

Continuous distributions

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Different types of distribution

- ① *continuous distributions*: $f_Y(y|\theta)$, are usually defined on $(-\infty, +\infty)$, $(0, +\infty)$ or $(0, 1)$.
- ② *discrete distributions*: $P(Y = y|\theta)$ are defined on $y = 0, 1, 2, \dots, n$, where n is a known finite value or n is infinite, i.e. usually discrete (count) values.
- ③ *mixed distributions*: (finite mixture distributions) are mixtures of continuous and discrete distributions, i.e. continuous distributions where the range of Y has been expanded to include some discrete values with non-zero probabilities.

Different types of distribution: demos

- 1 demo.GA()
- 2 demo.PO()
- 3 demo.BI()
- 4 demo.BE()
- 5 demo.BEINF

Types of continuous distributions

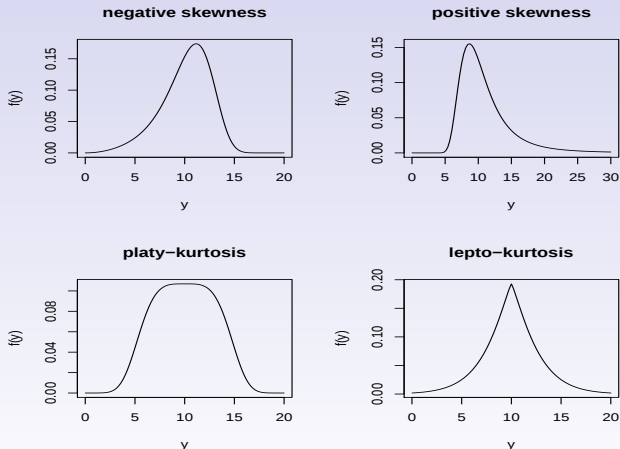


Figure: Showing different types of continuous distributions

Two parameter continuous distributions in GAMLSS

Two parameter distributions

BE Beta

GA Gamma

GU Gumbel

LO Logistic

LNO Log Normal

NO Normal

IG Inverse Gaussian

RG Reverse Gumbel

WEI Weibull (also WEI2, WEI3)

Three parameter continuous distributions in GAMLSS

Three parameter distributions

BCCG Box-Cox Normal

exGAUS Exponential-Gaussian

GG Generalized Gamma

GIG Generalized Inverse Gaussian

PE Power Exponential

TF t family

SN1 skew normal type 1

SN2 skew normal type 1

Two and three parameters comparison: demos

① `demo.PE.NO()`

② `demo.TF.NO()`

Four parameter continuous distributions in GAMLSS

Four parameter distributions

BCT Box-Cox t

BCPE Box-Cox Power Exponential

EGB1-EGB2 Exponential Generalized Beta

GT Generalized t

JSU Johnson Su

SHASH Sinh Arc-Sinh

SEP1-SEP4 Skew Exponential Power

ST1-ST5 Skew t

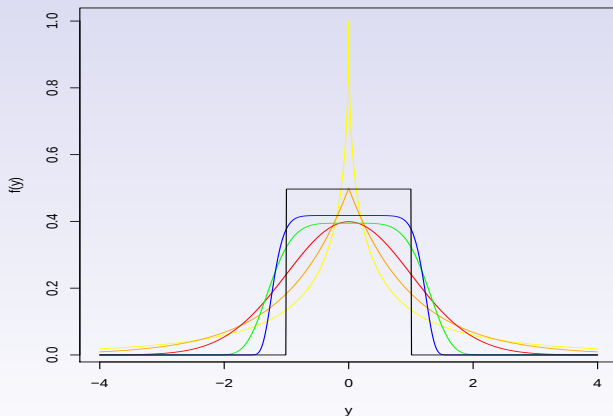
Continuous GAMLSS family distributions defined on $(-\infty, +\infty)$

Distributions	family	no par.	skewness	kurtosis
Exp. Gaussian	exGAUS	3	positive	-
Exp.G. beta 2	EGB2	4	both	lepto
Gen. t	GT	4	(symmetric)	lepto
Gumbel	GU	2	(negative)	-
Johnson's SU	JSU, JSUo	4	both	lepto
Logistic	LO	2	(symmetric)	(lepto)
Normal-Expon.- t	NET	2,(2)	(symmetric)	lepto
Normal	NO-NO2	2	(symmetric)	
Normal Family	NOF	3	(symmetric)	(meso)

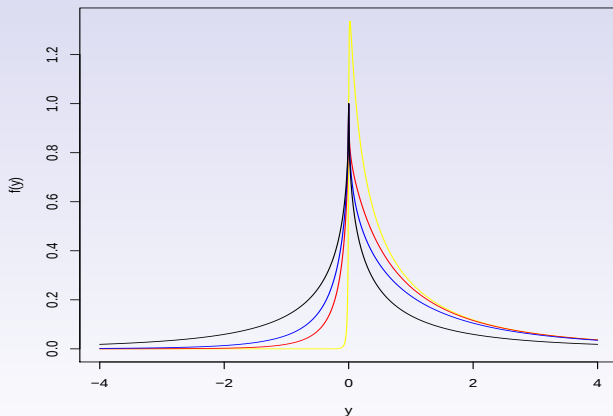
Continuous GAMLSS family distributions defined on $(-\infty, +\infty)$

