

Quantile Regression - Chapter 12 on Handbook

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Quantile Regression

```
library(gamlss.data)  
library(lattice)  
library("quantreg")
```

Introduction

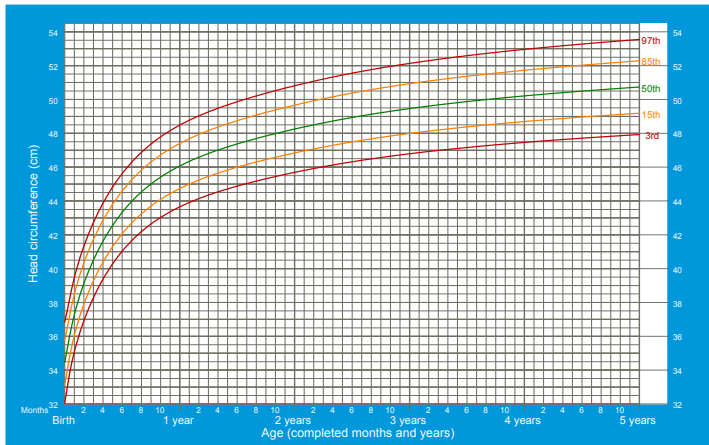
During ultrasound examination of an as-yet-unborn baby, anthropometric measurements are taken.

- ▶ For a given gestational age one can directly compare, say the femur length of the examined fetus with the femur length of all fetuses in the reference population.
- ▶ Too small or too large values may indicate developmental problems and require an intervention.
- ▶ From a statistical point of view : what does *too small* or *too large* mean?

Head Circumference for Age - WHO

Head circumference-for-age **BOYS**

Birth to 5 years (percentiles)



WHO Child Growth Standards

Figure 1: Boys Head Circumference Quantiles

Head Circumference for Age

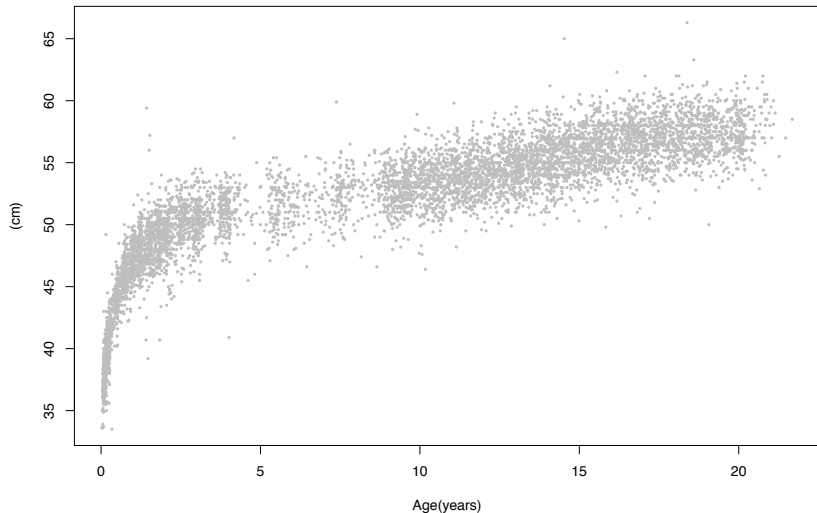
- ▶ The data contains head circumference for boys older than 24 months
- ▶ Aim: to construct a *growth chart*
- ▶ *i.e* Conditional distribution of head circumference given age.
- ▶ Age specific quantiles tells us how many boys in the reference population have a smaller head circumference compared to the single boy a physician is looking at.
- ▶ Quantile regression - method to estimate conditional quantiles

Head Circumference for Age

```
#library(gamlss.data)  
data(db)  
head(db)  
dim(db)  
plot(db$head ~ db$age, xlab = "Age(months)",  
      ylab = "Head circumference",  
      pch = 16, cex = 0.5, col = "gray")
```

Head Circumference for Age

```
## head age
## 1 33.6 0.03
## 2 33.6 0.04
## 3 33.7 0.04
```



Common regression models

- ▶ To date, our *Linear* or *additive models* have focused on describing the conditional **mean**, $E(y|x_1, x_2, \dots, x_q)$ of the response y as a linear or additive function of the explanatory variables x_1, x_2, \dots, x_q .
- ▶ For non normal response - link function of the conditional mean is modeled
 - ▶ $g(E(y|x_1, x_2, \dots, x_q))$
- ▶ Therefore for a linear model it follows that
 - ▶ $y \sim N(\alpha + \beta_1 x_1 + \dots + \beta_q x_q, \sigma^2)$
 - ▶ The conditional $\tau \times 100\%$ quantile for y is $\alpha + \beta_1 x_1 + \dots + \beta_q x_q + \sigma u_\tau$, where u_τ is the $\tau \times 100\%$ quantile for standard normal
 - ▶ For skewed or non-normal distribution the corresponding quantile will be misleading

Linear Quantile Regression

Simple linear quantile regression model (Koenker and Bassett, 1978)

$$y_i = \alpha_\tau + \beta_\tau x_i + \epsilon_{\tau i}$$

where $\epsilon_{\tau i} \sim F_{\tau i}$, $i = 1, \dots, n$, subject to $F_{\tau i}(0) = \tau$.

