

1 Modified R Functions from GAMLSS

Read in the NHANES data

```
require(epicalc, warn.conflicts = FALSE)

## Loading required package: epicalc
## Loading required package: foreign
## Loading required package: survival
## Loading required package: splines
## Loading required package: MASS
## Loading required package: nnet

require(gamlss, warn.conflicts = FALSE)

## Loading required package: gamlss
## Loading required package: gamlss.data
## Loading required package: gamlss.dist
## Loading required package: nlme
## ***** GAMLSS Version 4.3-0 *****
## For more on GAMLSS look at http://www.gamlss.org/
## Type gamlssNews() to see new features/changes/bug fixes.

require(lattice, warn.conflicts = FALSE)

## Loading required package: lattice

require(car, warn.conflicts = FALSE)

## Loading required package: car

require(xtable, warn.conflicts = FALSE)

## Loading required package: xtable

require(foreign, warn.conflicts = FALSE)
```

2 Modified Functions from GAMLSS Package

```
mycent <- function(obj, xvar = NULL, cent = c(1, 3, 5, 15, 25, 50, 75, 85, 95,
  97, 99), legend = TRUE, ylab = "y", xlab = "x", main = NULL, main.gsub = "@",
  xleg = min(xvar), yleg = max(obj$y), xlim = range(xvar), ylim = range(obj$y),
  save = FALSE, plot = TRUE, points = TRUE, pch = "+", col = "blue", col.centiles = 1:10,
  2, lty.centiles = 1, lwd.centiles = 1, ...) {
  if (!is.gamlss(obj))
    stop(paste("This is not an gamlss object", "\n", ""))
  if (is.null(xvar))
    stop(paste("The xvar argument is not specified", "\n", ""))
  fname <- obj$family[1]
  qfun <- paste("q", fname, sep = "")
  Title <- paste("Centile curves using", fname, sep = " ")
```

```

main <- if (is.null(main))
  paste("Centile curves using", fname, sep = " ") else gsub(main.gsub, Title, main)
oxvar <- xvar[order(xvar)]
oyvar <- obj$y[order(xvar)]
if (is.matrix(obj$y)) {
  oyvar <- obj$y[, 1][order(xvar)]
  ylim <- range(obj$y[, 1])
  yleg = max(obj$y[, 1])
}
if (plot) {
  lty.centiles <- rep(lty.centiles, length(cent))
  lwd.centiles <- rep(lwd.centiles, length(cent))
  col.centiles <- rep(col.centiles, length(cent))
  if (points == TRUE) {
    plot(oxvar, oyvar, type = "p", col = col, pch = pch, xlab = xlab,
         ylab = ylab, xlim = xlim, ylim, ...)
  } else {
    plot(oxvar, oyvar, type = "n", col = col, pch = pch, xlab = xlab,
         ylab = ylab, xlim = xlim, ylim, ...)
  }
  title(main)
}
col <- 3
lpar <- length(obj$parameters)
ii <- 0
per <- rep(0, length(cent))
for (var in cent) {
  if (lpar == 1) {
    newcall <- call(qfun, var/100, mu = fitted(obj, "mu")[order(xvar)])
  } else if (lpar == 2) {
    newcall <- call(qfun, var/100, mu = fitted(obj, "mu")[order(xvar)],
                   sigma = fitted(obj, "sigma")[order(xvar)])
  } else if (lpar == 3) {
    newcall <- call(qfun, var/100, mu = fitted(obj, "mu")[order(xvar)],
                   sigma = fitted(obj, "sigma")[order(xvar)], nu = fitted(obj,
                   "nu")[order(xvar)])
  } else {
    newcall <- call(qfun, var/100, mu = fitted(obj, "mu")[order(xvar)],
                   sigma = fitted(obj, "sigma")[order(xvar)], nu = fitted(obj,
                   "nu")[order(xvar)], tau = fitted(obj, "tau")[order(xvar)])
  }
  ii <- ii + 1
  ll <- eval(newcall)
  if (plot) {
    lines(oxvar, ll, col = col.centiles[ii], lty = lty.centiles[ii],
         lwd = lwd.centiles[ii], ...)
  }
  per[ii] <- (1 - sum(oyvar > ll)/length(oyvar)) * 100
  if (!save)
    cat("% of cases below ", var, "centile is ", per[ii], "\n")
}