Power Expon.	PE-PE2	3	(symmetric)	both
Reverse Gumbel	RG	2	positive	
Sinh Arcsinh	SHASH	4	both	both
Skew Exp. Power	SEP1-SEP4	4	both	both
Skew t	ST1-ST5	4	both	lepto
t Family	TF	3	(symmetric)	lepto

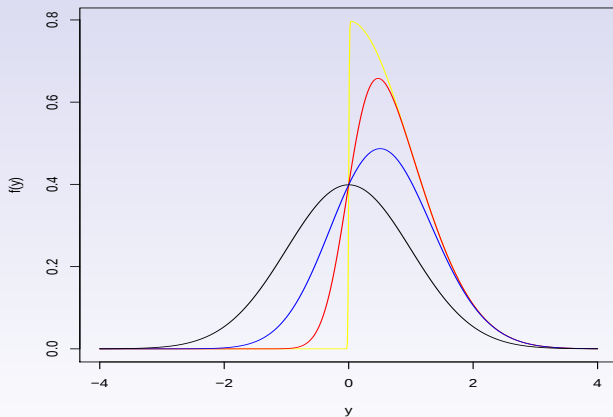
Symmetric SEP1 distribution ($\nu = 0$)



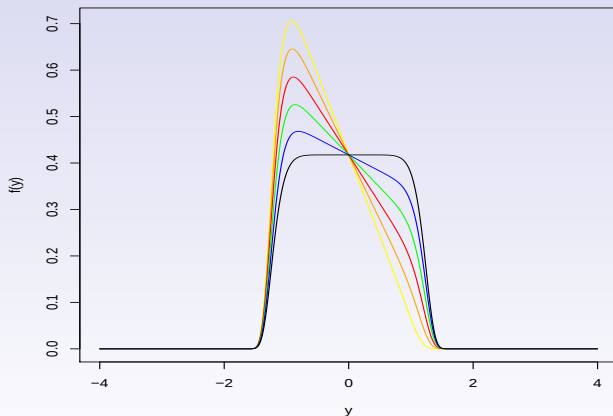
Skew SEP1 distributions ($\tau = 0.5$)



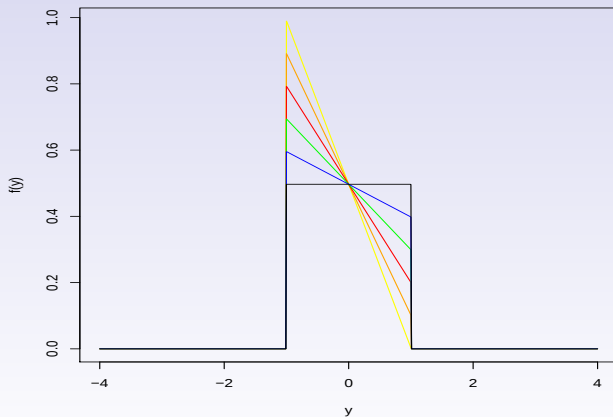
Skew SEP1 distributions ($\tau = 2$)



Skew SEP1 distributions ($\tau = 10$)



Skew SEP1 distributions ($\tau = 1000$)



Four parameters: demos

- 1 demo.SHASH()
- 2 demo.SEP1()

Continuous GAMLSS family distributions defined on $(0, +\infty)$

Distributions	family	no par.	skewness	kurtosis
BCCG	BCCG	3	both	
BCPE	BCPE	4	both	both
BCT	BCT	4	both	lepto
Exponential	EXP	1	(positive)	-
Gamma	GA	2	(positive)	-
Gen. Beta type 2	GB2	4	both	both
Gen. Gamma	GG-GG2	3	positive	-
Gen. Inv. Gaussian	GIG	3	positive	-
Inv. Gaussian	IG	2	(positive)	-
Log Normal	LOGNO	2	(positive)	-
Log Normal family	LNO	2,(1)	positive	
Reverse Gen. Extreme	RGE	3	positive	-
Weibull	WEI-WEI3	2	(positive)	

Positive response: demos

- 1 demo.NO.LOGNO()
- 2 demo.BCPE()

Continuous GAMLSS family distributions defined on $(0, 1)$

Distributions	family	no par.	skewness	kurtosis
Beta	BE	2	(both)	-
Beta original	BEo	2	(both)	-
Generalized beta type 1	GB1	4	(both)	(both)

Four parameters from 0 to 1: demos

1 demo.GB1()

Summary of methods of generating distributions

- ① univariate transformation from a single random variable
- ② transformation from two or more random variables
- ③ truncation distributions
- ④ a (continuous or finite) mixture of distributions
- ⑤ Azzalini type methods
- ⑥ splicing distributions
- ⑦ stopped sums
- ⑧ systems of distributions

Comparison of properties of distributions

- 1 Explicit pdf cdf and inverse cdf
- 2 Explicit centiles and centile based measures (e.g median)
- 3 Explicit moment based measures (e.g. mean)
- 4 Explicit mode(s)
- 5 Continuity of the pdf and its derivatives with respect to y
- 6 Continuity of the pdf with respect to μ , σ , ν and τ
- 7 Flexibility in modelling skewness and kurtosis
- 8 Range of Y

Comparison of properties

Univariate transformation: Explicit pdf, cdf, inverse cdf, centiles

Azzalini type methods: Non-explicit cdf, inverse cdf, centiles, but flexible

Splicing methods: Explicit pdf, cdf, inverse cdf, centiles Lack of continuity of second derivative of pdf.

