Introdução ao GAMLSS

Gustavo Thomas e Cristian Villegas

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Sumário

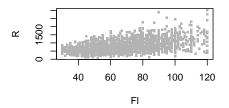
1	Introdução			
	1.1	exercises Chapter 1	17	
	1.2	$\operatorname{gamlss.demo}()$	17	

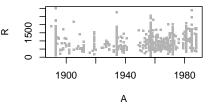
4 SUMÁRIO

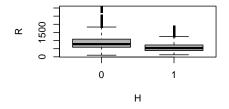
Capítulo 1

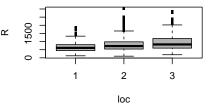
Introdução

```
library(knitr)
library(gamlss)
library(xtable)
PPP <- par(mfrow=c(2,2))
plot(R~F1, data=rent, col=gray(0.7), pch=15, cex=0.5)
plot(R~A, data=rent, col=gray(0.7), pch=15, cex=0.5)
plot(R~H, data=rent, col=gray(0.7), pch=15, cex=0.5)
plot(R~loc, data=rent, col=gray(0.7), pch=15, cex=0.5)</pre>
```



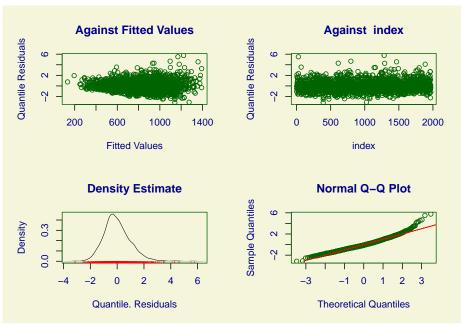






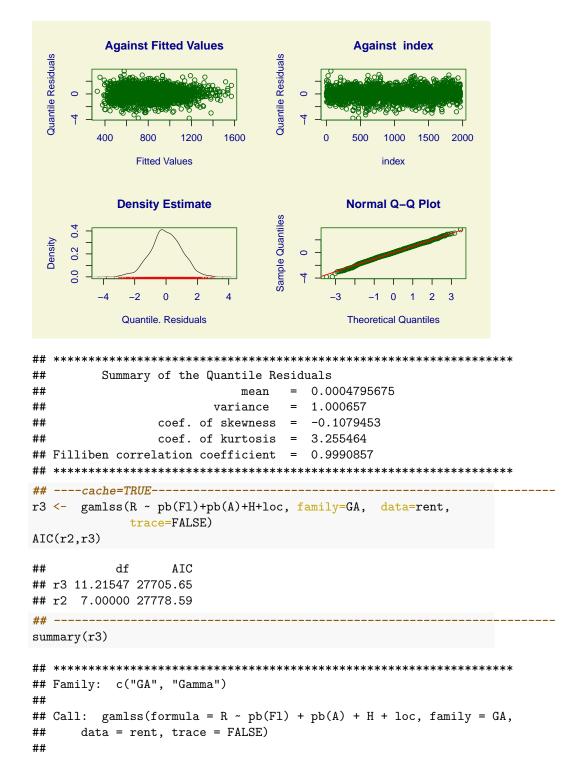
```
par(PPP)
# ----eval=FALSE----
## R \sim F1+A+H+loc
## ----cache=TRUE-----
r1 <- gamlss(R ~ Fl+A+H+loc, family=NO, data=rent, trace=FALSE)
11 <- lm(R ~ Fl+A+H+loc, data=rent)</pre>
coef(r1)
## (Intercept)
                     Fl
                                          H1
                                                   loc2
                                                              1oc3
## -2775.038803 8.839445
                         1.480755 -204.759562 134.052349
                                                         209.581472
coef(11)
## (Intercept)
                     Fl
                                Α
                                          H1
                                                   loc2
                                                              loc3
## -2775.038803 8.839445 1.480755 -204.759562
                                              134.052349
                                                         209.581472
fitted(r1, "sigma")[1]
##
## 308.4768
summary(r1)
## **********************
## Family: c("NO", "Normal")
##
## Call: gamlss(formula = R ~ Fl + A + H + loc, family = NO,
   data = rent, trace = FALSE)
##
## Fitting method: RS()
##
## -----
## Mu link function: identity
## Mu Coefficients:
##
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) -2775.0388 470.1352 -5.903 4.20e-09 ***
                       0.3370 26.228 < 2e-16 ***
## Fl
               8.8394
## A
               1.4808
                       0.2385
                               6.208 6.55e-10 ***
## H1
            -204.7596 18.9858 -10.785 < 2e-16 ***
## loc2
            134.0523
                       25.1430 5.332 1.09e-07 ***
## loc3
             209.5815
                       27.1286 7.725 1.76e-14 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## -----
## Sigma link function: log
## Sigma Coefficients:
            Estimate Std. Error t value Pr(>|t|)
##
```

```
## (Intercept) 5.73165
                      0.01594
                               359.7
                                     <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## No. of observations in the fit: 1969
## Degrees of Freedom for the fit:
##
       Residual Deg. of Freedom:
                             1962
##
                     at cycle: 2
##
## Global Deviance:
                    28159
##
            AIC:
                    28173
            SBC:
                    28212.1
Rsq(r1)
## [1] 0.3372028
## ----LM_residual_plot,echo=TRUE,fig.show='hide', fig.asp=1-----
plot(r1)
```