- ▶ α_τ and β_τ are the intercept and slope effects and $\tau \in (0, 1)$ is a fixed-known quantile
- ▶ $F_{\tau i}$ has no specific distributional assumption except that the distribution function at 0 is τ
- ▶ Equivalent to $Q_{y_i}(\tau|x_i) = F_{y_i}^{-1}(\tau|x_i) = \alpha_\tau + \beta_\tau x_i$

Linear Quantile Regression

Minimization problem

- ▶ $\operatorname{argmin}_{\alpha_{\tau}\beta_{\tau}} \sum_{i=1}^n \rho_{\tau}(y_i - (\alpha_{\tau} + \beta_{\tau}x_i))$
 - ▶ where $\rho_{\tau}(z) = z\tau$ for $z \geq 0$ and $z(\tau - 1)$ for $z < 0$

For median $\tau = 0.5$, $\rho_{0.5}(z) \propto |z|$ therefore

- ▶ $\operatorname{argmin}_{\alpha_{\tau}\beta_{\tau}} \sum_{i=1}^n |y_i - (\alpha_{\tau} + \beta_{\tau}x_i)|$

The minimization problem above is formulated as a set of linear constraints and estimation of parameters is conducted by linear programming. This will lead to the $\tau \times 100\%$ quantiles of the response variable

- ▶ Compare with $\operatorname{argmin}_{\alpha\beta} \sum_{i=1}^n |y_i - (\alpha + \beta x_i)|^2$ for simple linear regression
- ▶ Quantile regression is more robust towards extreme outliers as compared to least square regression

Additive Quantile Regression

- ▶ For cases where non-linear relationship between explanatory variables and quantiles of the response variable
- ▶ $Q_{y_i}(\tau|x_i) = f_\tau(x_i)$,
 - ▶ where f is a smooth function of x
- ▶ The minimization problem is extended by a penalty term to

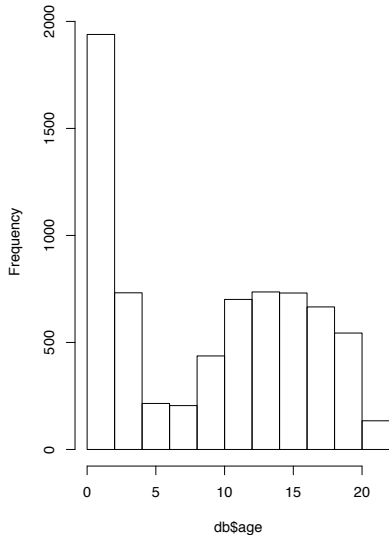
$$\operatorname{argmin}_{f_\tau} \sum_{i=1}^n \rho_\tau(y_i - f_\tau(x_i)) + \lambda V(f'_\tau),$$

- ▶ where $V(f'_\tau) = \sup \sum |f'_\tau(x_i + 1) - f'_\tau(x_i)|$ - the total variation of f'_τ and λ is the tuning parameter
- ▶ Solutions are obtained using linear programming (See Koenker et al. 1994 and Koenker 2005)

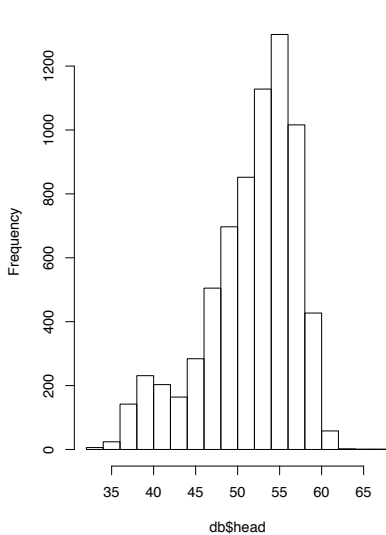
Dutch boys head circumference

```
layout(matrix(1:2, nrow = 1))  
hist(db$age)  
hist(db$head)
```

Histogram of db\$age

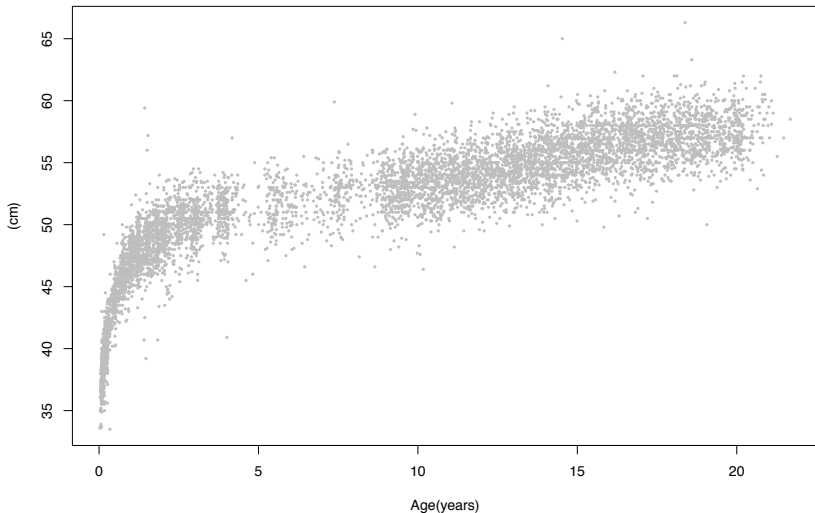


Histogram of db\$head



Head Circumference for Age

```
## head age
## 1 33.6 0.03
## 2 33.6 0.04
## 3 33.7 0.04
```