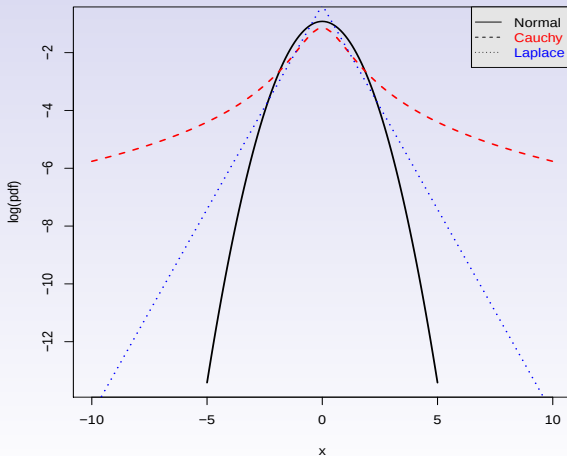
Theoretical comparison of distributions

- OK we have a lot of distributions in GAMLSS.
Do we need any more?
- Do we need all of them or some of them are redundant?

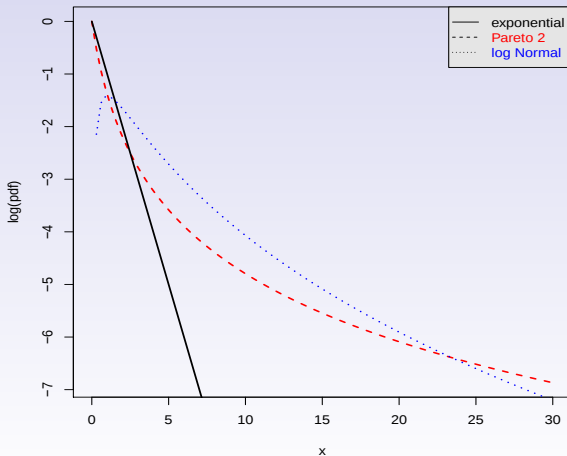
Is there any way to compare them theoretically?

- We can compare the **tail** behaviour in of the distributions
- Since most of them are location-scale we can compare their flexibility in terms of **skewness and kurtosis**

Comparing tails: real line



Comparing tails: positive real line



Types of tails

There are three main forms for $\log f_Y(y)$ for a tail of Y

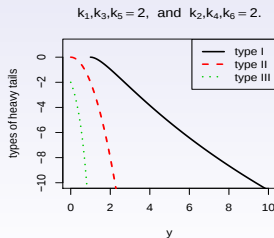
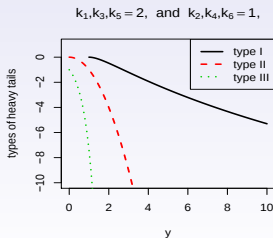
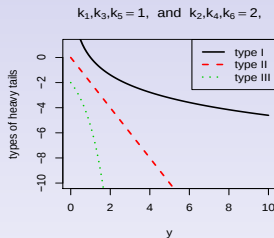
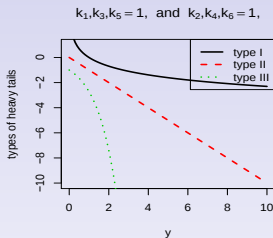
Type I: $-k_2 (\log |y|)^{k_1},$

Type II: $-k_4 |y|^{k_3},$

Type III: $-k_6 e^{k_5 |y|},$

decreasing k 's results in a heavier tail,

Types of tails



Types of tails: Bob's classification (real line)

Value of k_1-k_6	Distribution name	Distribution
$k_1 = 1$	Cauchy	CA(μ, σ)
	Generalized t	GT(μ, σ, ν, τ)
	Skew t type 3	ST3(μ, σ, ν, τ)
	Skew t type 4	ST4(μ, σ, ν, τ)
	Stable t	SB(μ, σ, ν, τ) TF(μ, σ, ν)
$k_1 = 2$	Johnson's SU	JSU(μ, σ, ν, τ)
	Johnson's SU original	JSUo(μ, σ, ν, τ)
$0 < k_3 < \infty$	Power exponential	PE(μ, σ, ν)
	Power exponential type 2	PE2(μ, σ, ν)
	Sinh-arcsinh original	SHASHo(μ, σ, ν, τ)
	Sinh-arcsinh	SHASH(μ, σ, ν, τ)
	Skew exponential power type 3	SEP3(μ, σ, ν, τ)
	Skew exponential power type 4	SEP4(μ, σ, ν, τ)
	Exponential generalized beta type 2	EGB2(μ, σ, ν, τ)
	Gumbel	GU(μ, σ)
	Laplace	LA(μ, σ)
	Logistic	LG(μ, σ)
	Reverse Gumbel	RG(μ, σ)
$k_3 = 2$	Normal	NO(μ, σ)
$0 < k_5 < \infty$	Gumbel	GU(μ, σ)
	Reverse Gumbel	RG(μ, σ)

Types of tails: Bob's classification (positive real line)