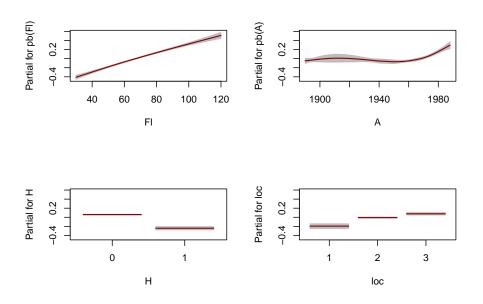
```
##
               coef. of kurtosis = 4.844416
## Filliben correlation coefficient = 0.9859819
## ----cache=TRUE-----
### using gamlss
r2 <- gamlss(R ~ Fl+A+H+loc, family=GA, data=rent)
## GAMLSS-RS iteration 1: Global Deviance = 27764.59
## GAMLSS-RS iteration 2: Global Deviance = 27764.59
coef(r2)
## (Intercept)
                    Fl
                                         H1
                                                 loc2
                                                            loc3
## 2.86497701 0.01062319 0.00151005 -0.30007446 0.19076406 0.26408285
coef(r2, "sigma") ### extract log(sigma)
## (Intercept)
## -0.9821991
deviance(r2)
## [1] 27764.59
### using glm
12 <- glm(R ~ Fl+A+H+loc, family=Gamma(link="log"), data=rent)
coef(12)
## (Intercept)
                      Fl
                                            Н1
                                                      loc2
                                                                 loc3
## 2.864943806 0.010623194 0.001510066 -0.300074001 0.190764594 0.264083376
summary(12)$dispersion ### extract phi
## [1] 0.1377881
deviance(12)
## [1] 282.5747
## -----
summary(r2)
## Family: c("GA", "Gamma")
## Call: gamlss(formula = R ~ Fl + A + H + loc, family = GA, data = rent)
## Fitting method: RS()
## -----
## Mu link function: log
```

```
## Mu Coefficients:
           Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.8649770 0.5681289 5.043 5.01e-07 ***
         0.0106232 0.0004128 25.735 < 2e-16 ***
## Fl
          0.0015100 0.0002886
## A
                         5.232 1.85e-07 ***
## H1
         -0.3000745 0.0231153 -12.982 < 2e-16 ***
## loc2
         0.1907641 0.0305203 6.250 5.01e-10 ***
## loc3
         ## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## -----
## Sigma link function: log
## Sigma Coefficients:
         Estimate Std. Error t value Pr(>|t|)
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## -----
## No. of observations in the fit: 1969
## Degrees of Freedom for the fit: 7
     Residual Deg. of Freedom: 1962
##
                at cycle: 2
##
## Global Deviance:
                27764.59
##
         AIC:
                27778.59
          SBC:
##
                27817.69
## -----
r22 <- gamlss(R ~ Fl+A+H+loc, family=IG, data=rent, trace=FALSE)
GAIC(r1, r2, r22, k=0) # GD
```