```

```

}
if (plot) {
  if (legend == TRUE)
    legend(list(x = xleg, y = yleg), legend = cent, col = col.centiles,
             lty = lty.centiles, lwd = lwd.centiles, ncol = 1, ...)
}
if (save) {
  # return(cbind(cent, per))
  TAU = tau
  L = nu
  Median = mu
  S = sigma
  return(cbind(L, Median, S, cent, per))
}
}

```

```

mycent.pred <- function(obj, type = c("centiles", "z-scores", "standard-centiles"),
  xname = NULL, xvalues = NULL, power = NULL, yval = NULL, cent = c(1, 3,
    5, 15, 25, 50, 75, 85, 95, 97, 99), dev = c(-4, -3, -2, -1, 0, 1, 2,
    3, 4), plot = FALSE, legend = TRUE, ...) {
  calc.cent <- function(xvar, cent) {
    o <- order(xvar)
    mat <- xvar[o]
    cent <- cent
    for (var in cent) {
      if (lpar == 1) {
        newcall <- call(qfun, var/100, mu = mu[o])
      } else if (lpar == 2) {
        newcall <- call(qfun, var/100, mu = mu[o], sigma = sigma[o])
      } else if (lpar == 3) {
        newcall <- call(qfun, var/100, mu = mu[o], sigma = sigma[o],
          nu = nu[o])
      } else {
        newcall <- call(qfun, var/100, mu = mu[o], sigma = sigma[o],
          nu = nu[o], tau = tau[o])
      }
      ll <- eval(newcall)
      mat <- cbind(mat, ll)
    }
    mat <- as.data.frame(mat)
    nnn <- paste("C", as.character(cent), sep = "")
    names(mat) <- c(xname, nnn)
    return(mat)
  }
  plot.mat <- function(mat, cent, legend, ...) {
    lcent <- dim(mat)[2]
    xleg <- min(mat[, 1])
    yleg <- max(mat[, 2:lcent])
    plot(mat[, 1], mat[, 2], type = "n", ...)
    for (i in 2:lcent) lines(mat[, 1], mat[, i], col = i)
  }
}

```

```

    if (legend)
      legend(list(x = xleg, y = yleg), legend = cent, col = c(2, 3, 4,
        5, 6, 7, 8, 9, 10, 11, 12), lty = 1, ncol = 1, bg = "white")
    invisible()
  }
  if (!is.gamlss(obj))
    stop(paste("This is not an gamlss object", "\n", ""))
  if (is.null(xvalues))
    stop(paste("The xvalues argument is not specified", "\n", ""))
  if (is.null(xname))
    stop(paste("The xname argument is not specified", "\n", ""))
  if (!is.character(xname))
    stop(paste("The xname argument is not a character", "\n", ""))
  xvar <- if (!is.null(power))
    xvar <- xvalues^power else xvalues
  newx <- data.frame(xvar)
  colnames(newx) <- xname
  lpar <- length(obj$parameters)
  if ("mu" %in% obj$parameters) {
    if (is.null(obj$mu.fix))
      mu <- predict(obj, what = "mu", newdata = newx, type = "response",
        ...) else if (obj$mu.fix == TRUE)
      mu <- rep(fitted(obj, "mu")[1], length(xvar))
  }
  if ("sigma" %in% obj$parameters) {
    if (is.null(obj$sigma.fix))
      sigma <- predict(obj, what = "sigma", newdata = newx, type = "response",
        ...) else if (obj$sigma.fix == TRUE)
      sigma <- rep(fitted(obj, "sigma")[1], length(xvar))
  }
  if ("nu" %in% obj$parameters) {
    if (is.null(obj$nu.fix))
      nu <- predict(obj, what = "nu", newdata = newx, type = "response",
        ...) else if (obj$nu.fix == TRUE)
      nu <- rep(fitted(obj, "nu")[1], length(xvar))
  }
  if ("tau" %in% obj$parameters) {
    if (is.null(obj$tau.fix))
      tau <- predict(obj, what = "tau", newdata = newx, type = "response",
        ...) else if (obj$tau.fix == TRUE)
      tau <- rep(fitted(obj, "tau")[1], length(xvar))
  }
  type <- match.arg(type)
  if (type == "centiles") {
    fname <- obj$family[1]
    qfun <- paste("q", fname, sep = "")
    xvar <- xvalues
    mat <- calc.cent(xvar = xvar, cent = cent)
    if (plot)
      plot.mat(mat, cent, legend, ...)
  }