Value of k_1-k_6	Distribution name	Distribution
$k_1 = 1$	Box-Cox Cole-Green	BCCG(μ, σ, ν)
	Box-Cox power exponential	BCPE(μ, σ, ν, τ)
	Box-Cox t	BCT(μ, σ, ν, τ)
	Generalized beta type 2	GB2(μ, σ, ν, τ)
	Generalized gamma	GG(μ, σ, ν)
	Inverse gamma	IGA(μ, σ)
	log t	LOGT(μ, σ, ν)
$k_1 = 2$	Pareto Type 2	PA2o(μ, σ)
	Box-Cox Cole-Green	BCCG(μ, σ, ν)
	Lognormal	LOGNO(μ, σ)
	Log Weibull	LOGWEI(μ, σ)
$1 \leq k_1 < \infty$	Box-Cox power exponential	BCPE(μ, σ, ν, τ)
$0 < k_3 < \infty$	Box-Cox Cole-Green	BCCG(μ, σ, ν)
	Box-Cox power exponential	BCPE(μ, σ, ν, τ)
	Generalized gamma	GG(μ, σ, ν)
	Weibull	WEI(μ, σ)
$k_3 = 1$	Exponential	EX(μ)
	Gamma	GA(μ, σ)
	Generalized inverse Gaussian	GIG(μ, σ, ν)
	Inverse Gaussian	IG(μ, σ)

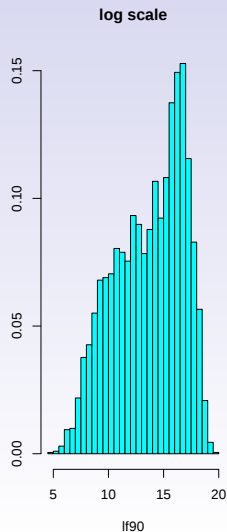
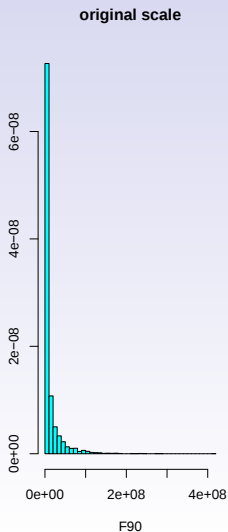
Types of tails

- Given we are dealing with a long tail situation can we use the tail classification in order to chose an appropriate distribution for data?
- For regression problem: ?? (research under progress)
- For single response variable (no explanatory variables): YES

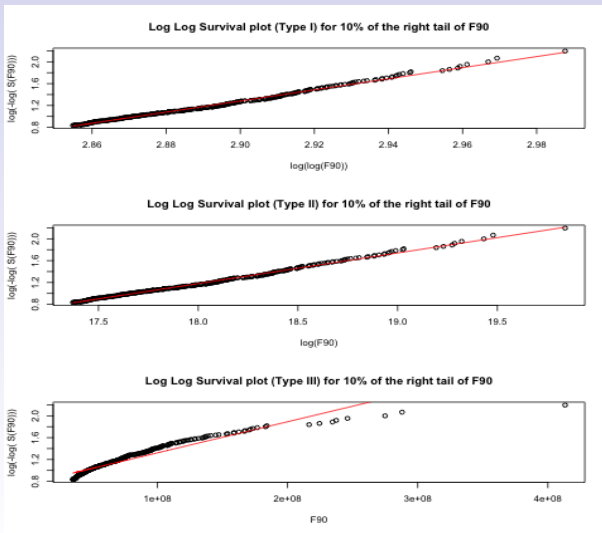
Types of tails: single variable

- complementary cumulative distribution function (CCDF) ($\log [S_Y(y)]$ against $\log y$).
- complementary log log plot of the survival function.
- Hill 'type' plot

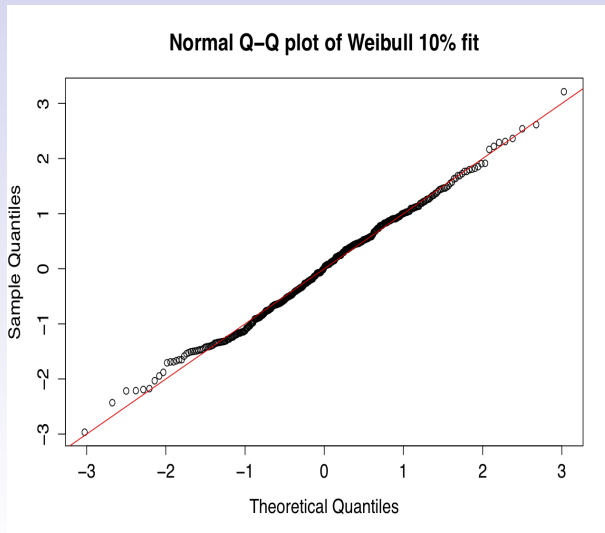
The 90's Film data



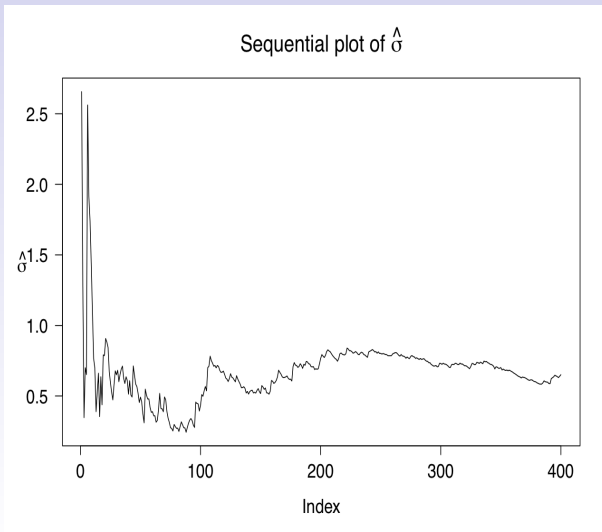
Comparing tails: complementary log log survival.



