

# Gompertz

## Parametrisation

The Gompertz distribution has log survival function

$$\log S(y) = -\frac{\mu}{\alpha} (\exp(\alpha y) - 1)$$

for response  $y \geq 0$ ,  $\mu > 0$  and  $\alpha > 0$ . The cumulative distribution function and the density then follows as

$$F(y) = 1 - \exp \left[ -\frac{\mu}{\alpha} (\exp(\alpha y) - 1) \right]$$

and

$$f(y) = \mu \exp \left[ \alpha y - \frac{\mu}{\alpha} (\exp(\alpha y) - 1) \right].$$

## Link-function

The parameter  $\mu$  is linked to the linear predictor  $\eta$  as:

$$\mu = \exp(\eta)$$

## Hyperparameters

The shape parameter  $\alpha$  is represented as

$$\alpha = \exp(S\theta)$$

and the prior is defined on  $\theta$ . The constant  $S$  currently set to 0.1 to avoid numerical instabilities in the optimization, since small changes of  $\alpha$  can make a huge difference.

## Specification

- `family="gompertz"` for regression models and `family="gompertz.surv"` for survival models.
- Required arguments:  $y$  (to be given in a format by using `inla.surv()` for survival models )

## Hyperparameter spesification and default values

**doc** gompertz distribution

**hyper**

**theta**

**hyperid** 105101

**name** shape

**short.name** alpha

**initial** -1

**fixed** FALSE

**prior** normal

**param** 0 1

**to.theta** function(x, sc = 0.1) log(x) / sc

**from.theta** function(x, sc = 0.1) exp(sc \* x)

**status** experimental

**survival** FALSE

**discrete** FALSE

**link** default log neglog

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**hyper**

**theta1**

**hyperid** 106101

**name** shape

**short.name** alpha

**initial** -10

**fixed** FALSE

**prior** normal

**param** 0 1

**to.theta** function(x, sc = 0.1) log(x) / sc

**from.theta** function(x, sc = 0.1) exp(sc \* x)

**theta2**

**hyperid** 106102

**name** beta1

**short.name** beta1

**initial** -5

**fixed** FALSE

**prior** normal

**param** -4 100

**to.theta** function(x) x

**from.theta** function(x) x

**theta3**

**hyperid** 106103

**name** beta2

**short.name** beta2

**initial** 0

**fixed** FALSE

**prior** normal

**param** 0 100

**to.theta** function(x) x

**from.theta** function(x) x

**theta4**

**hyperid** 106104

**name** beta3

**short.name** beta3

**initial** 0

```

    fixed FALSE
    prior normal
    param 0 100
    to.theta function(x) x
    from.theta function(x) x
theta5
    hyperid 106105
    name beta4
    short.name beta4
    initial 0
    fixed FALSE
    prior normal
    param 0 100
    to.theta function(x) x
    from.theta function(x) x
theta6
    hyperid 106106
    name beta5
    short.name beta5
    initial 0
    fixed FALSE
    prior normal
    param 0 100
    to.theta function(x) x
    from.theta function(x) x
theta7
    hyperid 106107
    name beta6
    short.name beta6
    initial 0
    fixed FALSE
    prior normal
    param 0 100
    to.theta function(x) x
    from.theta function(x) x
theta8
    hyperid 106108
    name beta7
    short.name beta7
    initial 0
    fixed FALSE
    prior normal
    param 0 100
    to.theta function(x) x

```

```

    from.theta function(x) x
theta9
  hyperid 106109
  name beta8
  short.name beta8
  initial 0
  fixed FALSE
  prior normal
  param 0 100
  to.theta function(x) x
  from.theta function(x) x
theta10
  hyperid 106110
  name beta9
  short.name beta9
  initial 0
  fixed FALSE
  prior normal
  param 0 100
  to.theta function(x) x
  from.theta function(x) x
theta11
  hyperid 106111
  name beta10
  short.name beta10
  initial 0
  fixed FALSE
  prior normal
  param 0 100
  to.theta function(x) x
  from.theta function(x) x

```

**status** experimental

**survival** TRUE

**discrete** FALSE

**link** default log neglog

**pdf** gompertz

## Example

In the following example we estimate the parameters in a simulated case

```

library(flexsurv)
library(INLA)

n <- 1000
alpha <- 1.0
intercept <- 1.1
beta <- 1.2
x <- rnorm(n, sd = 0.2)
eta <- intercept + beta*x
mu <- exp(eta)
event <- rep(1,n)
y <- rgompertz(n, rate = mu, shape = alpha)

r <- inla(y ~ 1 + x,
          family = "gompertz", data=data.frame(y, x))
r.surv <- inla(inla.surv(y, event) ~ 1 + x,
               family = "gompertzsurv", data=data.frame(y, event, x))

## should be 'small'
print(r$mlik - r.surv$mlik)

```

Notes