

# Generalized Pareto distribution

## Parametrisation

The generalized Pareto (GP) distribution with positive shape parameter has cumulative distribution function

$$F(y; \sigma, \xi) = 1 - \left(1 + \xi \frac{y}{\sigma}\right)^{-1/\xi}, \quad y > 0,$$

for a continuous response  $y$  where

$\xi$ : is the shape parameter,  $\xi > 0$

$\sigma$ : is the scale parameter,  $\sigma > 0$

The limit for  $\xi \downarrow 0$  is  $F(y; \sigma, 0) = 1 - \exp(-x/\sigma)$ .

## Link function

The linear predictor  $\eta$  controls the  $\alpha$  quantile of the GP

$$P(y \leq q_\alpha) = \alpha$$

and  $q_\alpha = \exp(\eta)$ . The scaling  $\sigma$ , is then a function of  $(q_\alpha, \xi)$ , as

$$\sigma = \frac{\xi \exp(\eta)}{(1 - \alpha)^{-\xi} - 1}$$

## Hyperparameters

The GP model has one hyperparameter. The shape  $\xi > 0$  is represented as

$$\theta = \log \xi,$$

and the prior is defined on  $\theta$ .

## Specification

- family=gp
- Required arguments:  $y$  and the quantile  $\alpha$ .

The quantile is given as `control.family=list(quantile= $\alpha$ )`.

## Hyperparameter spesification and default values

**doc** Generalized Pareto likelihood

**hyper**

**theta**

**hyperid** 101201  
**name** shape  
**short.name** xi  
**initial** -2.30258509299405  
**fixed** FALSE  
**prior** loggamma  
**param** 1 15  
**to.theta** function(x) log(x)  
**from.theta** function(x) exp(x)

**status** experimental

**survival** FALSE

**discrete** TRUE

**link** default log

**pdf** genPareto

### Example

```
rgp = function(n, sigma, eta, alpha, xi = 0.001)
{
  if (missing(sigma)) {
    stopifnot(!missing(eta) && !missing(alpha))
    sigma = exp(eta) * xi / ((1.0 - alpha)^(-xi) - 1.0)
  }
  return (sigma / xi * (runif(n)^(-xi) - 1.0))
}
```

```
n = 300
x = runif(n)-0.5
eta = 1+x
alpha = 0.99
xi = 0.3
y = rgp(n, eta = eta, alpha = alpha, xi=xi)

r = inla(y ~ 1+x,
```

```
data = data.frame(y, x),  
family = "gp",  
control.family = list(quantile = alpha),  
control.predictor = list(compute=TRUE),  
verbose=TRUE)  
  
rx = range(c(r$summary.fitted.values$mean, exp(eta)))  
plot(r$summary.fitted.values$mean, exp(eta),  
      xlim = rx, ylim = rx)  
abline(a=0,b=1)
```

## Notes