Comparing tails: QQ plot from a Weibull



Comparing tails: Hill 'type' plot for Weibull



Comparing skewness and kurtosis

Since moment based measures may not exist or may be unreliable, **centile based** measures of skewness and kurtosis are used for comparison

A general **measure of skewness**;

$$s_p = \frac{(y_p + y_{1-p})/2 - y_{0.5}}{(y_{1-p} - y_p)/2}$$

where $y_p = F_Y^{-1}(p)$. For example $p = 0.25$, giving **Galton's measure of skewness**

$$s_{0.25} = \frac{(Q_1 + Q_3)/2 - m}{(Q_3 - Q_1)/2}$$

a measure of central skewness

Comparing skewness and kurtosis

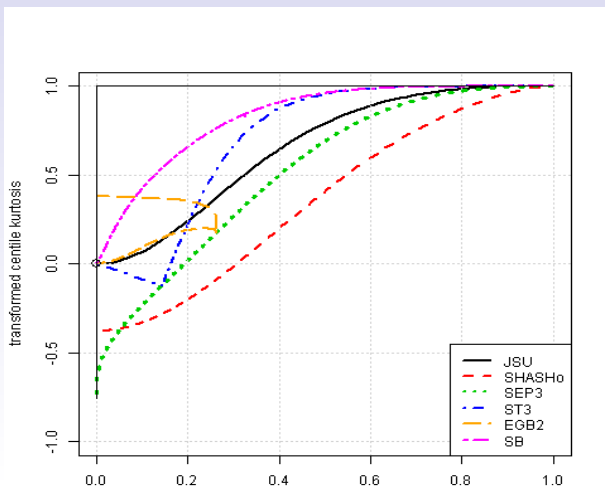
A general measure of kurtosis

$$k_p = \frac{(y_{1-p} - y_p)}{(Q_3 - Q_1)}$$

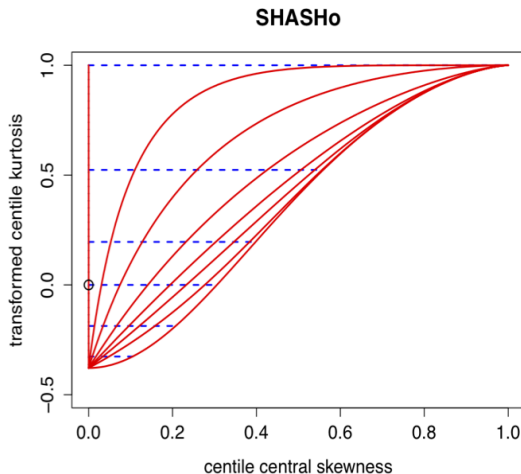
standardised

$$s k_{0.01} = \frac{(y_{0.99} - y_{0.01})}{3.49 (Q_3 - Q_1)}, \quad (1)$$

Comparing skewness and kurtosis: a plot of the boundary of central skewness against transformed kurtosis



Comparing skewness and kurtosis: The SHASHo distribution



Comparing skewness and kurtosis

- There is some progress in classifying the continuous `gamlss.family` distributions
- the results apply to all continuous distributions (new or old)

Theoretical considerations for fitting distributions

$$L(\boldsymbol{\theta}) = \prod_{i=1}^n f_Y(y_i|\boldsymbol{\theta})$$

$$L(\boldsymbol{\theta}) = \prod_{i=1}^n [F_Y(y_i + \Delta_i|\boldsymbol{\theta}) - F_Y(y_i - \Delta_i|\boldsymbol{\theta})]$$

$$L(\boldsymbol{\theta}) \approx \prod_{i=1}^n f_Y(y_i|\boldsymbol{\theta})\Delta_i = \left[\prod_{i=1}^n \Delta_i \right] \left[\prod_{i=1}^n f_Y(y_i|\boldsymbol{\theta}) \right]$$

$$\ell(\boldsymbol{\theta}) = \sum_{i=1}^n \log f_Y(y_i|\boldsymbol{\theta}) + \sum_{i=1}^n \log \Delta_i$$

How to fit distributions in R

`optim()` or `mle()` requires initial parameter values

`gamlssML()` mle a using variation of the `mle()` function

`gamlss()`: mle using RS or CG or mixed algorithms,

`histdist()` mle using RS or CG or mixed algorithms, for a univariate sample only, and plots a histogram of the data with the fitted distribution

`fitDist()` fits a set of distributions and chooses the one with the smaller GAIC

Strength of glass fibres data

Data summary:

R data file: glass in package `gamlss.data` of dimensions 63×1

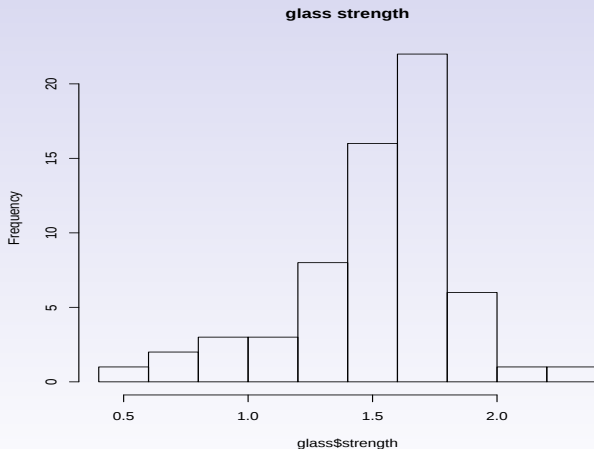
source: Smith and Naylor (1987)

strength: the strength of glass fibres (the unit of measurement are not given).

