.

The regression models

- Weibull:

$$f(t|\rho, \kappa) = \kappa \rho (\rho t)^{\kappa-1} \exp[-(\rho t)^\kappa]$$

$$T_i \stackrel{\text{iid.}}{\sim} \text{WEI2}(\rho, \kappa),$$

$$\log(\rho) = \beta_0 + \beta_1 x_1,$$

$$\log(\kappa) = \alpha_0,$$

- Exponentiated Weibull:

$$g(t|\rho, \kappa) = \kappa \rho \gamma (\rho t)^{\kappa-1} \exp[-(\rho t)^\kappa] \{1 - \exp[-(\rho t)^\kappa]\}$$

$$T_i \stackrel{\text{iid.}}{\sim} \text{EW}(\rho, \kappa, \gamma),$$

$$\log(\rho) = \beta_0 + \beta_1 x_1,$$

$$\log(\kappa) = \alpha_0,$$

$$\log(\gamma) = \lambda_0,$$

Weibull regression

```
## -----
## Optimization routine: nlminb
## Standard Error calculation: Hessian from optim
## -----
##           AIC       BIC
##    1082.519 1090.212
## -----
## Fixed effects for g(rho)
## -----
##           Estimate Std. Error  Z value  Pr(>|z|)
## (Intercept) -6.82473    0.21110 -32.3294 < 2.2e-16 ***
## Therapy      0.78603    0.27880   2.8193  0.004813 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## -----
## Fixed effects for g(k)
## -----
##           Estimate Std. Error  Z value  Pr(>|z|)
## (Intercept) -0.16173    0.09210  -1.756   0.07909 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## -----
## Note: p-values valid under asymptotic normality of estimators
## ---
```

Estimates by Khan: $\hat{\beta}_0 = -6.825$, $\hat{\beta}_1 = 0.786$, $\hat{\alpha}_0 = 0.162$

EW regression

Implement the distribution

```
dEW <- function(x, rho, k, gam, log = FALSE) {  
  prod1 <- -(rho*x)^k  
  prod2 <- (rho*x)^(k-1)  
  f <- k*rho*gam*prod2*exp(prod1)*( -expm1(prod1) )^(gam-1)  
  if (log == FALSE)  
    density <- f  
  else density <- log(f)  
  return(density)  
}  
  
pEW <- function(q, rho, k, gam, lower.tail = TRUE, log.p = FALSE) {  
  prod <- -(rho*q)^k  
  cdf <- (-expm1(prod))^gam  
  if (lower.tail == TRUE)  
    cdf <- cdf  
  else cdf <- 1 - cdf  
  if (log.p == FALSE)  
    cdf <- cdf  
  else cdf <- log(cdf)  
  cdf  
}
```

EW regression

```
formulas <- list(rho.fo = ~ Therapy, k.fo = ~ 1, gam.fo = ~ 1)

start <- list(
  rho = list(Intercept = 7, Therapy = 2),
  k = list(Intercept = -4), gam = list(Intercept = 4)
)
lower <- list(
  rho = list(Intercept = 0, Therapy = 0),
  k = list(Intercept = -6), gam = list(Intercept = 0)
)
upper <- list(
  rho = list(Intercept = 20, Therapy = 5),
  k = list(Intercept = 5), gam = list(Intercept = 10)
)

control <- list(step.max = 1e-5, step.min=1e-7, eval.max=500,
               iter.max=500, rel.tol=1e-8)

modEW <- maxloglreg(formulas, start = start, control = control,
                   lower = lower, upper = upper,
                   y_dist = Surv(Time, status) ~ dEW,
                   link = list(over = c("rho", "k", "gam"),
                               fun = rep("log_link", 3)))
```


