

Lab 4

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TODO: As this is a policy exercise, you should do your best to address the campaign's questions from a causal perspective. At the same time, you should clearly explain the limitations of your analysis, and provide discussion around whether your estimates suffer from endogeneity bias.

Introduction

TODO: A brief introduction

Exploratory Data Analysis

TODO: An initial exploratory analysis. Detect any anomalies, including missing values, top-coded or bottom-coded variables, etc.

```
library(car) # lm
library(ggplot2) # ggplot
library(lmtest) # bptest
```

```
## Loading required package: zoo
```

```
##
```

```
## Attaching package: 'zoo'
```

```
## The following objects are masked from 'package:base':
```

```
##
```

```
##      as.Date, as.Date.numeric
```

```
library(plm) # vcovHC
```

```
## Loading required package: Formula
```

```
library(sandwich)
```

```
library(stargazer) # stargazer
```

```
##
```

```
## Please cite as:
```

```
## Hlavac, Marek (2015). stargazer: Well-Formatted Regression and Summary Statistics Tables.
```

```
## R package version 5.2. http://CRAN.R-project.org/package=stargazer
```

```
library(tidyr) # gather
```

```
data <- read.csv("crime.csv")
```

```
str(data)
```

```
## 'data.frame':   90 obs. of  26 variables:
```

```
## $ X      : int  1 2 3 4 5 6 7 8 9 10 ...
```

```
## $ county : int  1 3 5 7 9 11 13 15 17 19 ...
```

```
## $ year   : int  87 87 87 87 87 87 87 87 87 87 ...
```

```
## $ crmrte : num 0.0356 0.0153 0.013 0.0268 0.0106 ...
## $ prbarr : num 0.298 0.132 0.444 0.365 0.518 ...
## $ prbconv : num 0.528 1.481 0.268 0.525 0.477 ...
## $ prbpris : num 0.436 0.45 0.6 0.435 0.443 ...
## $ avgscn : num 6.71 6.35 6.76 7.14 8.22 ...
## $ polpc : num 0.001828 0.000746 0.001234 0.00153 0.00086 ...
## $ density : num 2.423 1.046 0.413 0.492 0.547 ...
## $ taxpc : num 31 26.9 34.8 42.9 28.1 ...
## $ west : int 0 0 1 0 1 1 0 0 0 0 ...
## $ central : int 1 1 0 1 0 0 0 0 0 0 ...
## $ urban : int 0 0 0 0 0 0 0 0 0 0 ...
## $ pctmin80 : num 20.22 7.92 3.16 47.92 1.8 ...
## $ wcon : num 281 255 227 375 292 ...
## $ wtuc : num 409 376 372 398 377 ...
## $ wtrd : num 221 196 229 191 207 ...
## $ wfir : num 453 259 306 281 289 ...
## $ wser : num 274 192 210 257 215 ...
## $ wmfgr : num 335 300 238 282 291 ...
## $ wfed : num 478 410 359 412 377 ...
## $ wsta : num 292 363 332 328 367 ...
## $ wloc : num 312 301 281 299 343 ...
## $ mix : num 0.0802 0.0302 0.4651 0.2736 0.0601 ...
## $ pctymle : num 0.0779 0.0826 0.0721 0.0735 0.0707 ...
```

```
summary(data)
```

```
##      X      county      year      crmrte
## Min.   :1.00   Min.   : 1.0   Min.   :87   Min.   :0.005533
## 1st Qu.:23.25  1st Qu.: 51.5   1st Qu.:87   1st Qu.:0.020604
## Median :45.50  Median :103.0   Median :87   Median :0.030002
## Mean   :45.50  Mean   :100.6   Mean   :87   Mean   :0.033510
## 3rd Qu.:67.75  3rd Qu.:150.5   3rd Qu.:87   3rd Qu.:0.040249
## Max.   :90.00  Max.   :197.0   Max.   :87   Max.   :0.098966
##      prbarr      prbconv      prbpris      avgscn
## Min.   :0.09277   Min.   :0.06838   Min.   :0.1500   Min.   : 5.380
## 1st Qu.:0.20495   1st Qu.:0.34422   1st Qu.:0.3642   1st Qu.: 7.375
## Median :0.27146   Median :0.45170   Median :0.4222   Median : 9.110
## Mean   :0.29524   Mean   :0.55086   Mean   :0.4106   Mean   : 9.689
## 3rd Qu.:0.34487   3rd Qu.:0.58513   3rd Qu.:0.4576   3rd Qu.:11.465
## Max.   :1.09091   Max.   :2.12121   Max.   :0.6000   Max.   :20.700
##      polpc      density      taxpc      west
## Min.   :0.0007459   Min.   :0.2034   Min.   : 25.69   Min.   :0.0000
## 1st Qu.:0.0012378   1st Qu.:0.5472   1st Qu.: 30.73   1st Qu.:0.0000
## Median :0.0014897   Median :0.9792   Median : 34.92   Median :0.0000
## Mean   :0.0017080   Mean   :1.4379   Mean   : 38.16   Mean   :0.2333
## 3rd Qu.:0.0018856   3rd Qu.:1.5693   3rd Qu.: 41.01   3rd Qu.:0.0000
## Max.   :0.0090543   Max.   :8.8277   Max.   :119.76   Max.   :1.0000
##      central      urban      pctmin80      wcon
## Min.   :0.0000   Min.   :0.000000   Min.   : 1.284   Min.   :193.6
## 1st Qu.:0.0000   1st Qu.:0.000000   1st Qu.:10.024   1st Qu.:250.8
## Median :0.0000   Median :0.000000   Median :24.852   Median :281.2
## Mean   :0.3778   Mean   :0.08889   Mean   :25.713   Mean   :285.4
## 3rd Qu.:1.0000   3rd Qu.:0.000000   3rd Qu.:38.183   3rd Qu.:315.0
## Max.   :1.0000   Max.   :1.000000   Max.   :64.348   Max.   :436.8
##      wtuc      wtrd      wfir      wser
```

```
## Min. :187.6 Min. :154.2 Min. :170.9 Min. : 133.0
## 1st Qu.:374.3 1st Qu.:190.7 1st Qu.:285.6 1st Qu.: 229.3
## Median :404.8 Median :203.0 Median :317.1 Median : 253.1
## Mean :410.9 Mean :210.9 Mean :321.6 Mean : 275.3
## 3rd Qu.:440.7 3rd Qu.:224.3 3rd Qu.:342.6 3rd Qu.: 277.6
## Max. :613.2 Max. :354.7 Max. :509.5 Max. :2177.1
##      wmfg      wfed      wsta      wloc
## Min. :157.4 Min. :326.1 Min. :258.3 Min. :239.2
## 1st Qu.:288.6 1st Qu.:398.8 1st Qu.:329.3 1st Qu.:297.2
## Median :321.1 Median :448.9 Median :358.4 Median :307.6
## Mean :336.0 Mean :442.6 Mean :357.7 Mean :312.3
## 3rd Qu.:359.9 3rd Qu.:478.3 3rd Qu.:383.2 3rd Qu.:328.8
## Max. :646.9 Max. :598.0 Max. :499.6 Max. :388.1
##      mix      pctymle
## Min. :0.01961 Min. :0.06216
## 1st Qu.:0.08060 1st Qu.:0.07437
## Median :0.10095 Median :0.07770
## Mean :0.12905 Mean :0.08403
## 3rd Qu.:0.15206 3rd Qu.:0.08352
## Max. :0.46512 Max. :0.24871
```

```
# sample size n = 90
nrow(data)
```

```
## [1] 90
```

```
# number of variables = 26
ncol(data)
```

```
## [1] 26
```

```
# number of prbarr where probability is greater than 1
sum(data$prbarr > 1)
```

```
## [1] 1
```

```
# number of prbconv where probability is greater than 1
sum(data$prbconv > 1)
```