Dutch boys head circumference

```
db <- db[db$age>2,] # subset data by age>2  
summary(db)
```

##	head	age
##	Min. :40.90	Min. : 2.01
##	1st Qu.:52.30	1st Qu.: 9.08
##	Median :54.50	Median :12.76
##	Mean :54.34	Mean :12.02
##	3rd Qu.:56.50	3rd Qu.:16.23
##	Max. :66.30	Max. :21.68

Dutch boys head circumference

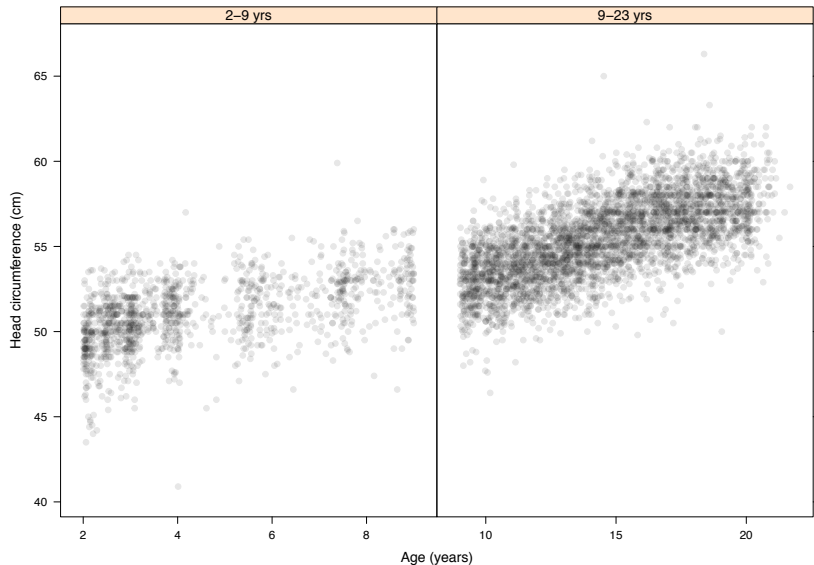
#add a cut variable in data to subset data

```
db$cut <- cut(db$age, breaks = c(2, 9, 23),  
  labels = c("2-9 yrs", "9-23 yrs"))
```

#different scatterplot by age group

```
xyplot(head ~ age | cut, data = db, xlab = "Age (years)",  
  ylab = "Head circumference (cm)",  
  scales = list(x = list(relation = "free")),  
  layout = c(2, 1), pch = 19,  
  col = rgb(.1, .1, .1, .1))
```

Dutch boys head circumference



Dutch boys head circumference

Simple linear regression model by age group

```
lm2.9 <- lm(head ~ age, data = db, subset = age < 9)  
lm2.9$coef
```

```
lm9.23 <- lm(head ~ age, data = db, subset = age > 9)  
lm9.23$coef
```

Equivalent to

```
lm_mod <- lm(head ~ age:I(age < 9) + I(age < 9) - 1,  
  data = db)  
lm_mod$coef
```

Under the normal assumption the mean is equal to median hence the models can be interpreted as conditional median models under normal assumption

Dutch boys head circumference

Simple linear regression model by age group

```
## (Intercept)          age
##  48.9233698    0.4734876
```

```
## (Intercept)          age
##  48.6194278    0.4689793
```

```
##      I(age < 9)FALSE      I(age < 9)TRUE age:I(age < 9)FALSE
##      48.6201100      48.9233698      0.4689376
## age:I(age < 9)TRUE
##      0.4734876
```

The model states that within one year, the **average** head circumference for boys less than nine years old increases by 0.473 cm and by 0.469 for older boys.