purpose: to demonstrate the fitting of a parametric distribution to the data.

conclusion: a SEP4 distribution fits adequately

Strength of glass fibres data: SBC



Strength of glass fibres data: creating truncated distribution

```
data(glass)
library(gamlss.tr)
gen.trun(par = 0, family = TF)
```

A truncated family of distributions from TF
has been generated
and saved under the names:
dTFtr pTFtr qTFtr rTFtr TFtr

The type of truncation is left and the
truncation parameter is 0

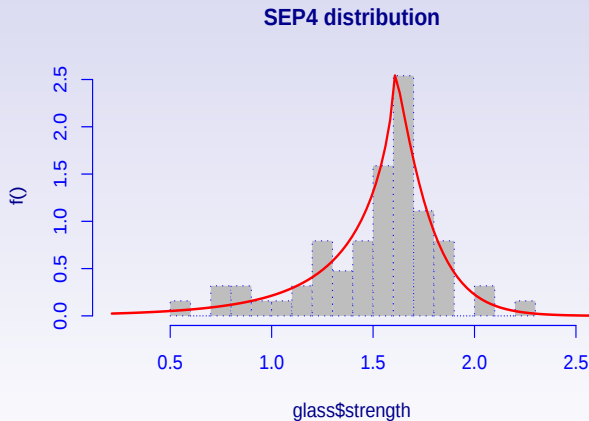
Strength of glass fibres data: fitting the models

```
> m1<-fitDist(strength, data=glass, k=2, extra="TFtr")
      # AIC
> m2<-fitDist(strength, data=glass, k=log(length(strength))
      extra="TFtr") # SBC
> m1$fit[1:8]
      SEP4      SEP3    SHASHo      EGB2      JSU      BCPEo
27.68790 27.98191 28.01800 28.38691 30.41380 31.17693 31.40
> m2$fit[1:8]
      SEP4      SEP3    SHASHo      EGB2      GU      WEI3
36.26044 36.55444 36.59054 36.95945 38.19839 38.69995 38.98
```

Strength of glass fibres data: Results

```
histDist(glass$strength, SEP4, nbins = 13,  
         main = "SEP4 distribution",  
         method = mixed(20, 50))
```

Strength of glass fibres data: SBC



Strength of glass fibres data: summary

```
Call:  gamlss(formula = y~1, family = FA)
```

```
Mu Coefficients:
```

```
(Intercept)
```

```
1.581
```

```
Sigma Coefficients:
```

```
(Intercept)
```

```
-1.437
```

```
Nu Coefficients:
```

```
(Intercept)
```

```
-0.1280
```

```
Tau Coefficients:
```

```
(Intercept)
```

```
0.3183
```

```
Degrees of Freedom for the fit: 4 Residual Deg. of Freedom
```

```
Global Deviance:      19.8111
```

```
AIC:      27.8111
```

Example of fitting regression models: CD4 count data

Data summary:

R data file: CD4 in package **gamlss.data** of dimensions 609×2

source: Wade and Ades (1994)

variables:

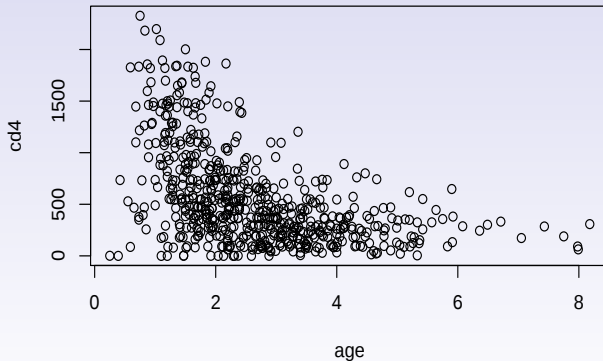
cd4 : CD4 counts from uninfected children born to HIV-1 mothers

age : age in years of the child.

purpose: to demonstrate regression fitting of different functional forms for the explanatory variable age.

conclusion: models for both μ and σ are needed with a 4-parameter distribution for the response variable