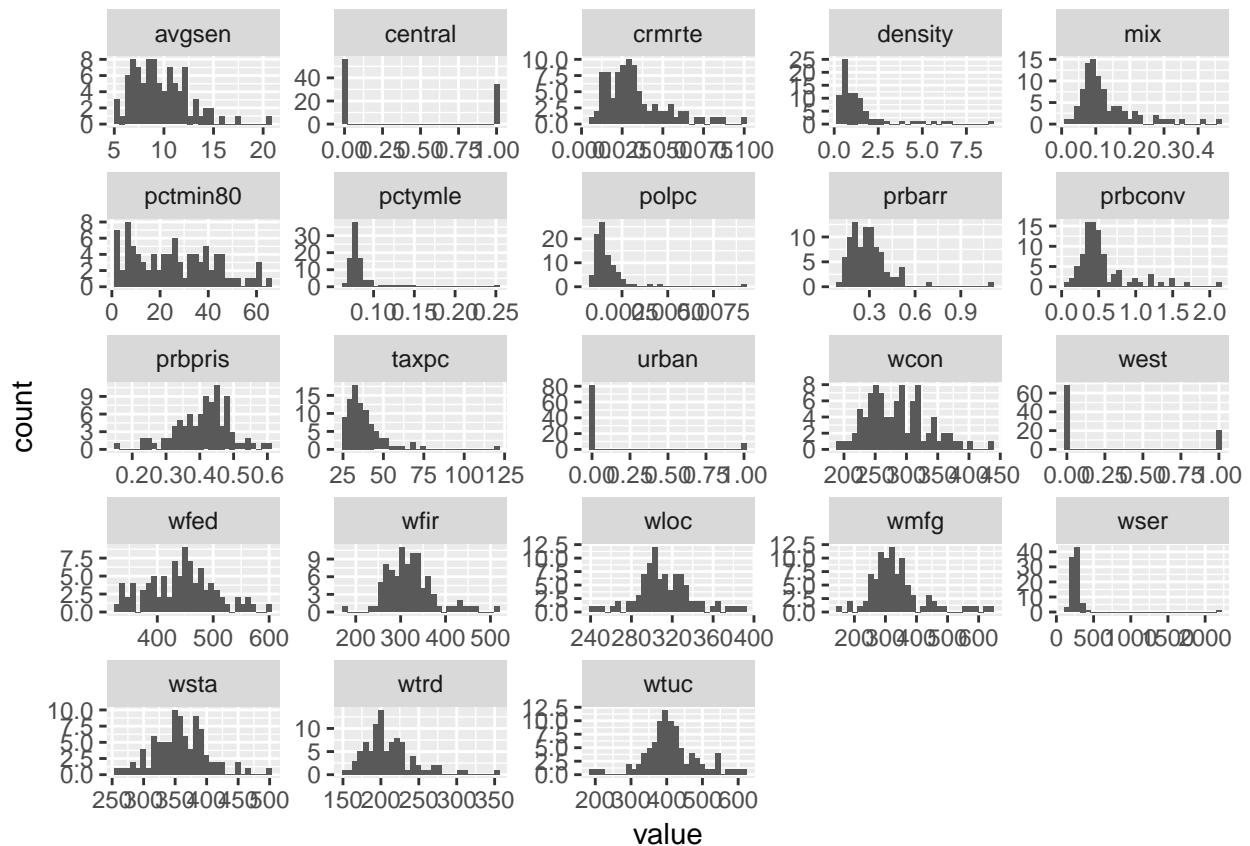
```
## [1] 10
```

```
# verify number of missing values = 0
colSums(sapply(data, is.na))
```

```
##      X      county      year      crmrte      prbarr      prbconv      prbpris      avgse
##      0          0          0          0          0          0          0          0
##      polpc      density      taxpc      west      central      urban      pctmin80      wcon
##      0          0          0          0          0          0          0          0
##      wtuc      wtrd      wfir      wser      wmfg      wfed      wsta      wloc
##      0          0          0          0          0          0          0          0
##      mix      pctymle
##      0          0
```

```
# plot every variable except x, county, year
plot.data <- data[!(names(data) %in% c("X", "county", "year"))]
ggplot(gather(plot.data), aes(value)) +
  facet_wrap(~key, scales="free") +
  geom_histogram()
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```



```
# just list outliers for variables that are non-categorical
subset.data <- data[!(names(data) %in% c("X", "county", "year", "central", "urban", "west"))]

# list all outliers
list_all_outliers <- function (var) {
  outliers <- sort(boxplot.stats(var)$out)
  return(paste(length(outliers), ": ", paste(outliers, collapse=" ")))
}
lapply(subset.data, list_all_outliers)
```

```
## $crmrte
## [1] "5 : 0.0729478970170021, 0.0790162980556488, 0.0834982022643089, 0.0883848965167999, 0.09896589"
##
## $prbarr
## [1] "2 : 0.689023971557617, 1.09090995788574"
##
## $prbconv
## [1] "11 : 0.972972989082336, 1.01538002490997, 1.06896996498108, 1.18292999267578, 1.22561001777649"
##
## $prbpris
## [1] "1 : 0.150000005960464"
##
## $avgsen
## [1] "1 : 20.7000007629395"
##
```

```
## $polpc
## [1] "4 : 0.00316379009746015, 0.00400961982086301, 0.00445923022925854, 0.00905433017760515"
##
## $density
## [1] "8 : 3.93455100059509, 4.38875865936279, 4.8347339630127, 5.12442398071289, 5.6744966506958, 6.12442398071289"
##
## $taxpc
## [1] "6 : 56.8621063232422, 61.1525115966797, 67.6796340942383, 67.8479766845703, 75.6724319458008, 75.6724319458008"
##
## $pctmin80
## [1] "0 : "
##
## $wcon
## [1] "1 : 436.766632080078"
##
## $wtuc
## [1] "9 : 187.617263793945, 202.429153442383, 213.675216674805, 548.323852539062, 548.986511230469, 548.986511230469"
##
## $wtrd
## [1] "5 : 277.292510986328, 279.227264404297, 306.083526611328, 308.576232910156, 354.676116943359, 354.676116943359"
##
## $wfir
## [1] "7 : 170.940170288086, 430.069702148438, 435.110717773438, 441.141296386719, 453.172210693359, 453.172210693359"
##
## $wser
## [1] "4 : 133.043060302734, 354.300720214844, 391.308074951172, 2177.06811523438"
##
## $wmfg
## [1] "7 : 157.410003662109, 494.299987792969, 560.780029296875, 567.059997558594, 588.989990234375, 588.989990234375"
##
## $wfed
## [1] "0 : "
##
## $wsta
## [1] "1 : 499.589996337891"
##
## $wloc
## [1] "5 : 239.169998168945, 246.649993896484, 379.769989013672, 386.119995117188, 388.089996337891, 388.089996337891"
##
## $mix
## [1] "8 : 0.273622035980225, 0.28078818321228, 0.290488421916962, 0.311355322599411, 0.3195266425609, 0.3195266425609"
##
## $pctymle
## [1] "10 : 0.0989191979169846, 0.0993558466434479, 0.106941692531109, 0.114216551184654, 0.122244790, 0.122244790"
```

Glaring outliers: 1. Probability > 1 for prbarr and prbconv 2. Outlier 2177 in wser which happens to be in central instead of urban (weird)

```
# number of urban
sum(data$urban == 1)
```

```
## [1] 8
```

```
# number of central
sum(data$central == 1)
```

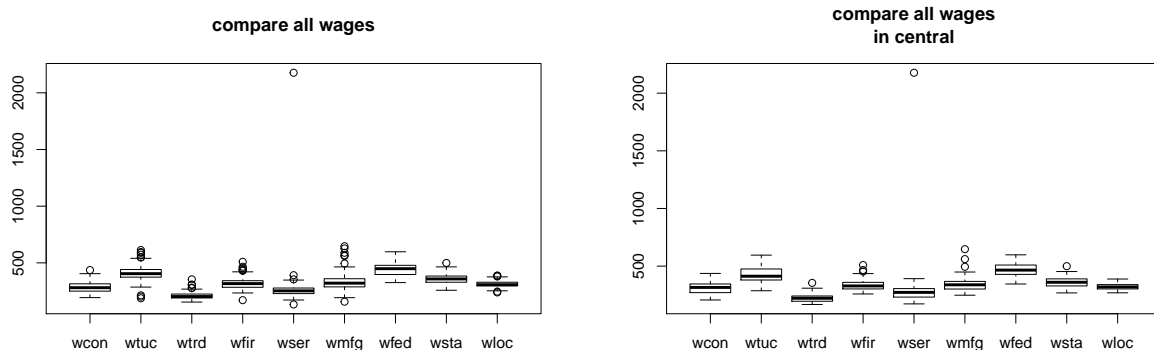
```
## [1] 34
# number of west
sum(data$west == 1)