Dutch boys head circumference - rq

- ▶ Relax the distributional assumption (use **rq** function)
- ▶ Conditional median ($\tau = 0.5$)

```
rq_med2.9 <- rq(head ~ age, data = db, tau = 0.5,  
subset = age < 9)  
rq_med2.9$coef
```

```
## (Intercept)          age  
##  48.9282511    0.4932735
```

```
rq_med9.23 <- rq(head ~ age, data = db, tau = 0.5,  
subset = age > 9)  
rq_med9.23$coef
```

```
## (Intercept)          age  
##  48.5791795    0.4717949
```

Dutch boys head circumference - lm vs rq

- Calculate Confidence intervals for the intercept and slope:- Younger boys confidence interval : similar intercept but different slopes using *lm* vs *rq*

```
cbind(coef(lm2.9)[1], confint(lm2.9, parm = "(Intercept)"))
```

```
##                2.5 %    97.5 %  
## (Intercept) 48.92337 48.70166 49.14508
```

```
cbind(coef(lm2.9)[2], confint(lm2.9, parm = "age"))
```

```
##                2.5 %    97.5 %  
## age 0.4734876 0.4282969 0.5186783
```

```
options(warn=-1) # turns off warning message  
summary(rq_med2.9, se = "rank")$coef
```

```
##                coefficients    lower bd    upper bd  
## (Intercept)    48.9282511 48.7567664 49.1160521  
## age            0.4932735  0.4326066  0.5493336
```

```
options(warn=0) # turns on warning message
```

Dutch boys head circumference - lm vs rq

- Calculate Confidence intervals for the intercept and slope:- Older boys confidence interval : similar intercept but different slopes using *lm* vs *rq*

```
cbind(coef(lm9.23)[1], confint(lm9.23, parm = "(Intercept)"))
```

```
##                2.5 %    97.5 %  
## (Intercept) 48.61943 48.36341 48.87545
```

```
cbind(coef(lm9.23)[2], confint(lm9.23, parm = "age"))
```

```
##                2.5 %    97.5 %  
## age 0.4689793 0.4517425 0.4862161
```

```
options(warn=-1) # turns off warning message  
summary(rq_med9.23, se = "rank")$coef
```

```
##                coefficients    lower bd    upper bd  
## (Intercept)  48.5791795 48.3907933 48.8928025  
## age          0.4717949 0.4299378 0.4858946
```

```
options(warn=0) # turns on warning message
```

Dutch boys head circumference - growth curve lm

- ▶ Use linear model for construction of growth curves
- ▶ Based on the normal linear models, we can compute the quantiles of head circumference for age.
- ▶ Here we consider the following values of τ

```
tau <- c(.01, .1, .25, .5, .75, .9, .99)
gage <- c(2:9, 9:23)
i <- 1:8
idf <- data.frame(age = gage[i])
p <- predict(lm2.9, newdata = idf, level = 0.5,
  interval = "prediction") # level - coverage
colnames(p) <- c("0.5", "0.25", "0.75")
p
```

Dutch boys head circumference - growth curve lm

##		0.5	0.25	0.75
## 1	49.87034	48.69777	51.04292	
## 2	50.34383	49.17165	51.51602	
## 3	50.81732	49.64533	51.98931	
## 4	51.29081	50.11880	52.46282	
## 5	51.76430	50.59206	52.93653	
## 6	52.23778	51.06512	53.41044	
## 7	52.71127	51.53797	53.88457	
## 8	53.18476	52.01062	54.35889	

Dutch boys head circumference

Find 80% and 98% prediction intervals

```
p <- cbind(p, predict(lm2.9, newdata = idf, level = 0.8,  
  interval = "prediction")[,-1])  
colnames(p)[4:5] <- c("0.1", "0.9")  
p <- cbind(p, predict(lm2.9, newdata = idf, level = 0.98,  
  interval = "prediction")[,-1])  
colnames(p)[6:7] <- c("0.01", "0.99")  
p2.9 <- p[, c("0.01", "0.1", "0.25", "0.5",  
  "0.75", "0.9", "0.99")]  
head(p2.9)
```

##		0.01	0.1	0.25	0.5	0.75	0.9	0.99
## 1	45.82205	47.64188	48.69777	49.87034	51.04292	52.09881	53.91864	
## 2	46.29691	48.11612	49.17165	50.34383	51.51602	52.57155	54.39076	
## 3	46.77105	48.58997	49.64533	50.81732	51.98931	53.04467	54.86359	
## 4	47.24448	49.06342	50.11880	51.29081	52.46282	53.51819	55.33713	
## 5	47.71720	49.53649	50.59206	51.76430	52.93653	53.99210	55.81139	
## 6	48.18921	50.00916	51.06512	52.23778	53.41044	54.46640	56.28636	

Dutch boys head circumference

Repeat the same for older boys

```
idf <- data.frame(age = gage[-i])
p <- predict(lm9.23, newdata = idf, level = 0.5,
interval = "prediction")
colnames(p) <- c("0.5", "0.25", "0.75")
p <- cbind(p, predict(lm9.23, newdata = idf, level = 0.8,
interval = "prediction"), -1)
colnames(p)[4:5] <- c("0.1", "0.9")
p <- cbind(p, predict(lm9.23, newdata = idf, level = 0.98,
interval = "prediction"), -1)
colnames(p)[6:7] <- c("0.01", "0.99")
p9.23 <- p[, c("0.01", "0.1", "0.25", "0.5",
"0.75", "0.9", "0.99")]
p9.23
```