```

```

    # return(mat)
    TAU = tau
    L = nu
    Median = mu
    S = sigma
    Age <- mat[, 1]
    return(cbind(Age, L, Median, S, mat[, -1]))
}
if (type == "z-scores") {
  if (is.null(yval))
    stop("the y values should be set if type=z-scores is used")
  if (length(yval) != length(xvalues))
    stop("length of xvalues and yval is not the same")
  fname <- obj$family[1]
  qfun <- paste("p", fname, sep = "")
  if (lpar == 1) {
    newcall <- call(qfun, yval, mu = mu)
  } else if (lpar == 2) {
    newcall <- call(qfun, yval, mu = mu, sigma = sigma)
  } else if (lpar == 3) {
    newcall <- call(qfun, yval, mu = mu, sigma = sigma, nu = nu)
  } else {
    newcall <- call(qfun, yval, mu = mu, sigma = sigma, nu = nu, tau = tau)
  }
  cdf <- eval(newcall)
  rqres <- qnorm(cdf)
  return(rqres)
}
if (type == "standard-centiles") {
  cent <- pnorm(dev) * 100
  fname <- obj$family[1]
  qfun <- paste("q", fname, sep = "")
  xvar <- xvalues
  mat <- calc.cent(xvar = xvar, cent = cent)
  nnn <- paste(as.character(dev), sep = "")
  names(mat) <- c(xname, nnn)
  if (plot)
    plot.mat(mat, dev, legend, ...)
  # return(mat)
  TAU = tau
  L = nu
  Median = mu
  S = sigma
  Age <- mat[, 1]
  return(cbind(Age, L, Median, S, mat[, -1]))
}
}

```

3 Reading NHANES data

```
d2 <- read.dta("hes2hes3nhanes1.dta")
attach(d2)
des(d2)

##
## No. of observations = 23118
## Variable      Class
## 1 muac         integer
## 2 muac_left_arm integer
## 3 dataset      character
## 4 agey         numeric
## 5 sex          factor
## 6 muac_female  integer
## 7 muac_male    integer
## 8 l_r_diff     numeric
## Description
## 1 R arm was standard measure'
## 2 subset left arm measured
## 3
## 4 age calculated from dob/date (NHANES) or months of age (HES 2 &3)
## 5
## 6 muac, sex == female
## 7 muac, sex == male'
## 8 difference between R and L arm

str(d2)

## 'data.frame': 23118 obs. of 8 variables:
## $ muac : int 151 173 152 125 159 160 151 164 142 162 ...
## $ muac_left_arm: int NA NA NA NA NA NA NA NA 166 NA NA ...
## $ dataset : chr "NHANES 1" "NHANES 1" "NHANES 1" "NHANES 1" ...
## $ agey : num 1.98 1.98 1.99 1.99 1.99 ...
## $ sex : Factor w/ 2 levels "female","male": 2 2 2 1 1 1 2 1 2 1 ...
## $ muac_female : int NA NA NA 125 159 160 NA 164 NA 162 ...
## $ muac_male : int 151 173 152 NA NA NA 151 NA 142 NA ...
## $ l_r_diff : num NA NA NA NA NA NA NA NA -2 NA NA ...
## - attr(*, "datalabel")= chr ""
## - attr(*, "time.stamp")= chr "23 Mar 2011 10:11"
## - attr(*, "formats")= chr "%9.0g" "%9.0g" "%9s" "%9.0g" ...
## - attr(*, "types")= int 252 252 8 254 251 252 252 254
## - attr(*, "val.labels")= chr "" "" "" "" ...
## - attr(*, "var.labels")= chr "R arm was standard measure'" "subset left arm measured"
## - attr(*, "expansion.fields")=List of 2
## ..$ : chr "_dta" "_lang_c" "default"
## ..$ : chr "_dta" "_lang_list" "default"
## - attr(*, "version")= int 12
## - attr(*, "label.table")=List of 1
## ..$ sex: Named int 1 2
## .. ..- attr(*, "names")= chr "female" "male"
```

```
summ(d2)

##
##
## No. of observations = 23118
##
##   Var. name      obs.   mean   median  s.d.   min.   max.
## 1 muac           23118 227.29  222     50.35  59     499
## 2 muac_left_arm 1788   229.82 226     57.41  107    485
## 3 dataset
## 4 agey           23118 12.29   12.08   5.34   1.98   26.02
## 5 sex            23118 1.491   1       0.5    1      2
## 6 muac_female    11761 226.35 224     48.22  59     499
## 7 muac_male      11357 228.26 220     52.45  105    483
## 8 l_r_diff       1788   1.62   1       8.48   -92    110
```