## [1] 21
# determine if a county can have multiple categorical label: urban, central, west
urban.county <- data[data$urban == 1, ]$county
central.county <- data[data$central == 1, ]$county
west.county <- data[data$west == 1, ]$county
# can a county be labeled as both urban and central?
uc.county <- c(urban.county, central.county)
uc.county[duplicated(uc.county)]

## [1] 63 67 81 119 183
# can a county be labeled as both urban and west?
uw.county <- c(urban.county, west.county)
uw.county[duplicated(uw.county)]

## [1] 21
# I have verified that there is no county labeled as both central and west
# counties that have more than one categorical label
all.county <- c(urban.county, central.county, west.county)
all.county[duplicated(all.county)]

## [1] 63 67 81 119 183 21
# compare all the wages
boxplot(as.vector(data[c(16:24)]), main="compare all wages")
# compare all wages in central
central.data <- data[data$central == 1, ]
boxplot(as.vector(central.data[c(16:24)]), main="compare all wages\nin central")
```

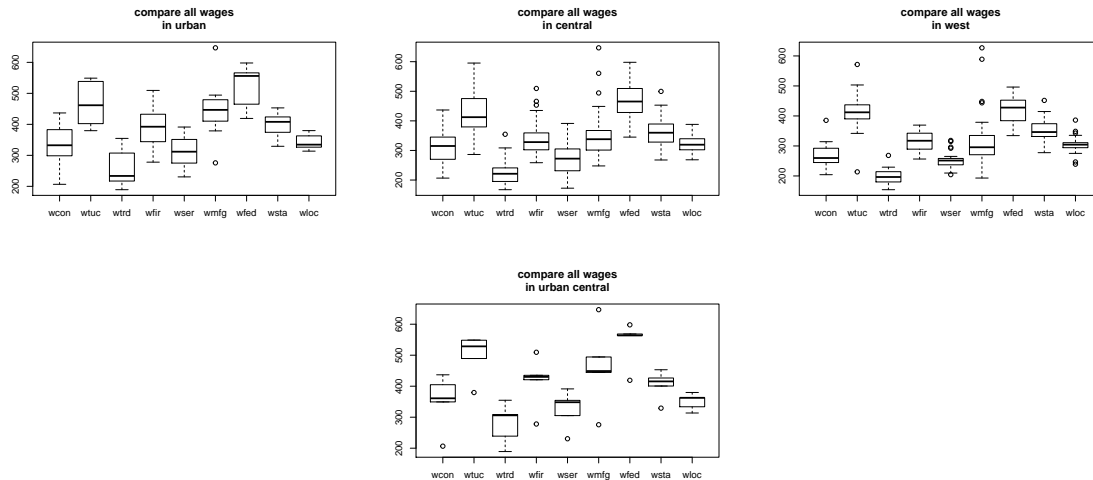


```
# compare all wages in urban
urban.data <- data[data$urban == 1, ]
boxplot(as.vector(urban.data[c(16:24)]), main="compare all wages\nin urban")
# compare all wages in central with outlier removed
central.data[central.data$wser == 2177.06811523438, ]$wser <- mean(central.data$wser)
boxplot(as.vector(central.data[c(16:24)]), main="compare all wages\nin central")
# compare all wages in west
west.data <- data[data$west == 1, ]
```

```

boxplot(as.vector(west.data[c(16:24)]), main="compare all wages\nin west")
# compare all wages in urban and central
urban.central.data <- data[data$urban == 1 & data$central == 1, ]
boxplot(as.vector(urban.central.data[c(16:24)]), main="compare all wages\nin urban central")
# compare all wages in urban and west
# not worth it because it's only one dataset

```



```

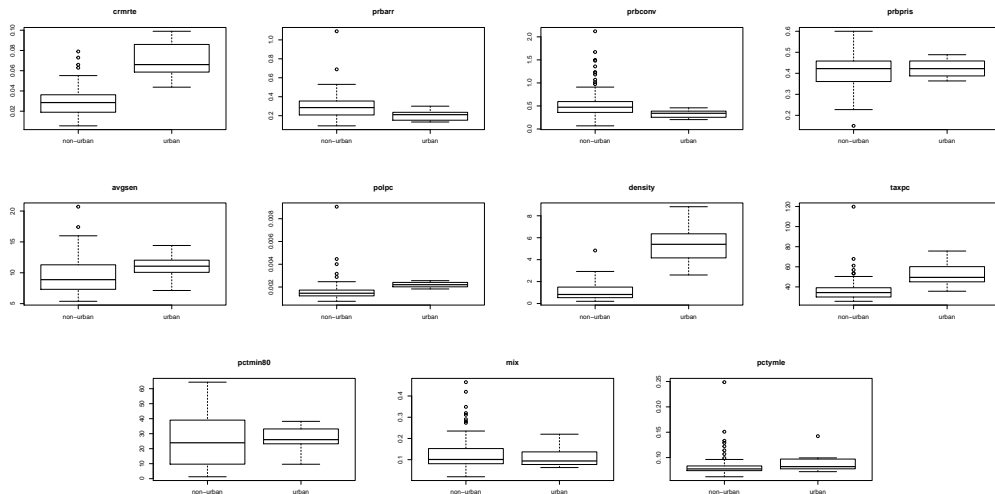
# in urban area, is there higher ...?
independents <- c("crmte", "prbarr", "prbconv", "prbpris", "avgsen", "polpc", "density", "taxpc", "pctmle")
mapply(function(var, var.name) {
  boxplot(var ~ urban, data=data, names=c("non-urban", "urban"), main=var.name)
}, subset(data, select=independents), independents)

```

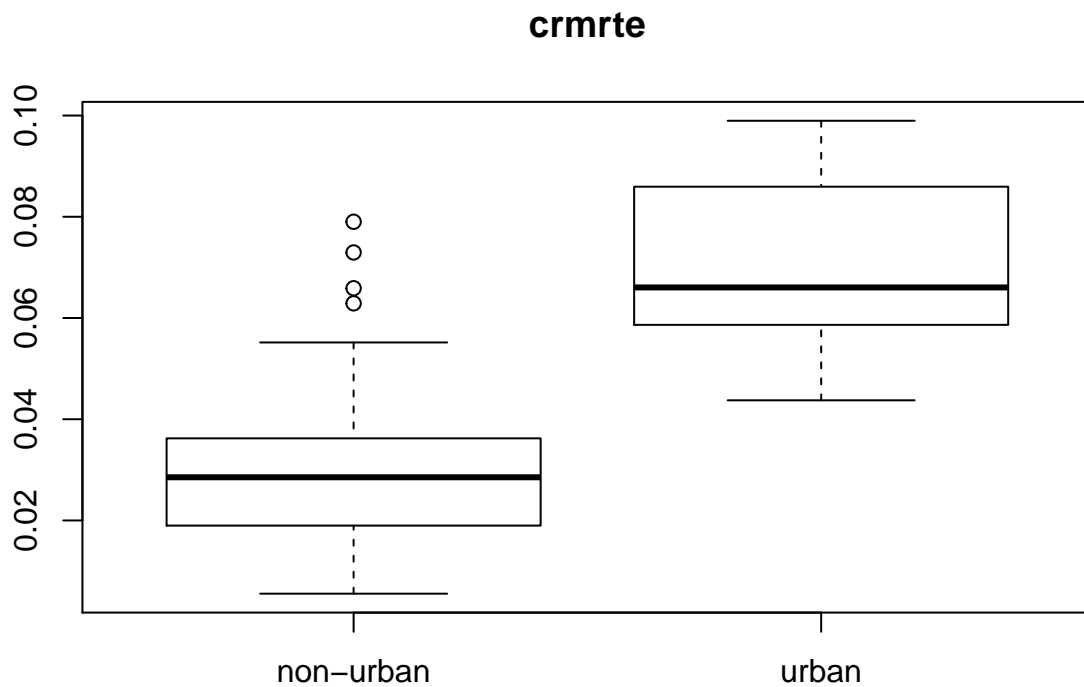
```