Dutch boys head circumference

Quantiles for older boys

##		0.01	0.1	0.25	0.5	0.75	0.9	0.99
## 1		48.78475	50.60668	51.66479	52.84024	54.01569	55.07381	56.89574
## 2		49.25424	51.07594	52.13392	53.30922	54.48452	55.54250	57.36420
## 3		49.72363	51.54515	52.60302	53.77820	54.95338	56.01125	57.83277
## 4		50.19292	52.01430	53.07209	54.24718	55.42227	56.48006	58.30143
## 5		50.66211	52.48339	53.54113	54.71616	55.89119	56.94893	58.77021
## 6		51.13119	52.95243	54.01014	55.18514	56.36014	57.41785	59.23908
## 7		51.60017	53.42141	54.47912	55.65412	56.82912	57.88683	59.70806
## 8		52.06905	53.89033	54.94807	56.12310	57.29813	58.35586	60.17714
## 9		52.53782	54.35919	55.41699	56.59208	57.76717	58.82496	60.64633
## 10		53.00649	54.82800	55.88588	57.06106	58.23624	59.29411	61.11562
## 11		53.47506	55.29676	56.35473	57.53003	58.70533	59.76331	61.58501
## 12		53.94352	55.76545	56.82356	57.99901	59.17446	60.23258	62.05450
## 13		54.41188	56.23409	57.29236	58.46799	59.64362	60.70190	62.52410
## 14		54.88014	56.70267	57.76113	58.93697	60.11281	61.17127	62.99380
## 15		55.34830	57.17120	58.22988	59.40595	60.58203	61.64071	63.46361

Dutch boys head circumference

Conditional quantiles estimated under the normal assumption of head circumference

```
q2.23 <- rbind(p2.9, p9.23)  
head(round(q2.23, 3), n = 14)
```

##		0.01	0.1	0.25	0.5	0.75	0.9	0.99
## 1	45.822	47.642	48.698	49.870	51.043	52.099	53.919	
## 2	46.297	48.116	49.172	50.344	51.516	52.572	54.391	
## 3	46.771	48.590	49.645	50.817	51.989	53.045	54.864	
## 4	47.244	49.063	50.119	51.291	52.463	53.518	55.337	
## 5	47.717	49.536	50.592	51.764	52.937	53.992	55.811	
## 6	48.189	50.009	51.065	52.238	53.410	54.466	56.286	
## 7	48.661	50.481	51.538	52.711	53.885	54.941	56.762	
## 8	49.131	50.953	52.011	53.185	54.359	55.416	57.238	
## 1	48.785	50.607	51.665	52.840	54.016	55.074	56.896	
## 2	49.254	51.076	52.134	53.309	54.485	55.543	57.364	
## 3	49.724	51.545	52.603	53.778	54.953	56.011	57.833	
## 4	50.193	52.014	53.072	54.247	55.422	56.480	58.301	
## 5	50.662	52.483	53.541	54.716	55.891	56.949	58.770	
## 6	51.131	52.952	54.010	55.185	56.360	57.418	59.239	

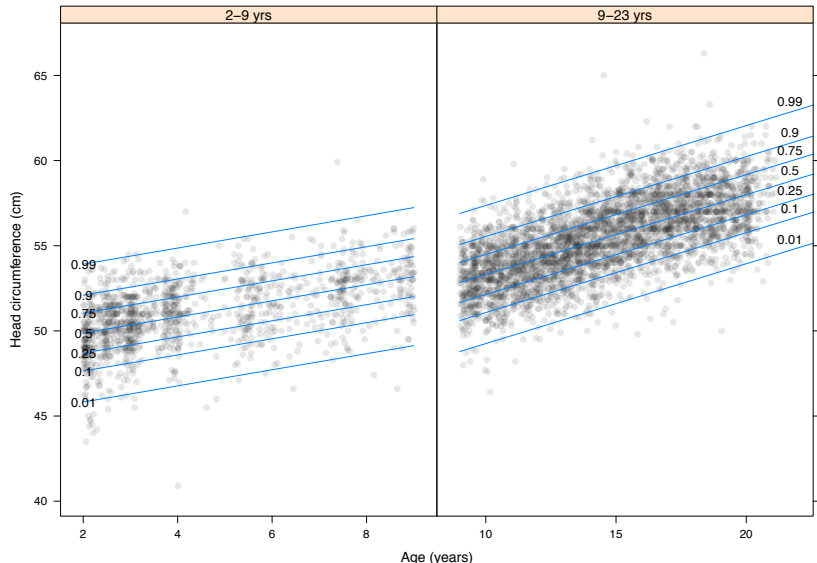
Dutch boys head circumference

Superimpose these conditional quantiles on our scatterplot

```
pfun <- function(x, y, ...) {  
  panel.xyplot(x = x, y = y, ...)  
  if (max(x) <= 9) {  
    apply(q2.23, 2, function(x)  
      panel.lines(gage[i], x[i]))  
  } else {  
    apply(q2.23, 2, function(x)  
      panel.lines(gage[-i], x[-i]))  
  }  
  panel.text(rep(max(db$age), length(tau)),  
    q2.23[nrow(q2.23),], label = tau, cex = 0.9)  
  panel.text(rep(min(db$age), length(tau)),  
    q2.23[1,], label = tau, cex = 0.9)  
}  
xyplot(head ~ age | cut, data = db, xlab = "Age (years)",  
  ylab = "Head circumference (cm)", pch = 19,  
  scales = list(x = list(relation = "free")),  
  layout = c(2, 1), col = rgb(.1, .1, .1, .1),  
  panel = pfun)
```