Convert muac to cm and age to months. Drop if age is less than 5 and greater than 25

```
d2$muac <- d2$muac/10 # convert to mm
d2$agem <- d2$agey * 12 # convert age to months
summ(d2)

##
##
## No. of observations = 23118
##
##   Var. name      obs.   mean   median  s.d.   min.   max.
## 1 muac           23118 22.73   22.2    5.03   5.9    49.9
## 2 muac_left_arm 1788   229.82 226     57.41  107    485
## 3 dataset
## 4 agey           23118 12.29   12.08   5.34   1.98   26.02
## 5 sex            23118 1.491   1       0.5    1      2
## 6 muac_female    11761 226.35 224     48.22  59     499
## 7 muac_male      11357 228.26 220     52.45  105    483
## 8 l_r_diff       1788   1.62   1       8.48   -92    110
## 9 agem           23118 147.53 145     64.07  23.75  312.25
```

Keep data only for ages 2-25 years

```
d2 = d2[d2$agem >= 24 & d2$agem <= 300, c(5, 1, 9)]
summ(d2)

##
##
## No. of observations = 22699
##
##   Var. name  obs.   mean   median  s.d.   min.   max.
## 1 sex        22699 1.494   1       0.5    1      2
## 2 muac       22699 22.62   22.1    4.98   5.9    49.9
## 3 agem       22699 144.72 143     60.92  24.05  299.99

tab1(d2$sex, graph = FALSE)
```

```
## d2$sex :
##           Frequency Percent Cum. percent
## female      11481     50.6         50.6
## male'       11218     49.4         100.0
##   Total      22699    100.0         100.0

d2$sex <- factor(d2$sex, labels = c("Female", "Male"))
tab1(d2$sex, graph = FALSE)

## d2$sex :
##           Frequency Percent Cum. percent
## Female      11481     50.6         50.6
## Male       11218     49.4         100.0
##   Total      22699    100.0         100.0
```

3.1 Create two datasets, one for boys and the other for girls

```
boysd = d2[d2$sex == "Male", ]
des(boysd)

##
## No. of observations = 11218
##   Variable      Class      Description
## 1 sex          factor
## 2 muac         numeric
## 3 agem         numeric

summ(boysd)

##
##
## No. of observations = 11218
##
##   Var. name obs.  mean  median  s.d.  min.  max.
## 1 sex        11218  2      2      0     2     2
## 2 muac       11218 22.73  21.8   5.18  10.5  48.3
## 3 agem       11218 140.58 140.25 57.07 24.11 299.93

girlsd = d2[d2$sex == "Female", ]
des(girlsd)

##
## No. of observations = 11481
##   Variable      Class      Description
## 1 sex          factor
## 2 muac         numeric
## 3 agem         numeric

summ(girlsd)
```