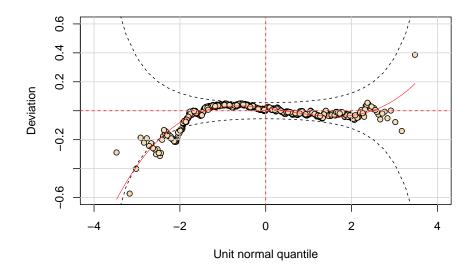
```
## Fitting method: RS()
## -----
## Mu link function: log
## Mu Coefficients:
            Estimate Std. Error t value Pr(>|t|)
## (Intercept) 3.0851197 0.5692315 5.420 6.70e-08 ***
## pb(F1) 0.0103084 0.0004031 25.573 < 2e-16 ***
          0.0014062 0.0002893 4.861 1.26e-06 ***
## pb(A)
## H1
          -0.3008111 0.0225869 -13.318 < 2e-16 ***
## loc2
          0.1886692  0.0299295  6.304  3.58e-10 ***
## loc3
          0.2719856 0.0322862 8.424 < 2e-16 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## -----
## Sigma link function: log
## Sigma Coefficients:
          Estimate Std. Error t value Pr(>|t|)
## (Intercept) -1.00196 0.01559 -64.27 <2e-16 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## -----
## NOTE: Additive smoothing terms exist in the formulas:
## i) Std. Error for smoothers are for the linear effect only.
## ii) Std. Error for the linear terms maybe are not accurate.
## -----
## No. of observations in the fit: 1969
## Degrees of Freedom for the fit: 11.21547
##
      Residual Deg. of Freedom: 1957.785
##
                  at cycle: 3
##
## Global Deviance:
                 27683.22
                 27705.65
           AIC:
           SBC:
                  27768.29
## *********************
## ----cache=TRUE------
drop1(r3)
## Single term deletions for
## mu
##
## Model:
## R ~ pb(F1) + pb(A) + H + loc
```

```
Pr(Chi)
##
             Df
                  AIC
                         LRT
## <none>
                 27706
## pb(Fl) 1.4680 28261 558.59 < 2.2e-16 ***
         4.3149 27798 101.14 < 2.2e-16 ***
          1.8445 27862 160.39 < 2.2e-16 ***
## H
          2.0346 27770 68.02 1.825e-15 ***
## loc
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## ----GAM_termplot, echo=TRUE,fig.show='hide', fig.asp=1,cache=TRUE------
term.plot(r3, pages=1, ask=FALSE)
```



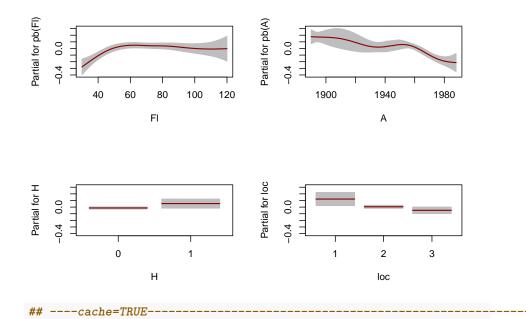
wp(r3, ylim.all=.6)

----GAM_wormplot, echo=TRUE,fig.show='hide', fig.asp=1,cache=TRUE------

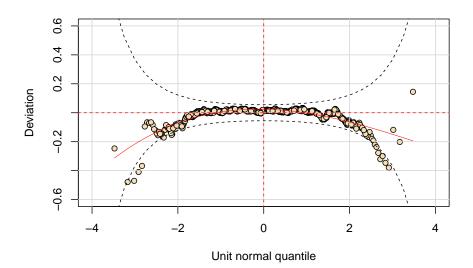


```
## df AIC
## r4 22.25035 27614.78
## r3 11.21547 27705.65
## r5 21.82318 27716.66
## ----MADAM_termplot, echo=TRUE,fig.show='hide', fig.asp=1,cache=TRUE-----
term.plot(r4, pages=1, what="sigma", ask=FALSE)
```