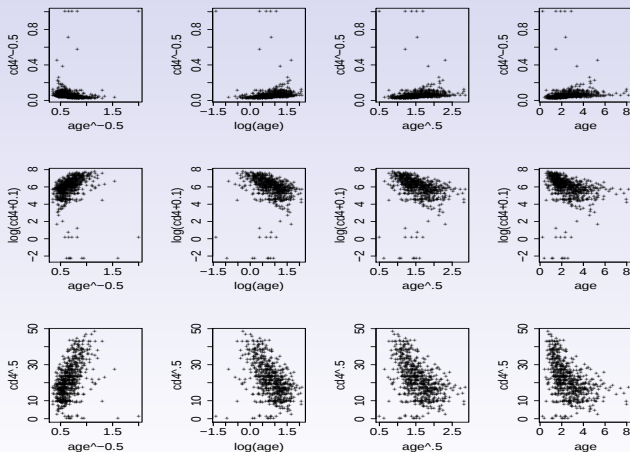
CD4 count data: plotting the data



Examples

Example of fitting regression models: CD4 count data

CD4 count data: transforming the data



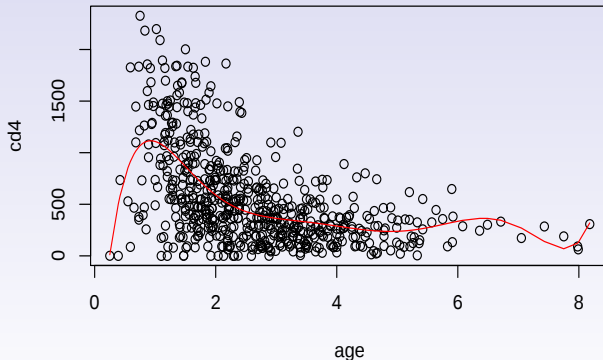
CD4 count data: fitting polynomials

```
m1 <- gamlss(cd4 ~ age, sigma.fo = ~1, data = CD4, control
m2 <- gamlss(cd4 ~ poly(age, 2), sigma.fo = ~1, data = CD4)
m3 <- gamlss(cd4 ~ poly(age, 3), sigma.fo = ~1, data = CD4)
m4 <- gamlss(cd4 ~ poly(age, 4), sigma.fo = ~1, data = CD4)
m5 <- gamlss(cd4 ~ poly(age, 5), sigma.fo = ~1, data = CD4)
m6 <- gamlss(cd4 ~ poly(age, 6), sigma.fo = ~1, data = CD4)
m7 <- gamlss(cd4 ~ poly(age, 7), sigma.fo = ~1, data = CD4)
m8 <- gamlss(cd4 ~ poly(age, 8), sigma.fo = ~1, data = CD4)
```

CD4 count data: fitting polynomials

	df	AIC
m7	9	8963.263
m8	10	8963.874
m5	7	8977.383
m4	6	8988.105
m3	5	8993.351
m2	4	8995.636
m1	3	9044.145

CD4 count data: fitting polynomials of degree 7



CD4 count data: fractional polynomials

```
m1f <- gamlss(cd4 ~ fp(age, 1), sigma.fo = ~1, data = CD4)
m2f <- gamlss(cd4 ~ fp(age, 2), sigma.fo = ~1, data = CD4)
m3f <- gamlss(cd4 ~ fp(age, 3), sigma.fo = ~1, data = CD4)
```

CD4 count data: fractional polynomials

	df	AIC
m3f	8	8966.375
m2f	6	8978.469
m1f	4	9015.321

CD4 count data: fractional polynomials

```
m3f$mu.coefSmo
```

```
[[1]]
```

```
[[1]]$coef
```

```
one
```

```
-599.2970 1116.7924 1776.1937 698.6097
```

```
[[1]]$power
```

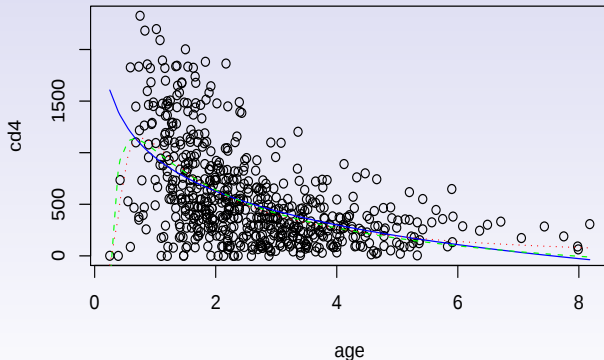
```
[1] -2 -2 -2
```

```
[[1]]$varcoeff
```

```
[1] 2016.238 4316.743 22835.146 7521.417
```

terms : age^{-2} , $age^{-2} \log(age)$ and $age^{-2} [\log(age)]^2$.

CD4 count data: fractional polynomials fits



CD4 count data: piecewise polynomials

```
m2b <- gamlss(cd4 ~ bs(age), data = CD4)
m3b <- gamlss(cd4 ~ bs(age, df = 3), data = CD4)
m4b <- gamlss(cd4 ~ bs(age, df = 4), data = CD4)
m5b <- gamlss(cd4 ~ bs(age, df = 5), data = CD4)
m6b <- gamlss(cd4 ~ bs(age, df = 6), data = CD4)
m7b <- gamlss(cd4 ~ bs(age, df = 7), data = CD4)
m8b <- gamlss(cd4 ~ bs(age, df = 8), data = CD4)
```

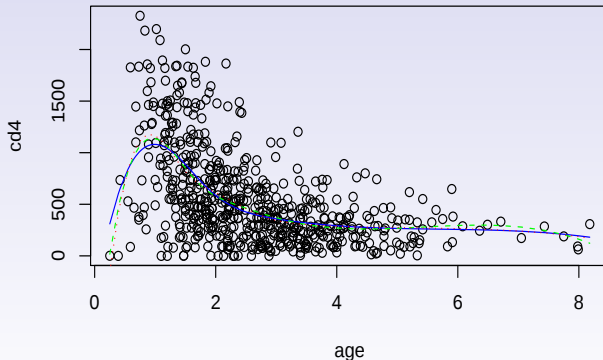
CD4 count data: piecewise polynomials GAIC(2)

	df	AIC
m7b	9	8959.519
m6b	8	8960.353
m8b	10	8961.073
m5b	7	8964.022
m4b	6	8977.475
m2b	5	8993.351
m3b	5	8993.351