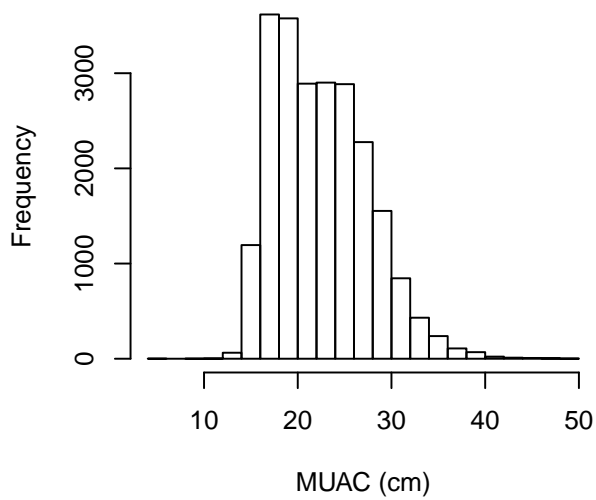
```
##
##
## No. of observations = 11481
##
##   Var. name obs.   mean   median  s.d.   min.   max.
## 1 sex         11481 1       1       0     1     1
## 2 muac        11481 22.51  22.3    4.77   5.9    49.9
## 3 agem        11481 148.76 145.28 64.22  24.05  299.99
```

4 Fitting models, calculating Z scores

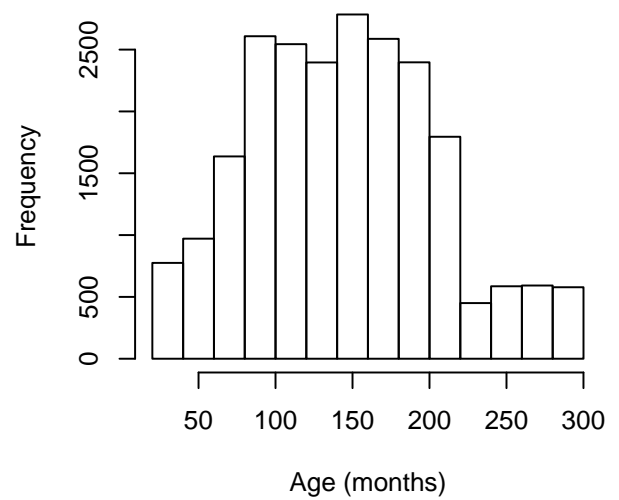
4.1 Data Explanatory Analysis

```
X11(height = 8, width = 8)
par(mfrow = c(2, 2))
hist(d2$muac, main = "MUAC distribution", xlab = "MUAC (cm)")
hist(d2$agem, main = "Age distribution", xlab = "Age (months)")
plot(muac ~ agem, data = girlsd, ylab = "MUAC (cm)", xlab = "Age (months)",
     pch = "x", col = "grey", main = "Girls data", ylim = c(0, 50))
plot(muac ~ agem, data = boysd, ylab = "MUAC (cm)", xlab = "Age (months)", pch = "x",
     col = "grey", main = "Boys data", ylim = c(0, 50))
```