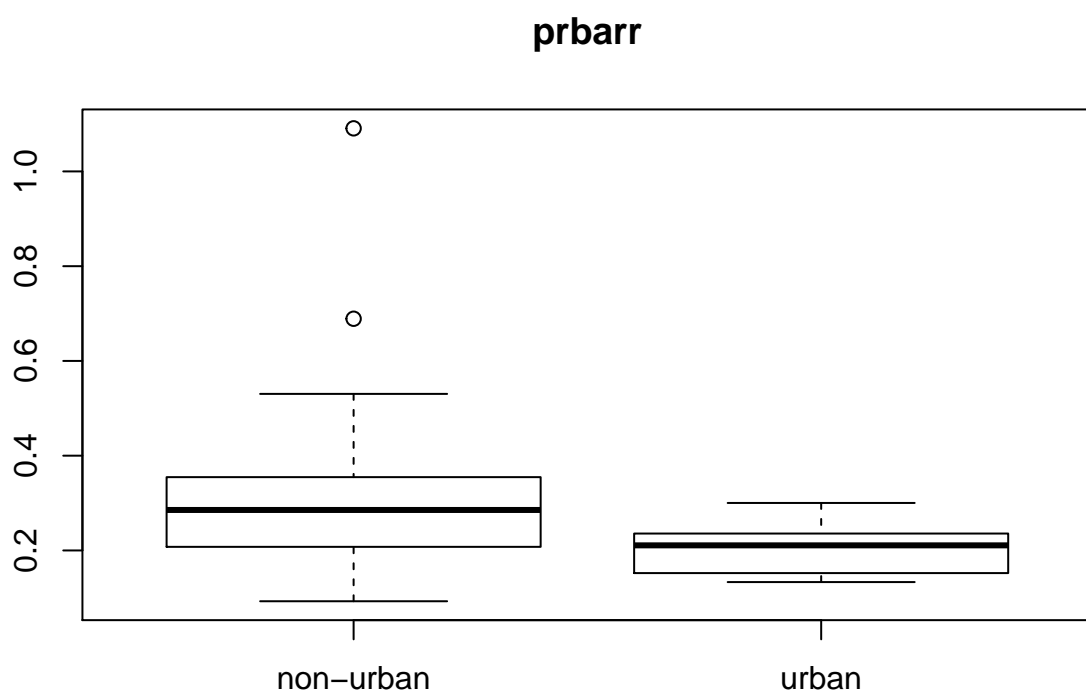
##      crmte      prbarr      prbconv      prbpris      avgsen
## stats Numeric,10 Numeric,10 Numeric,10 Numeric,10 Numeric,10
## n      Numeric,2  Numeric,2  Numeric,2  Numeric,2  Numeric,2
## conf   Numeric,4  Numeric,4  Numeric,4  Numeric,4  Numeric,4
## out    Numeric,4  Numeric,2  Numeric,11 0.15      Numeric,2
## group  Numeric,4  Numeric,2  