EW regression

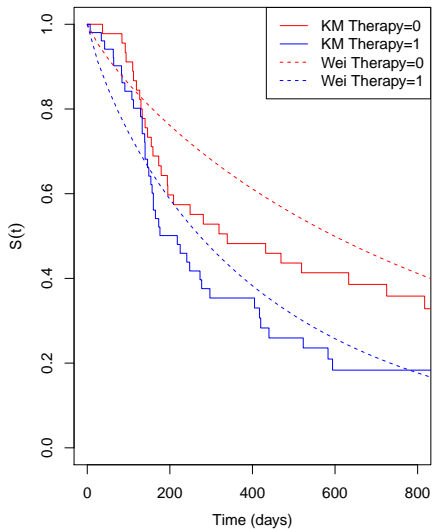
```
summary(modEW)
```

```
## -----
## Optimization routine: nlminb
## Standard Error calculation: Hessian from optim
## -----
##           AIC       BIC
##    1064.556 1074.813
## -----
## Fixed effects for g(rho)
## -----
##           Estimate Std. Error Z value Pr(>|z|)
## (Intercept) 10.17474   15.33130  0.6637  0.50688
## Therapy      0.56092    0.26730  2.0985  0.03586 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## -----
## Fixed effects for g(k)
## -----
##           Estimate Std. Error Z value Pr(>|z|)
## (Intercept) -2.1157     0.6429 -3.2908 0.000999 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## -----
## Fixed effects for g(gam)
## -----
##           Estimate Std. Error Z value Pr(>|z|)
## (Intercept)   6.7060     4.1885  1.6011  0.1094
## -----
## Note: p-values valid under asymptotic normality of estimators
## ---
```

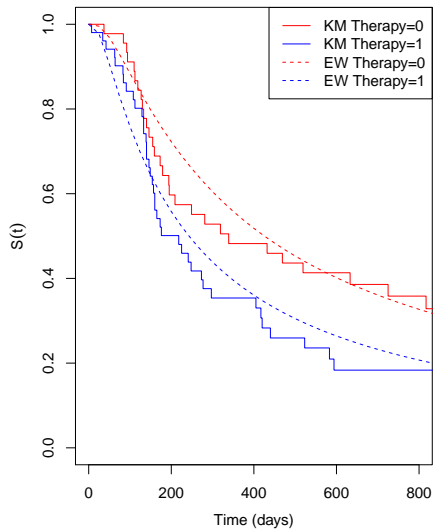
Estimates by Khan: $\hat{\beta}_0 = 10.183$, $\hat{\beta}_1 = 0.561$, $\hat{\alpha}_0 = 2.116$, $\hat{\lambda}_0 = 6.708$

Comparison

Weibull model



EW model



Comparison

```
AIC(modW, modEW)
```

```
##           df      AIC
## modW      3 1082.519
## modEW     4 1064.556
```

The AIC decreases, so EW model is better than Weibull model.

- Our routine performs a successfully estimation process.
- `maxlogLreg` facilitates the definition of initial values and bounds in estimation for the final user.
- This function is flexible: allows any distribution.
- Initial values are always a difficult issue.

References

- Efron, Bradley. 1988. "Logistic Regression, Survival Analysis, and the Kaplan-Meier Curve." *Journal of the American Statistical Association* 83 (402): 414. <https://doi.org/10.2307/2288857>.
- Khan, Shahedul A. 2018. "Exponentiated Weibull regression for time-to-event data." *Lifetime Data Analysis* 24 (2): 328–54. <https://doi.org/10.1007/s10985-017-9394-3>.
- Mullen, Katharine, David Ardia, David Gil, Donald Windover, and James Cline. 2011. "DEoptim : An R Package for Global Optimization by Differential Evolution." *Journal of Statistical Software* 40 (6). <https://doi.org/10.18637/jss.v040.i06>.
- Nash, John C. 1979. *Compact Numerical Methods for Computers. Linear Algebra and Function Minimisation*. 2nd Editio. Bristol: Adam Hilger.
- Pawitan, Yudi. 2013. *In all likelihood: statistical modelling and inference using likelihood*. Oxford University Press.