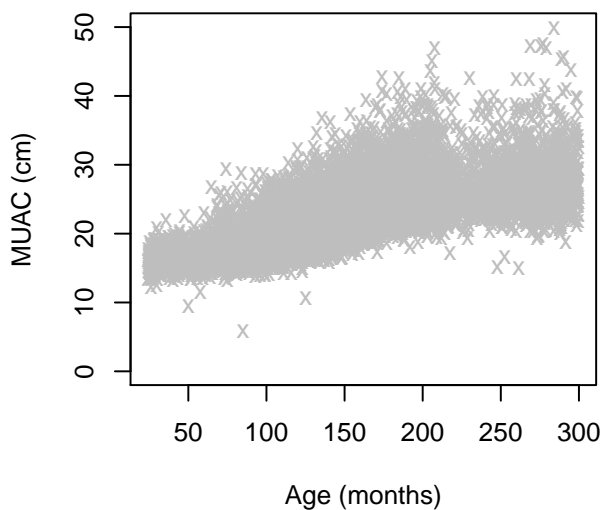
MUAC distribution



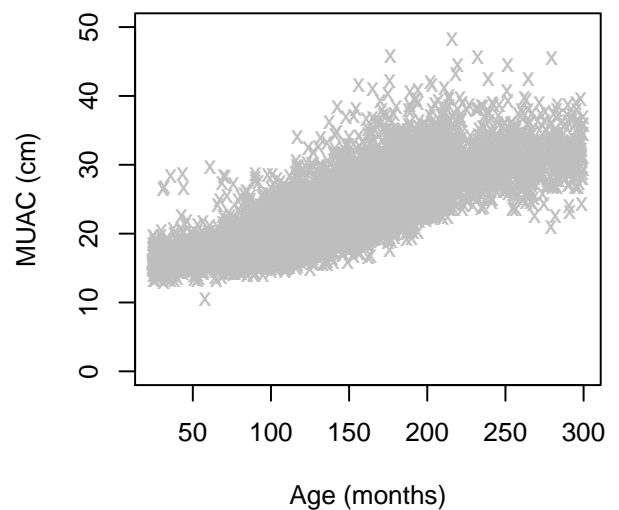
Age distribution



Girls data



Boys data

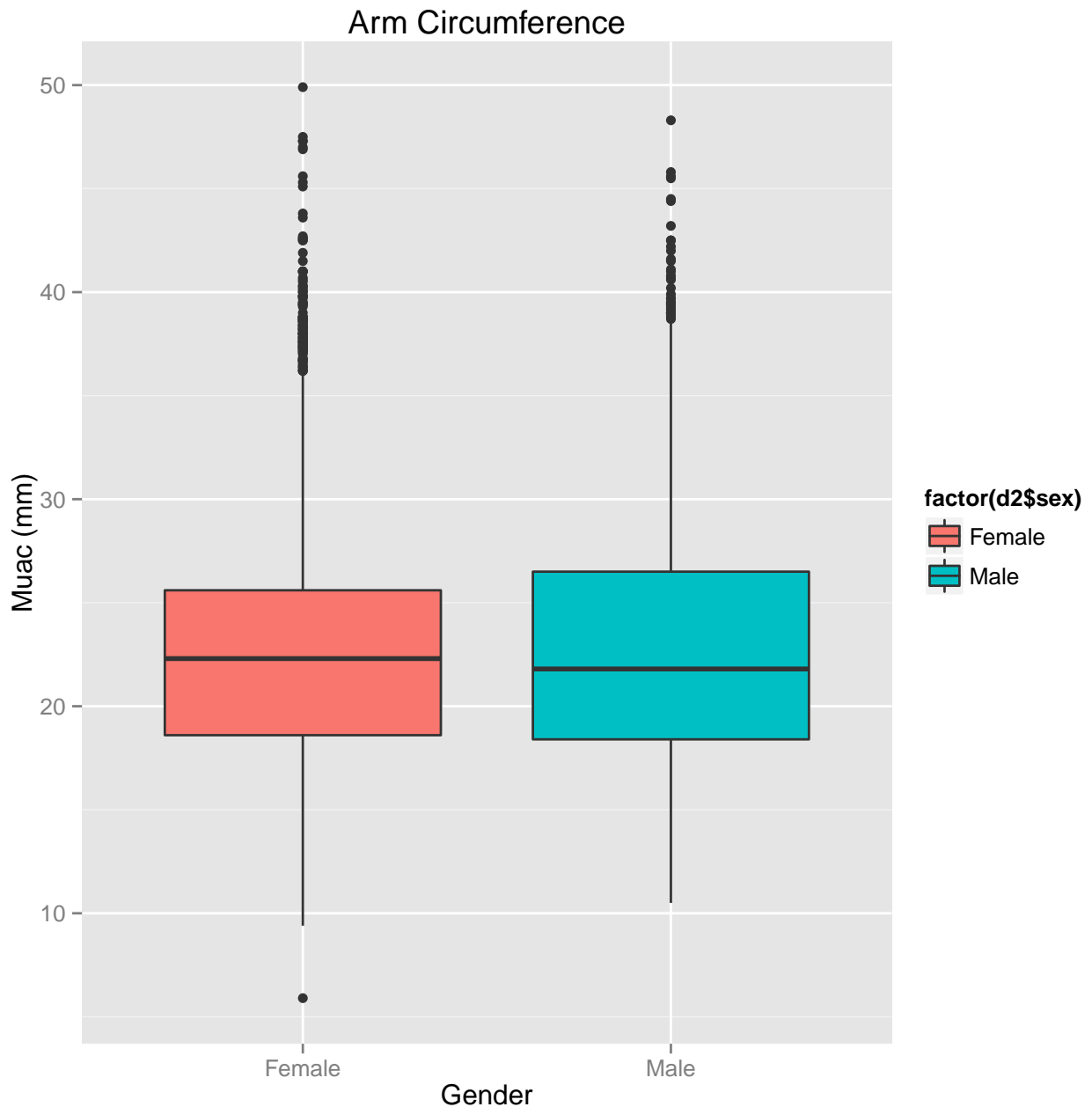


```
par(mfrow = c(1, 1))
```

```
require(ggplot2)
```

```
## Loading required package: ggplot2
```

```
qplot(factor(d2$sex), d2$muac, data = d2, geom = c("boxplot"), main = "Arm Circumference",  
       fill = factor(d2$sex), xlab = "Gender", ylab = "Muac (mm)")
```