Numeric,11 1          Numeric,2
## names  Character,2 Character,2 Character,2 Character,2 Character,2
##      polpc      density      taxpc      pctmin80      mix
## stats Numeric,10 Numeric,10 Numeric,10 Numeric,10 Numeric,10
## n      Numeric,2  Numeric,2  Numeric,2  Numeric,2  Numeric,2
## conf   Numeric,4  Numeric,4  Numeric,4  Numeric,4  Numeric,4
## out    Numeric,5  4.834734  Numeric,6  Numeric,0  Numeric,8
## group  Numeric,5  1          Numeric,6  Numeric,0  Numeric,8
## names  Character,2 Character,2 Character,2 Character,2 Character,2
##      pctmle
## stats Numeric,10
## n      Numeric,2
## conf   Numeric,4
## out    Numeric,9
## group  Numeric,9
## names  Character,2

```

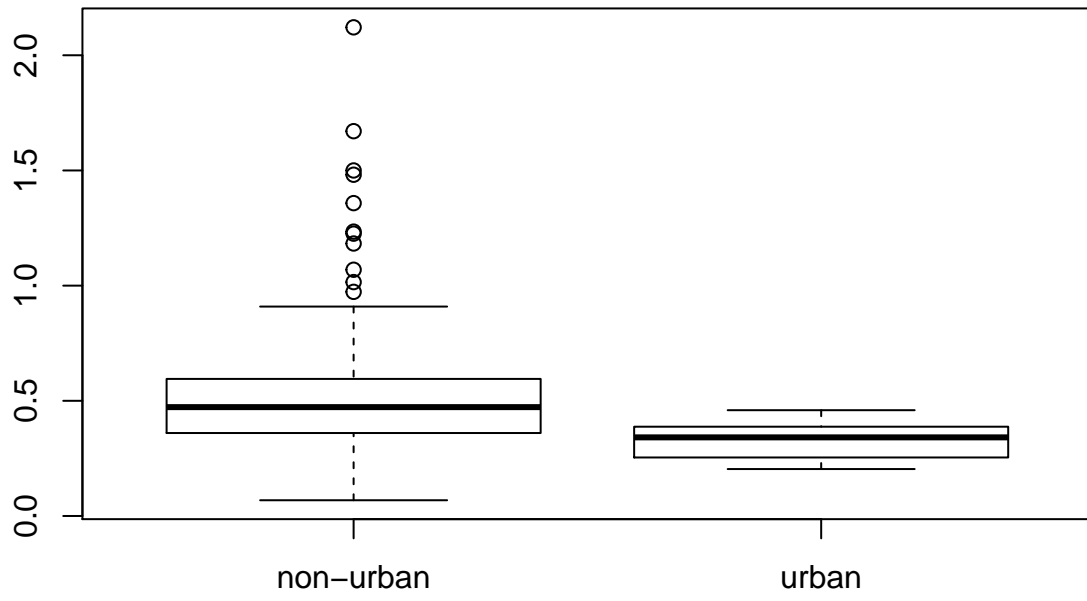


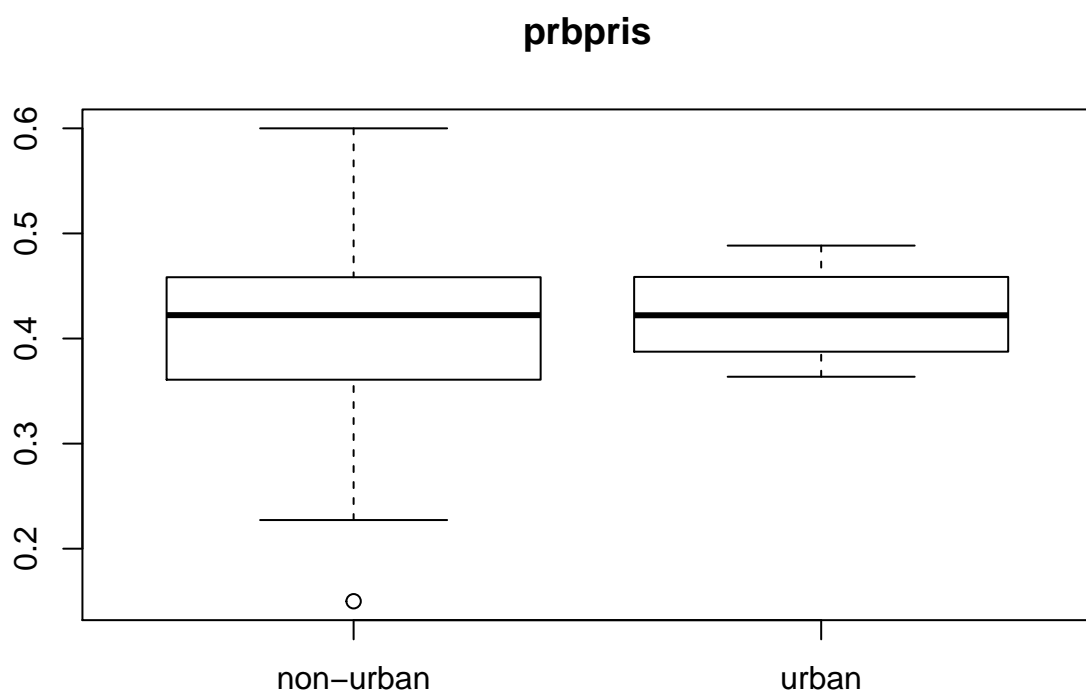
```