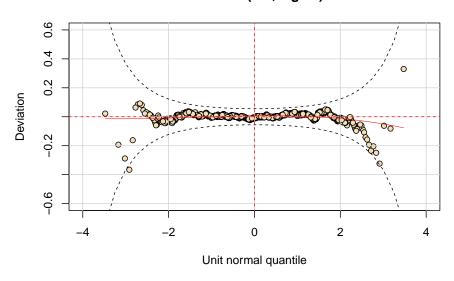
drop1(r4, what="sigma")



```
## Single term deletions for
## sigma
##
## Model:
## \simpb(F1) + pb(A) + H + loc
##
                   AIC
                                Pr(Chi)
              Df
                          LRT
## <none>
                  27615
## pb(F1) 4.02694 27631 24.683 5.997e-05 ***
## pb(A) 3.87807 27659 52.167 1.067e-10 ***
## H
          0.88335 27615
                        1.866
                                 0.14788
          2.03694 27619
## loc
                        8.036
                                 0.01872 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## ----MADAM_wormplot, echo=TRUE,fig.show='hide', fig.asp=1,cache=TRUE-----
wp(r4, ylim.all=.6)
```

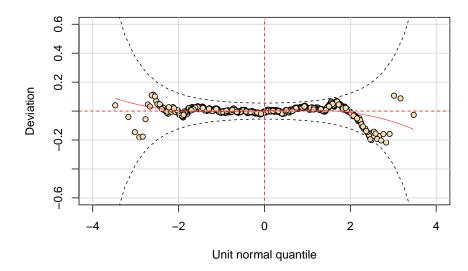


r6: BCCG(mu, sigma)



wp(r7, ylim.all=.6); title("r7: BCCG(mu, sigma, nu)")

r7: BCCG(mu, sigma, nu)



1.1 exercises Chapter 1

```
## data(airquality)
## plot(airquality[,-c(5,6)])
## # Fit the standard linear model
## air.lm <- lm(Ozone~Temp+Wind+Solar.R,data=airquality)</pre>
## summary(air.lm)
## op<-par(mfrow=c(1,3))
## termplot(air.lm,partial.resid=TRUE,se=T)
## par(op)
## op<-par(mfrow=c(1,2))
## plot(air.lm, which=1:2)
## par(op)
## library(gamlss)
## da <- na.omit(airquality) # clear the data of NA's
## mno<-gamlss(Ozone~Temp+Wind+Solar.R, data=da) # fit the model
## summary(mno)
## term.plot(mno, pages=1, partial=T) # plot the fitted terms
## plot(mno)
## wp(mno)
## # fit different distributions
## mga <- gamlss(Ozone~Temp+Wind+Solar.R, data=da, family=GA)</pre>
## mig <- gamlss(Ozone~Temp+Wind+Solar.R, data=da, family=IG)</pre>
## mbccg <- gamlss(Ozone~Temp+Wind+Solar.R, data=da, family=BCCGo)
## GAIC(mno, mga, mig, mbccg)
## # fit smoothers
## mga1=gamlss(Ozone~pb(Temp)+pb(Wind)+pb(Solar.R),data=da,
              family=GA)
## term.plot(mga1, pages=1)
## plot(mga1)
## wp(mga1)
```

1.2 gamlss.demo()

```
#install.packages("gamlss.demo", dep=TRUE)
#Based on rpanel
```

```
library(gamlss.demo)
#Examples
#t family distribution
demo.TF()
\#Skew\ Normal\ Type\ 1\ distribution
demo.SN1()
\#Box-Cox\ Power\ Exponential\ distribution
demo.BCPE()
#Demos for smoothing techniques
demo.BSplines()
demo.PSplines()
demo.interpolateSmo()
demo.histSmo()
#Interface for demonstrating the gamlss.family distributions
demoDist()
#Demo for local polynomial smoothing
demoLpolyS()
#The demo for gamlss distributions and smoothing
gamlss.demo()
```