4.2 Model Selection

Boys model selection

Note that BCT has been selected as the best other than BCCG and BCPE.

4.3 Generating Z-scores

Generating Z-scores for both of the datasets separately starting with the female dataset.

```
# Girls model selection and fitting the model for boys
hf1 <- gamlss(muac ~ cs(agem), sigma.fo = ~cs(agem), nu.fo = ~cs(agem), tau.fo = ~cs(agem),
  data = girlsd, family = BCT)

## GAMLSS-RS iteration 1: Global Deviance = 55080
## GAMLSS-RS iteration 2: Global Deviance = 55026
## GAMLSS-RS iteration 3: Global Deviance = 55025
```

```

## GAMLSS-RS iteration 4: Global Deviance = 55025
## GAMLSS-RS iteration 5: Global Deviance = 55025
## GAMLSS-RS iteration 6: Global Deviance = 55025
## GAMLSS-RS iteration 7: Global Deviance = 55025
## GAMLSS-RS iteration 8: Global Deviance = 55025
## GAMLSS-RS iteration 9: Global Deviance = 55025
## GAMLSS-RS iteration 10: Global Deviance = 55025
## GAMLSS-RS iteration 11: Global Deviance = 55025
## GAMLSS-RS iteration 12: Global Deviance = 55025
## GAMLSS-RS iteration 13: Global Deviance = 55025

hf0 <- gamlss(muac ~ cs(agem), sigma.fo = ~cs(agem), nu.fo = ~cs(agem), tau.fo = ~cs(agem),
  data = girlsd, family = BCPE)

## GAMLSS-RS iteration 1: Global Deviance = 55585
## GAMLSS-RS iteration 2: Global Deviance = 55124
## GAMLSS-RS iteration 3: Global Deviance = 55111
## GAMLSS-RS iteration 4: Global Deviance = 55111
## GAMLSS-RS iteration 5: Global Deviance = 55111
## GAMLSS-RS iteration 6: Global Deviance = 55112
## GAMLSS-RS iteration 7: Global Deviance = 55112
## GAMLSS-RS iteration 8: Global Deviance = 55112
## GAMLSS-RS iteration 9: Global Deviance = 55112

AIC(hf1, hf0, k = 3)

##      df    AIC
## hf1 20 55085
## hf0 20 55172

GAIC(hf1, hf0, k = 3)

##      df    AIC
## hf1 20 55085
## hf0 20 55172

```