# repeat of above but checking out all the wages
mapapply(function(var, var.name) {
  boxplot(var ~ urban, data=data, names=c("non-urban", "urban"), main=var.name)
}, subset.data, names(subset.data))
```

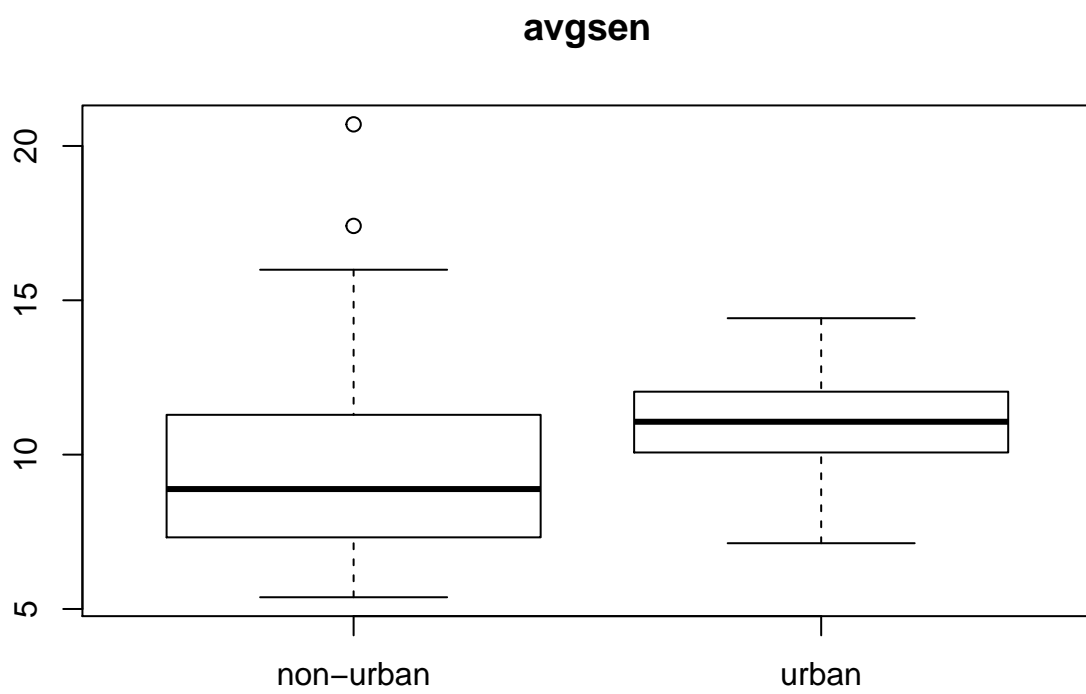


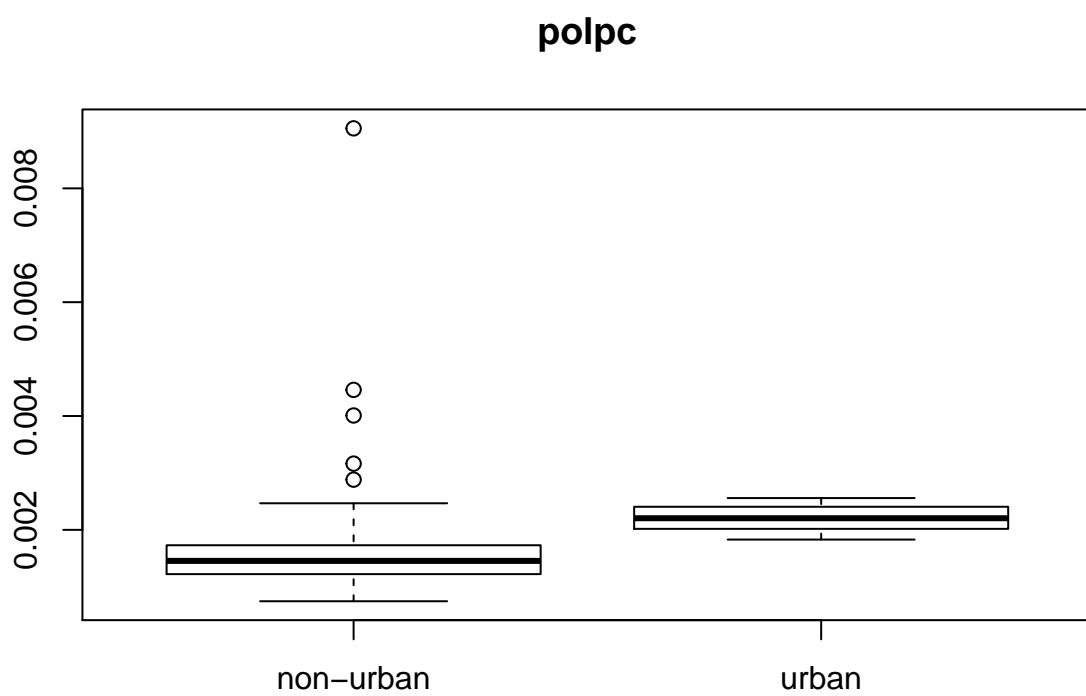


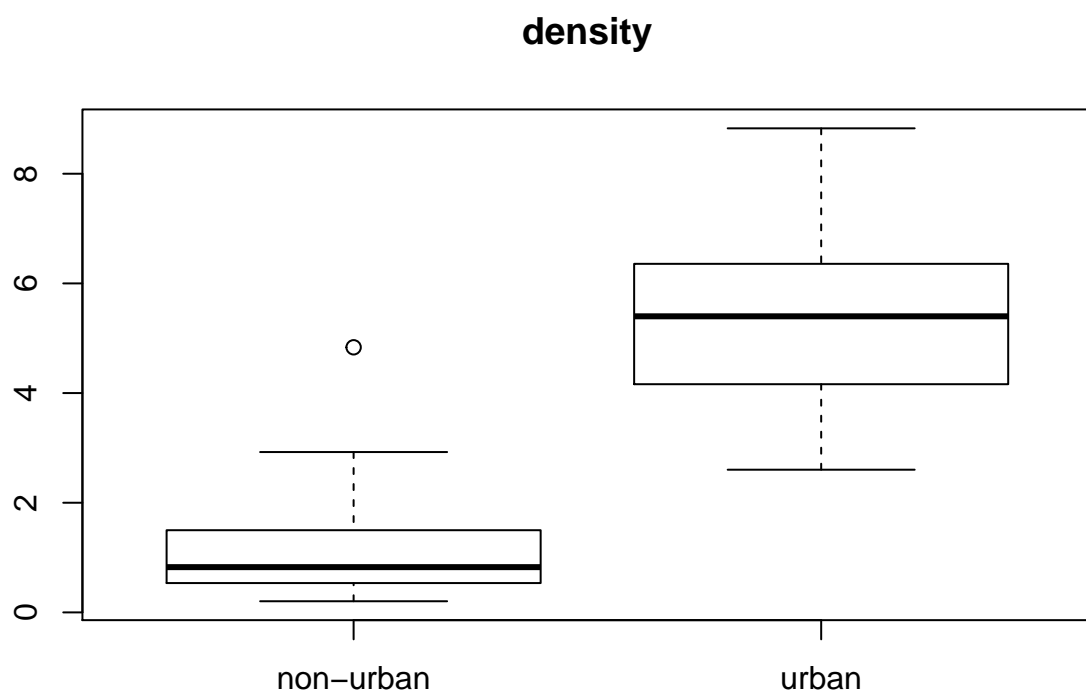
prbconv

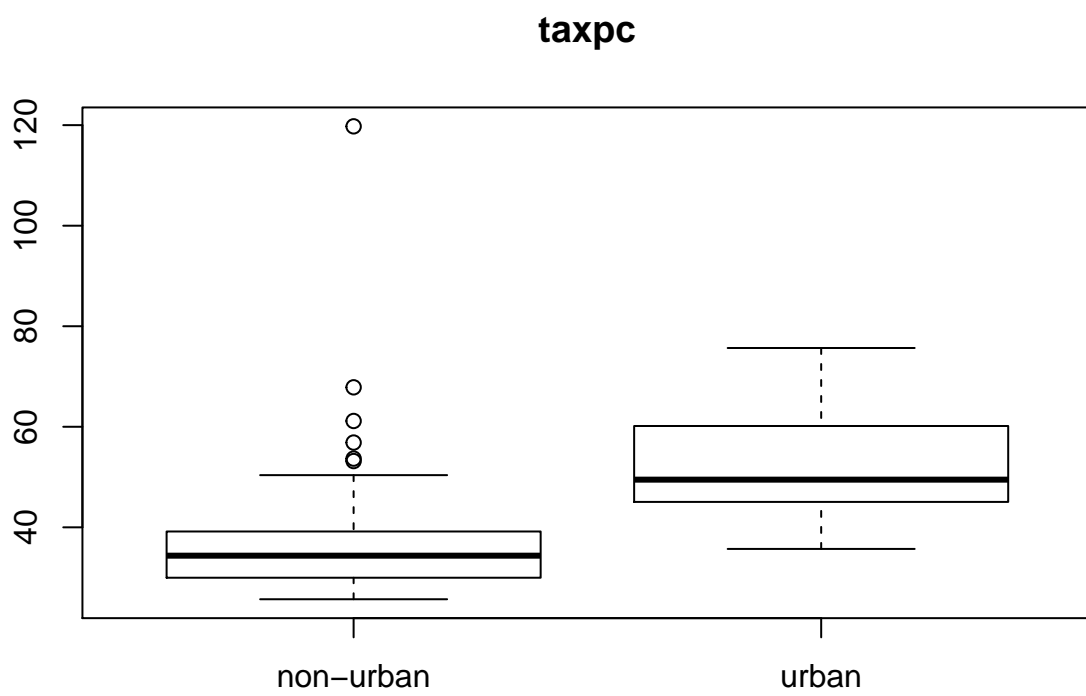




