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A Unified Approach for Probability Distribution FITTING WITH FITDISTRPLUS

M-L. Delignette-Muller¹, C. Dutang^{2,3}

¹ VetAgro Sud Campus Vétérinaire - Lyon ² ISFA - Lyon, ³ AXA GRM - Paris,



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OUTLINE

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- 4 MAXIMUM GOODNESS-OF-FIT ESTIMATION
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MAXIMUM LIKELIHOOD ESTIMATION - BRIEF REMINDER

Assuming a sample $(X_i)_{1 \leq i \leq n} \stackrel{i.i.d.}{\sim} X$, the likelihood

$$\mathcal{L}(\theta, x_1, \dots, x_n) = \prod_{i=1}^n f_X(x_i, \theta),$$

where f_X is the generic mass probability/density function. The MLE estimator $\theta_{\rm MLE}$ maximizes the likelihood

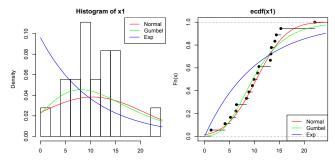
$$\theta_{\mathsf{MLE}} = \operatorname*{arg\,max}_{\theta \in \Theta} \mathcal{L}(\theta, x_1, \dots, x_n).$$

Example

```
> library(fitdistrplus)
                                                   $loglik
                                                   [11 -53.57625
> x1 <- c(6.4.13.3.4.1.1.3.14.1.
+ 10.6, 9.9, 9.6, 15.3, 22.1, 13.4,
                                                   Shessian
+ 13.2,8.4,6.3,8.9,5.2,10.9,14.4)
                                                              mean
                                                                          sd
                                                   mean 0.7987816 0.000000
> (f1 <- mledist(x1,"norm"))</pre>
                                                        0.0000000 1.597564
Sestimate
     mean
                                                   Soptim.function
10.411111 4.747033
                                                   [1] "optim"
$convergence
```

FUNCTION MLEDIST WITH NON R-BASE DISTRIBUTIONS

```
> dgumbel < -function(x,a,b) 1/b*exp((a-x)/b)*exp(-exp((a-x)/b))
> (f2 <- mledist(x1, "qumbel", start=list(a=10, b=5)))</pre>
$estimate
       а
8.094333 4.375401
$convergence
[1] 0
$loglik
[1] -54.09525
Shessian
a 0.9400408 -0.4418806
b -0.4418806 1.9124424
```



OPTIONAL ARGUMENTS OF MIEDIST

Fixed arguments

```
> (f4 <- mledist(x1, "gumbel",
start=list(b=5), fix.arg=list(a=7) ))
Sestimate
       h
4 248811
$convergence
[1] 0
$loglik
[11 -54.60187
Shessian
h 1 710569
Soptim.function
[1] "optim"
> f2
$estimate
```

Custom optimization

```
> fit1 <- mledist(x1, "gamma")
> fit1bis <- mledist(x1, "gamma", optim.method="BFGS")
> #wrap genoud function
> mygenoud <- function(fn, par, ...)
+ require (rgenoud)
+ res <- genoud(fn, starting.values=par, ...)
+ standardres <- c(res, convergence=0)
+ return(standardres)
+ }
> #custom optimization call
> fit2 <- mledist(x1, "gamma", custom.optim=mygenoud,
+ nvars=2, Domains=cbind(c(0,0), c(10, 10)),
+ boundary.enforcement=1, print.level=0, hessian=TRUE)
> cbind(NelderMead=fit1$estimate, BFGS=fit1bis$estimate,
+ Genoud=fit2Sestimate)
     NelderMead
                      BEGS
                              Genoud
shape 3.5747819 3.5768812 3.5742223
      0.3433516 0.3435683 0.3433094
rate
```

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8.094333 4.375401

MOMENT MATCHING ESTIMATION

It consists in equating the theoretical moments and the empirical moments

$$E[X^{k}; \theta] = \frac{1}{n} \sum_{i=1}^{n} X_{i}^{k}, \text{ for } k = 1, \dots, p.$$

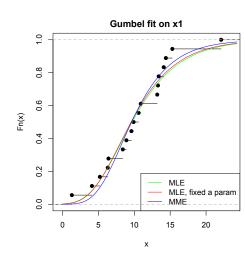
with $\theta \in \mathbb{R}^p$.