Dutch boys head circumference

- Parallel lines owing to the fact that the linear model assumes an error variance independent from age- variance homogeneity



Dutch boys head circumference - growth curves rq

Nonparametric version of our growth curves

```
rq2.9 <- rq(head ~ age, data = db, tau = tau,  
  subset = age < 9)  
rq2.9$coef
```

```
##           tau= 0.01  tau= 0.10  tau= 0.25  tau= 0.50  tau= 0.75  
## (Intercept) 43.2992424 46.9331190 48.0224215 48.9282511 50.1110357  
## age         0.6515152  0.4501608  0.4484305  0.4932735  0.4584041  
##           tau= 0.90  tau= 0.99  
## (Intercept) 50.765014 52.6367698  
## age         0.523416  0.4467354
```

```
rq9.23 <- rq(head ~ age, data = db, tau = tau,  
  subset = age > 9)  
rq9.23$coef
```

```
##           tau= 0.01  tau= 0.10  tau= 0.25  tau= 0.50  tau= 0.75  
## (Intercept) 44.3351899 46.4375451 47.5965517 48.5791795 49.6719626  
## age         0.4810127  0.4693141  0.4597701  0.4717949  0.4766355  
##           tau= 0.90  tau= 0.99  
## (Intercept) 50.7155801 52.6674762  
## age         0.4751381  0.4646251
```

Dutch boys head circumference- growth curves rq

Nonparametric version of our growth curves - prediction

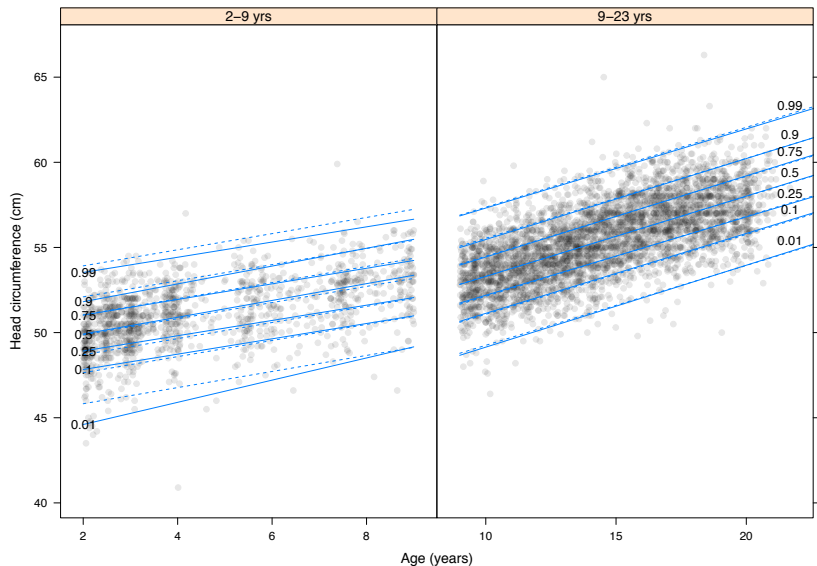
```
p2.23 <- rbind(predict(rq2.9,  
  newdata = data.frame(age = gage[i])),  
  predict(rq9.23,  
    newdata = data.frame(age = gage[-i])))  
head(p2.23)
```

##		tau= 0.01	tau= 0.10	tau= 0.25	tau= 0.50	tau= 0.75	tau= 0.90	tau= 0.99
## 1		44.60227	47.83344	48.91928	49.91480	51.02784	51.81185	53.53024
## 2		45.25379	48.28360	49.36771	50.40807	51.48625	52.33526	53.97698
## 3		45.90530	48.73376	49.81614	50.90135	51.94465	52.85868	54.42371
## 4		46.55682	49.18392	50.26457	51.39462	52.40306	53.38209	54.87045
## 5		47.20833	49.63408	50.71300	51.88789	52.86146	53.90551	55.31718
## 6		47.85985	50.08424	51.16143	52.38117	53.31986	54.42893	55.76392