CD4 count data: piecewise polynomials $\text{GAIC}(\log(n))$

	df	AIC
m5b	7	8994.904
m6b	8	8995.648
m7b	9	8999.225
m4b	6	9003.946
m8b	10	9005.191
m2b	5	9015.410
m3b	5	9015.410

CD4 count data: piecewise polynomials fits



CD4 count data: cubic smoothing splines

```
fn <- function(p) AIC(gamlss(cd4 ~ cs(age, df = p[1]),  
  data = CD4, trace = FALSE), k = 2)  
opAIC <- optim(par = c(3), fn, method = "L-BFGS-B",  
  lower = c(1), upper = c(15))  
fn <- function(p) AIC(gamlss(cd4 ~ cs(age, df = p[1]),  
  data = CD4, trace = FALSE), k = log(length(CD4$age)))  
opSBC <- optim(par = c(3), fn, method = "L-BFGS-B",  
  lower = c(1), upper = c(15))  
opAIC$par  
[1] 10.85157  
opSBC$par  
[1] 1.854689
```

effective df should be between 1.8 and 10.8

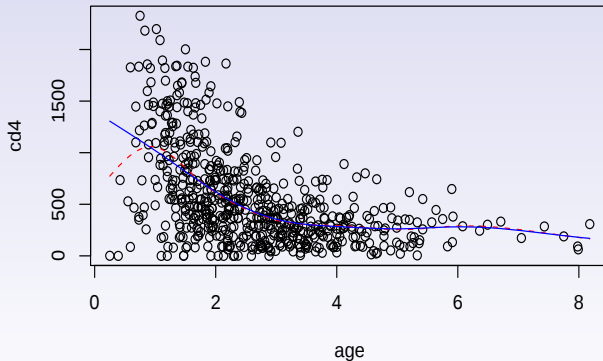
CD4 count data: cubic P-splines

```
mpb1 <- gamlss(cd4 ~ pb(age), data = CD4)
mpb1$mu.df}
[1] 8.606468
```

CD4 count data: Modelling sigma using cubic P-splines

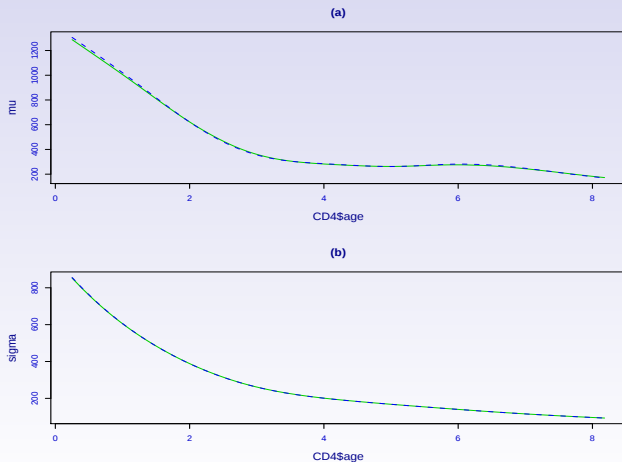
```
mpb2 <- gamlss(cd4 ~ pb(age), sigma.fo = ~pb(age),  
  data = CD4, gd.tol = 10)  
mpb2$mu.df  
[1] 6.200118      compare to 5.72 for cs()  
mpb2$sigma.df    compare to 3.81 for cs()  
[1] 3.803651
```

CD4 count data: P-spline fits



Example of fitting regression models: CD4 count data

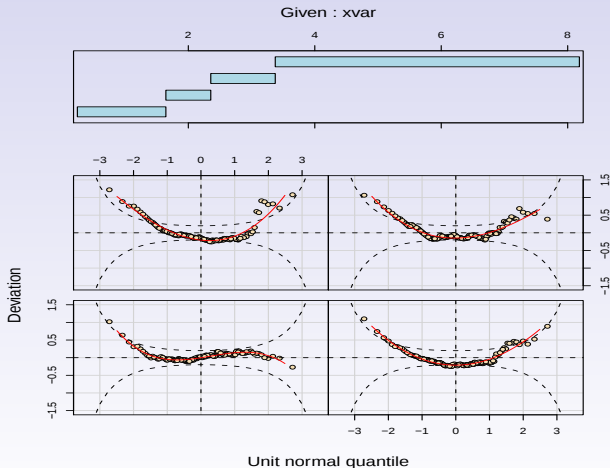
CD4 count data: Comparing P-splines with Cubic splines



CD4 count data: Validation Global Deviance

```
set.seed(1234)
rSample6040 <- sample(2, length(CD4$cd4), replace=T,
                      prob=c(0.6, 0.4))
fn <- function(p) VGD(cd4~cs(age, df=p[1]),
                      sigma.fo=~cs(age, df=p[2]),
                      data=CD4, rand=rSample6040)
op<-optim(par=c(3,1), fn, method="L-BFGS-B", lower=c(1,1),
          upper=c(10,10))
op$par
[1] 4.779947 1.376534
```

CD4 count data: Worm plot



CD4 count data: Fitting different distributions

distributions	df	AIC
SEP2	8.86	8685.69
SEP1	8.77	8688.08
SEP3	11.24	8693.68
ST2	9.31	8693.91
ST1	9.31	8693.91
ST3	11.33	8694.21
JSU	13.28	8701.05
SHASH	12.75	8705.26
ST5	8.99	8724.25
ST4	12.54	8749.01
TF	11.48	8785.20
PE	11.17	8790.16

END

for more information see

www.gamlss.org