 $\theta_{\rm MME}$ can be computed in two ways, either by closed formulas (e.g. exponential distribution) or by square residual numeric minimization. Example with closed formulas

```
> (g1 <- mmedist(x1, "norm"))
Sestimate
                                                  $loglik
     mean
                 sd
                                                  [1] -53.57625
10.411111 4.747033
                                                  $method
$convergence
                                                  [1] "closed formula"
[1] 0
                                                  > cbind(MLE=f1$estimate, MME=g1$estimate)
$order
                                                              MLE
                                                                        MMF.
[1] 1 2
                                                  mean 10.411111 10.411111
                                                        4.747033 4.747033
                                                  sd
$memp
NUT.T.
```

MMEDIST - EXAMPLE WITH NUMERICAL OPTIMIZATION

```
> #empirical raw moment
> memp <- function(x, order)
+ ifelse(order == 1, mean(x),
+ sum(x^order)/length(x))
> #euler constant
> euler <- 0.5772156649
> #theoretical raw moment
> mgumbel <- function(order, a, b)
   mean <- a + b*euler
  if(order == 1)
  return (mean)
   else
  return(mean^2 + pi^2*b^2/6)
+ }
>
> g2 <- mmedist(x1, "gumbel", order=c(1, 2),</pre>
+ memp="memp", start=c(10, 5))
>
> cbind(MLE=f2$estimate, MLEfix=c(8,
+ f4$estimate[1]), MME=g2$estimate)
       MLE MLEfix
                          MME
a 8.094333 8.000000 8.260669
b 4.375401 4.248811 3.713298
```



QUANTILE MATCHING ESTIMATION

It consists in equating the theoretical quantiles and the empirical quantiles

$$q_{n,p_k} = F_X^{-1}(p_k), \text{ for } k = 1, \dots, p$$

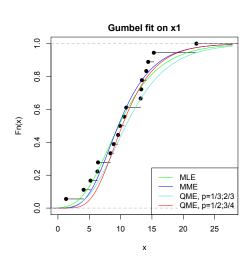
where q_{n,p_k} is the empirical quantile and $F_{\mathbf{x}}^{-1}(p_k)$ the theoretical one. p_k are given probabilities on which θ_{QME} is computed numerically. Example with normal distribution

```
> (h1 <- gmedist(x1, "norm", prob=c(1/2, 2/3))) $probs</pre>
                                                  [1] 0.5000000 0.6666667
$estimate
     mean
10 250030 6 926297
                                                 Soptim.function
                                                  [1] "optim"
$convergence
[1] 0
                                                 $loalik
                                                  [1] -55.60913
Švalue
[11 2.722893e-09
                                                 > h1bis <- qmedist(x1, "norm", prob=c(1/3, 2/3)
Shessian
                                                 > cbind(MLE=f1$estimate, MME=g1$estimate,
                                                  + OME1=h1$estimate, OME2=h1bis$estimate)
          mean
mean 4.0000000 0.8614546
                                                             MLE
                                                                       MME
                                                                                 OME 1
                                                                                           OME 2
sd 0.8614546 0.3710520
                                                 mean 10.411111 10.411111 10.250030 10.983327
                                                       4.747033 4.747033 6.926297 5.223794
```

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OMEDIST EXAMPLE

```
> #empirical quantiles computed with
> #the quantile() function
> #theoretical quantiles
> ggumbel <- function(p, a, b)
   a - b*log(-log(p))
>
> h2 <- gmedist(x1, "gumbel",
+ prob=c(1/3, 2/3), start=list(a=10, b=5))
> h2bis <- qmedist(x1, "gumbel",
+ prob=c(1/2, 3/4), start=list(a=10, b=5))
>
> cbind(MLE=f2$estimate, MME=g2$estimate,
+ QME1=h2$estimate, QME2=h2bis$estimate)
       MLE
                MME
                        OME 1
                                  QME2
a 8.094333 8.260669 9.157968 8.947923
b 4.375401 3.713298 4.514493 3.553285
```



MAXIMUM GOODNESS-OF-FIT ESTIMATION

It consists in maximizing a goodness of fit statistics, or equivalently minimizing a distance. Generally, we use the following statistics

Cramér-von Mises:

$$\Delta_{\mathsf{CvM}}^2 = \int_{\mathbb{R}} \left(F_n(x) - F_X(x) \right)^2 dx,$$

Kolmogorov Smirnov:

$$\Delta_{KS}^2 = \sup_{x} |F_n(x) - F_X(x)|,$$

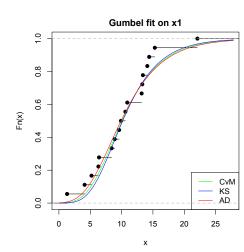
Anderson Darling:

$$\Delta_{AD}^{2} = n \int_{\mathbb{R}} \frac{(F_{n}(x) - F_{X}(x))^{2}}{F_{X}(x)(1 - F_{X}(x))} dx,$$

with F_n the empirical cdf and F_X the theoretical ones.