Dutch boys head circumference - growth curves rq

```
pfun <- function(x, y, ...) {  
  panel.xyplot(x = x, y = y, ...)  
  if (max(x) <= 9) {  
    apply(q2.23, 2, function(x)  
      panel.lines(gage[i], x[i], lty = 2))  
    apply(p2.23, 2, function(x)  
      panel.lines(gage[i], x[i]))  
  } else {  
    apply(q2.23, 2, function(x)  
      panel.lines(gage[-i], x[-i], lty = 2))  
    apply(p2.23, 2, function(x)  
      panel.lines(gage[-i], x[-i]))  
  }  
  panel.text(rep(max(db$age), length(tau)),  
    p2.23[nrow(p2.23),], label = tau, cex = 0.9)  
  panel.text(rep(min(db$age), length(tau)),  
    p2.23[1,], label = tau, cex = 0.9)  
}  
xyplot(head ~ age | cut, data = db, xlab = "Age (years)",  
  ylab = "Head circumference (cm)", pch = 19,  
  scales = list(x = list(relation = "free")),  
  layout = c(2, 1), col = rgb(.1, .1, .1, .1),  
  panel = pfun)
```


Dutch boys head circumference - growth curves rq



Dutch boys head circumference - non-linear qr

Non-linear quantile regression (use *rqss* function)

```
rqssmod <- vector(mode = "list", length = length(tau))
db$lage <- with(db, age^(1/3))
for (i in 1:length(tau))
  rqssmod[[i]] <- rqss(head ~ qss(lage, lambda = 1),
    data = db, tau = tau[i])

gage <- seq(from = min(db$age), to = max(db$age), length = 50)
p <- sapply(1:length(tau), function(i) { predict(rqssmod[[i]],
  newdata = data.frame(lage = gage^(1/3)))
})
```

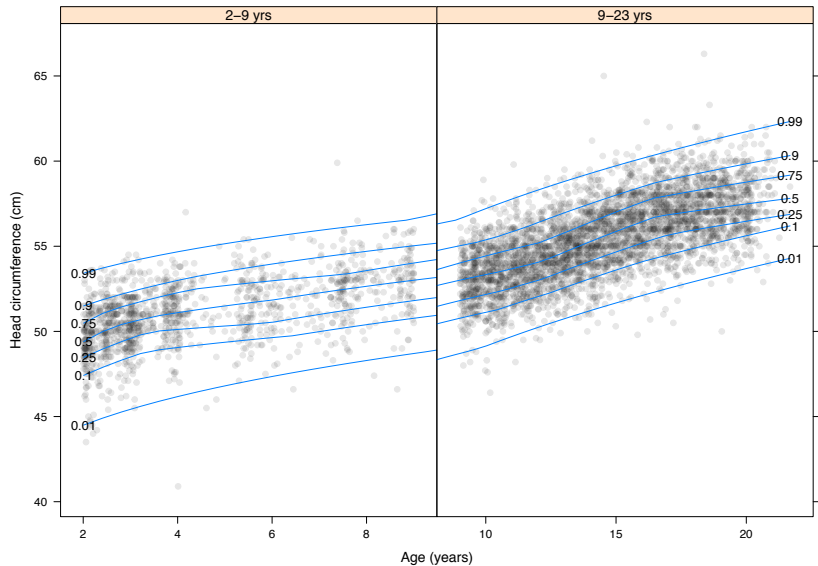
Dutch boys head circumference

Non-linear quantile regression

```
pfun <- function(x, y, ...) {  
  panel.xyplot(x = x, y = y, ...)  
  apply(p, 2, function(x) panel.lines(gage, x))  
  panel.text(rep(max(db$age), length(tau)),  
    p[nrow(p),], label = tau, cex = 0.9)  
  panel.text(rep(min(db$age), length(tau)),  
    p[1,], label = tau, cex = 0.9)  
}  
xyplot(head ~ age | cut, data = db, xlab = "Age (years)",  
  ylab = "Head circumference (cm)", pch = 19,  
  scales = list(x = list(relation = "free")),  
  layout = c(2, 1), col = rgb(.1, .1, .1, .1),  
  panel = pfun)
```

Dutch boys head circumference

Non-linear quantile regression

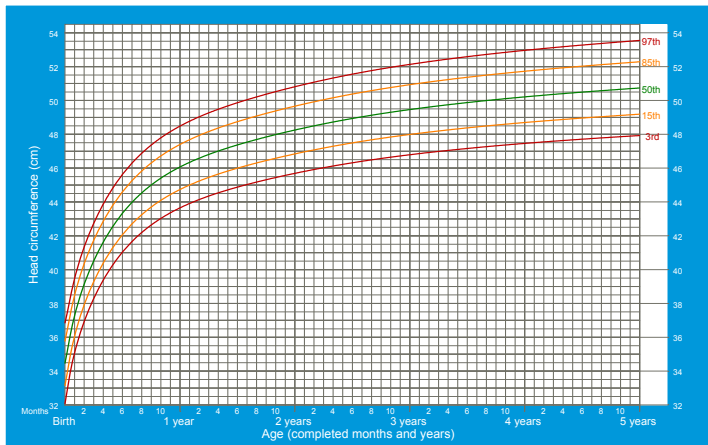


Playing with the whole db data

Let us now try to replicate the WHO plot

Head circumference-for-age BOYS

Birth to 5 years (percentiles)