MGEDIST EXAMPLES

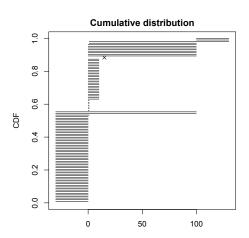
```
> i1 1 <- mgedist(x1, "norm", "CvM")
> i1_2 <- mgedist(x1, "norm", "KS")
> i1 3 <- mgedist(x1, "norm", "AD")
>
> cbind(MLE=f1$estimate, CvM= i1_1$estimate,
+ KS= i1 2$estimate, AD= i1 3$estimate)
           MLE
                    CvM
                               KS
                                          AΠ
mean 10.411111 10.34687 10.643541 10.336154
      4.747033 4.64827 4.595217 4.763116
sd
>
> i2 1 <- mgedist(x1, "gumbel", "CvM",
+ start=list(a=10, b=5))
> i2 2 <- mgedist(x1, "gumbel", "KS",</pre>
+ start=list(a=10, b=5))
> i2 3 <- mgedist(x1, "gumbel", "AD",
+ start=list(a=10, b=5))
> cbind(MLE=f2$estimate, CvM= i2 1$estimate,
+ KS= i2 2$estimate, AD= i2 3$estimate)
       MLE
                CvM
                          KS
                                    AΠ
a 8.094333 8.550061 8.736596 8.298506
b 4.375401 4.146123 3.916195 4.385616
```



CENSORED DATA

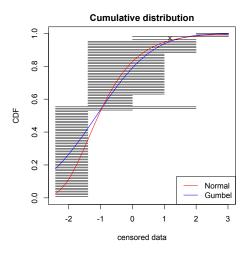
The *i*th obersvation x_i is not known exactly, but rather somewhere on an interval $x_i \in]l_i, u_i[$ with possible infinite bound. Non censored case is $l_i = u_i = x_i$.

```
> head(smokedfish, 10)
    left right
     NA 0.04
     NA 0.04
   0.04 10.00
   0.04 10.00
     NA 1.00
     NA 0.04
   0.04 10.00
    0.04 10.00
     NA 0.04
10 15.00 15.00
> plotdistcens(smokedfish)
```



ON THE USE OF MLEDISTCENS

Taking into account left and/or right censoring in the (log-)likelihood, maximum likelihood estimation can be carried out.



Conclusion (1/2)

Functionalities of the **fitdistrplus** package

- MLE: Extends the MASS fitdistr function with fixed arguments, custom optimization algorithms, possible censoring,
- MME: Provides a generic function to perform moment matching estimation with the raw or centered moments,
- QME: Based on the stats quantile function, provides the quantile matching estimation,
- MGE: Maximum goodness-of-fit is now available with the usual statistical distance and their variants.

So we can fit any probability distributions.

For specific probability distributions, please look at the task view http://cran.r-project.org/web/views/Distributions.html Introduction MLE MME QME MGE Censored Conclusion

CONCLUSION (2/2) - UNIFIED APPROACH WITH FITDIST

```
> f0 <- fitdist(x1, "gamma", method="mle")
> summary(f0)
Fitting of the distribution ' gamma '
by maximum likelihood
Parameters .
       estimate Std. Error
shape 3.5747819 1.1403248
rate 0.3433516 0.1175915
Loglikelihood: -54.44954
AIC: 112.8991
BTC: 114.6798
Correlation matrix:
          shape
                     rate
shape 1.0000000 0.9313999
rate 0.9313999 1.0000000
>
> plot(f0, col="turquoise")
> descdist(x1, boot=10, boot.col="turquoise")
summary statistics
min: 1.3 max: 22.1
median: 10.25
mean: 10.41111
estimated sd: 4.884657
estimated skewness: 0.3433588
estimated kurtosis: 3.755991
```