WHO Child Growth Standards

Playing with the whole db data

```
## use the tau values as given in the above plot
library(gamlss.data)
data(db)
db2 <- db
tau <- c(.03, .15, .5, .85, .97)

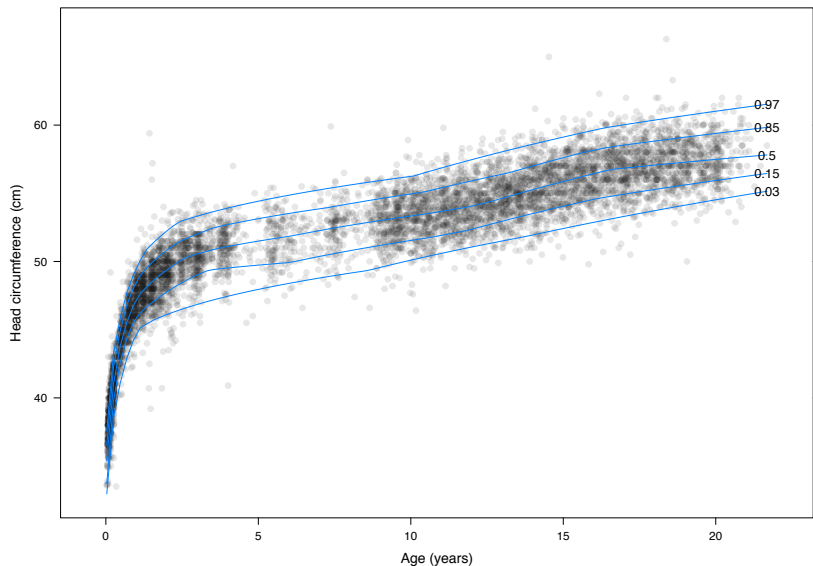
rqssmod <- vector(mode = "list", length = length(tau))
db2$lage <- with(db2, age^(1/3))
for (i in 1:length(tau))
  rqssmod[[i]] <- rqss(head ~ qss(lage, lambda = 1),
    data = db2, tau = tau[i])

gage <- seq(from = min(db2$age), to = max(db2$age), length = 100)
p <- sapply(1:length(tau), function(i) { predict(rqssmod[[i]],
  newdata = data.frame(lage = gage^(1/3)))
})
```

Playing with the whole db data

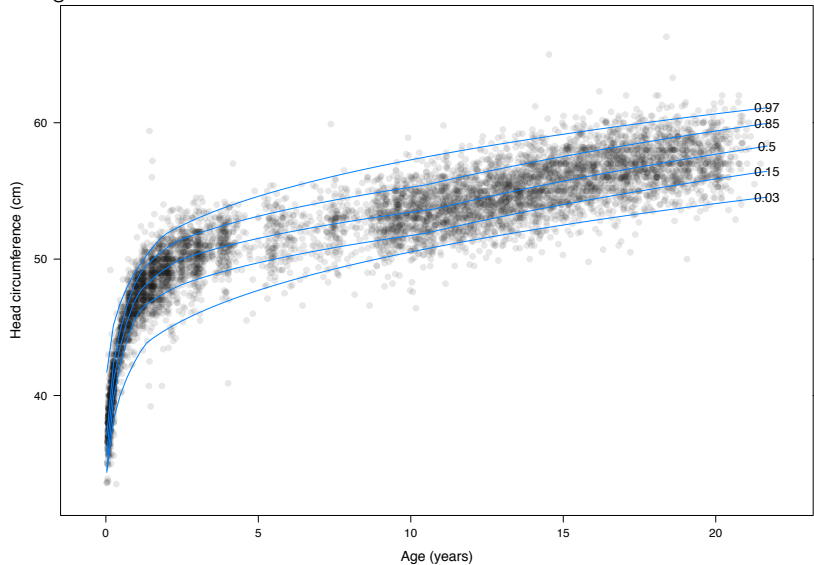
```
pfun <- function(x, y, ...) {  
  panel.xyplot(x = x, y = y, ...)  
  apply(p, 2, function(x) panel.lines(gage, x))  
  panel.text(rep(max(db2$age), length(tau)),  
    p[nrow(p),], label = tau, cex = 0.9)  
  #panel.text(rep(min(db2$age), length(tau)),  
    #p[1,], label = tau, cex = 0.9)  
}  
xyplot(head ~ age, data = db2, xlab = "Age (years)",  
  ylab = "Head circumference (cm)", pch = 19,  
  scales = list(x = list(relation = "free")),  
  layout = c(1, 1), col = rgb(.1, .1, .1, .1),  
  panel = pfun)
```

Playing with the whole db data



Playing with the whole db data

Change lambda = 20 for smoothness



Quantile regression final remark

- ▶ When estimating regression models, we have to be aware of the implications of model assumptions when interpreting the results. Symmetry, linearity, and variance homogeneity are among the strongest but common assumptions.
- ▶ Quantile regression, both in its linear and additive formulation, is an intellectually stimulating and practically very useful framework where such assumptions can be relaxed.
- ▶ At a more basic level, one should always ask Am I really interested in the mean? before using the regression models discussed in other chapters of this book.