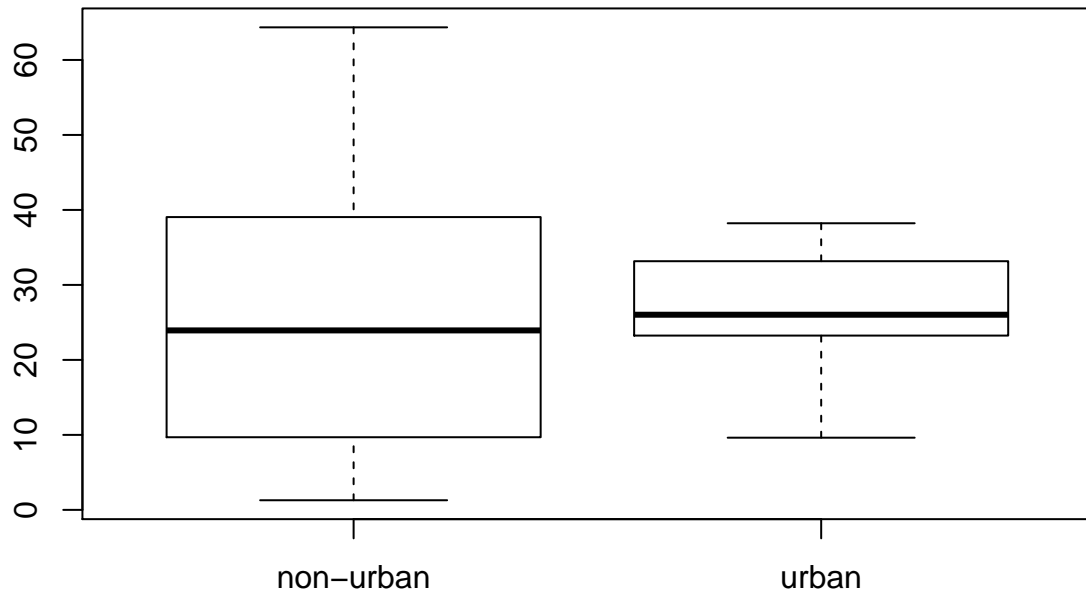


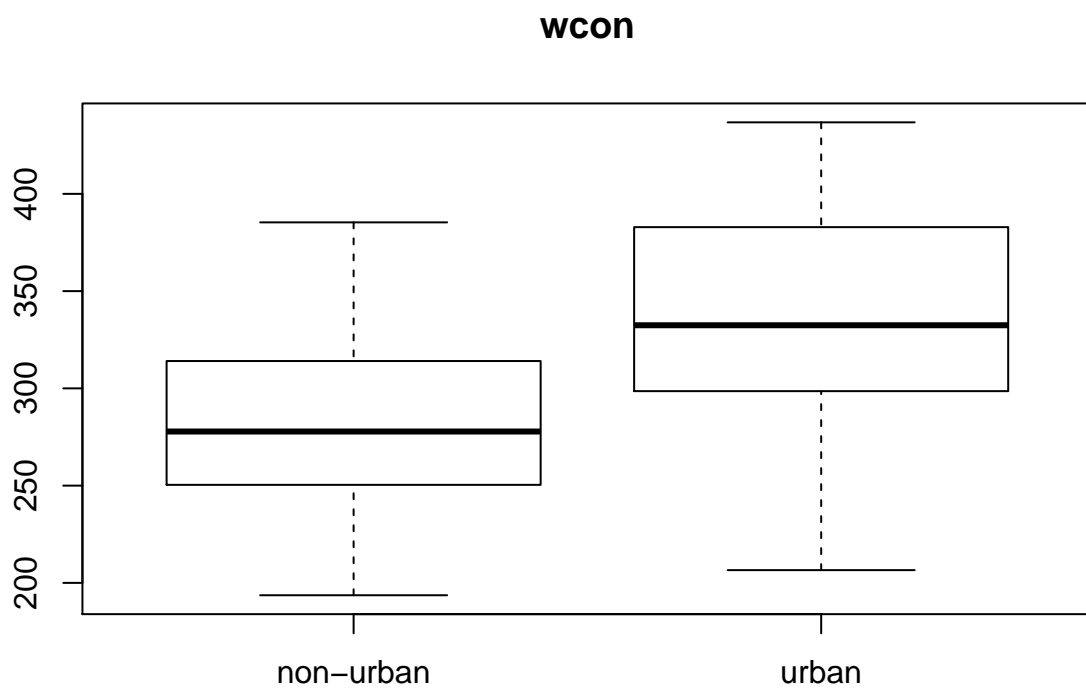


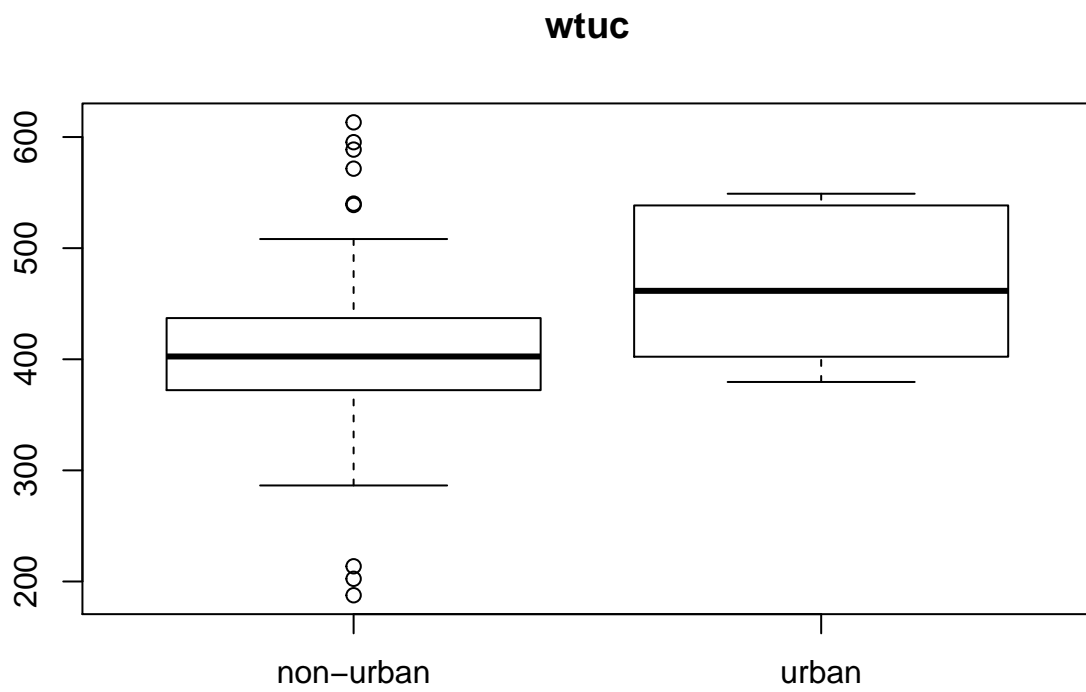


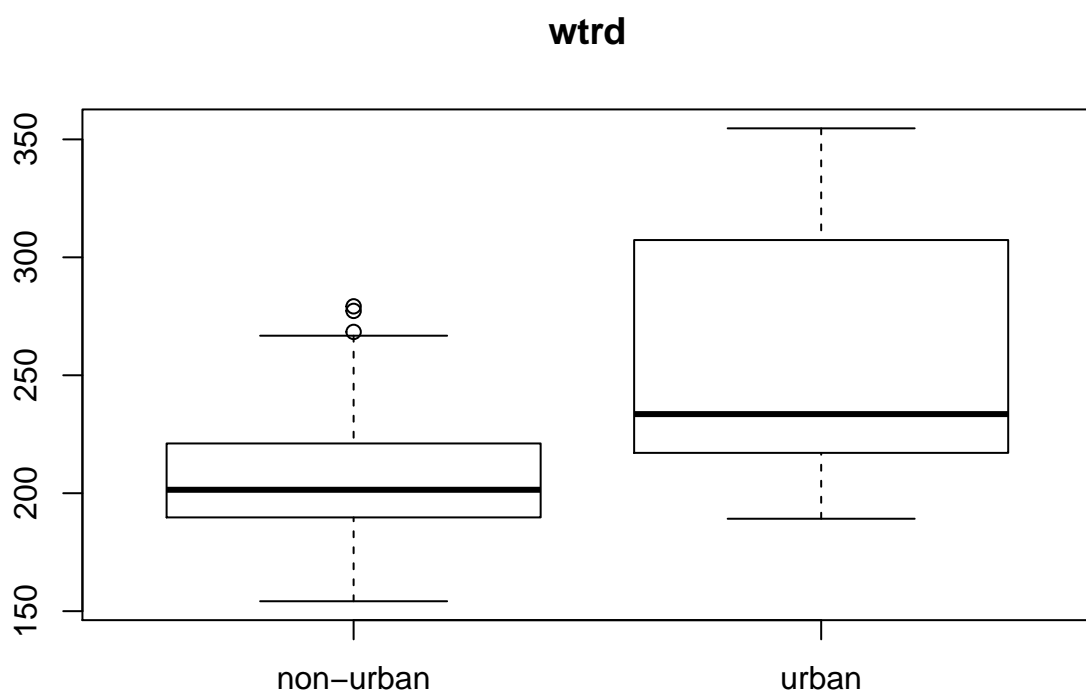


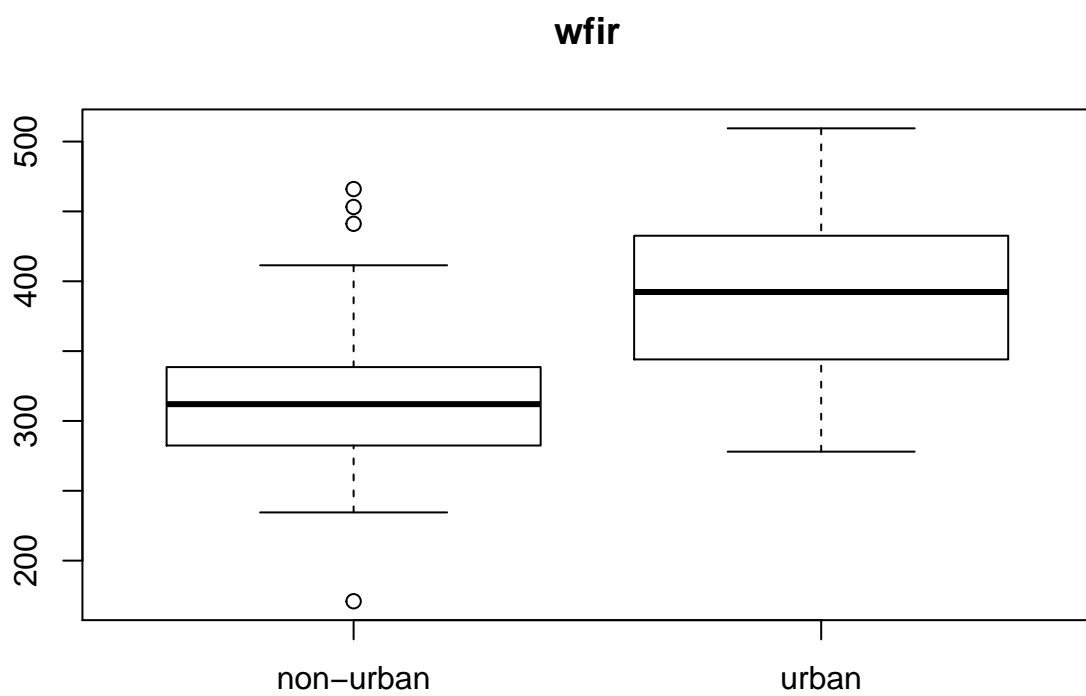
pctmin80

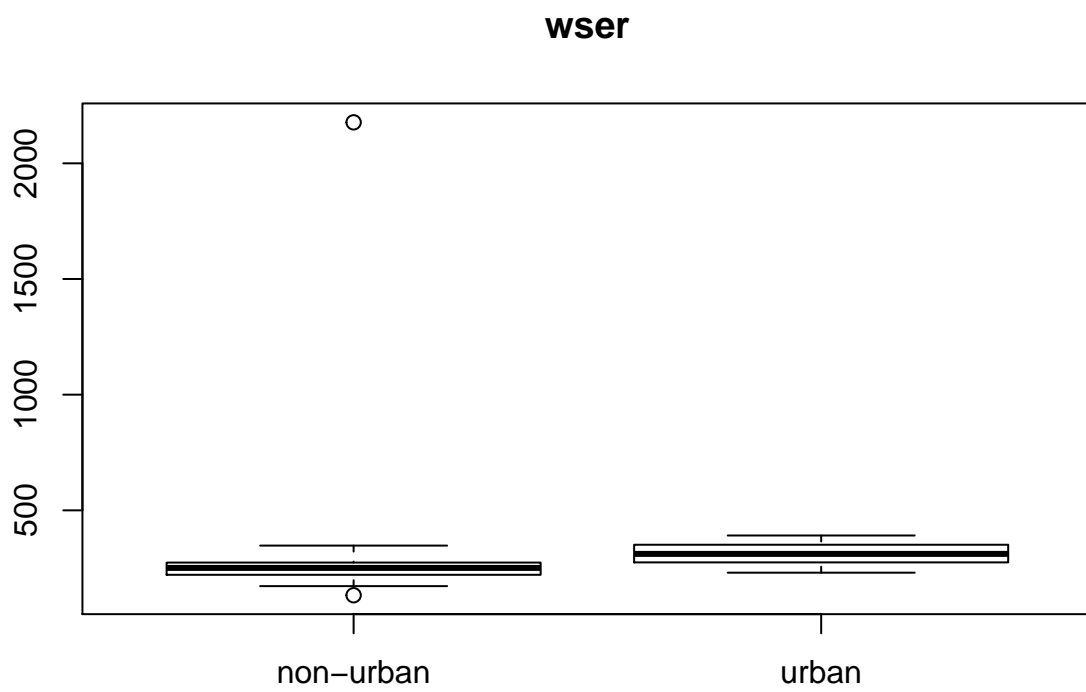


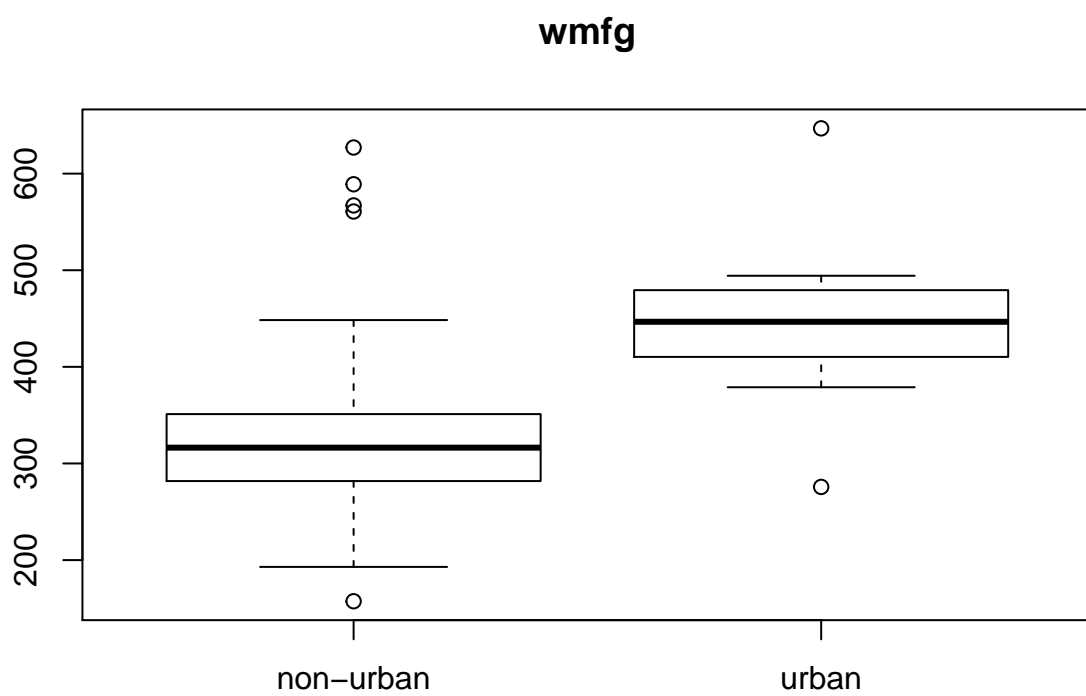


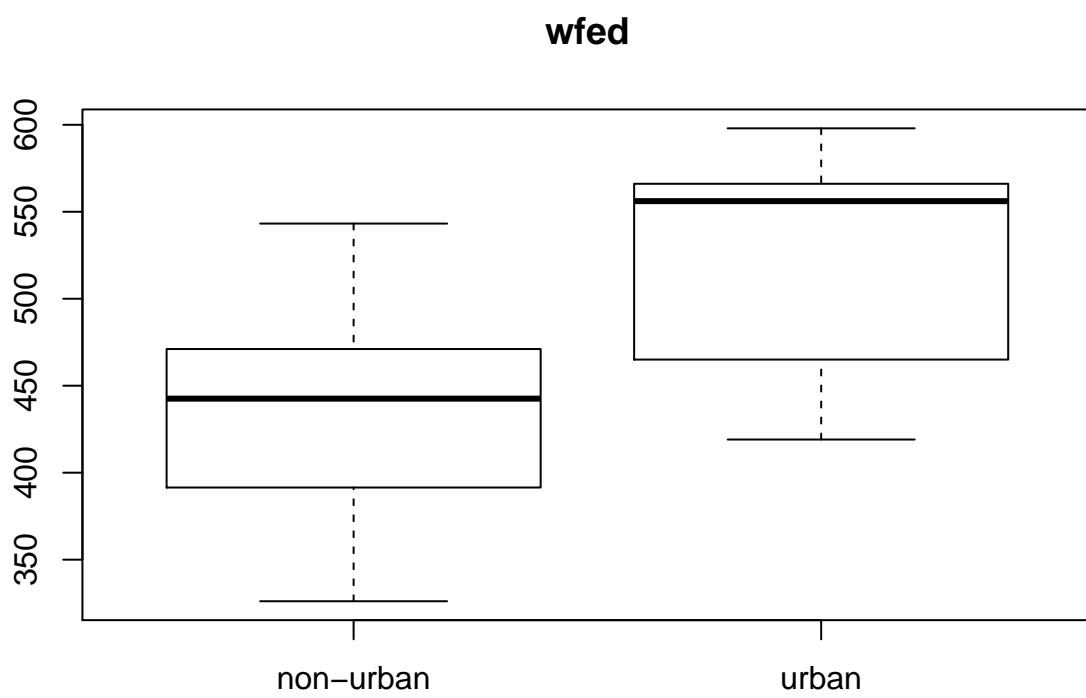


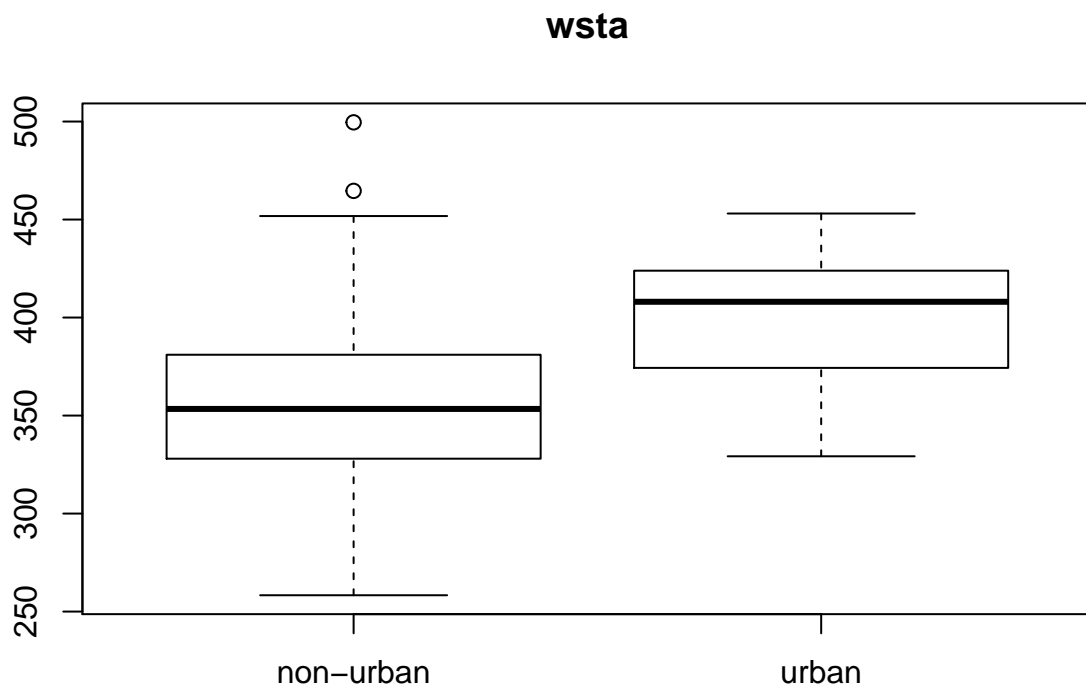


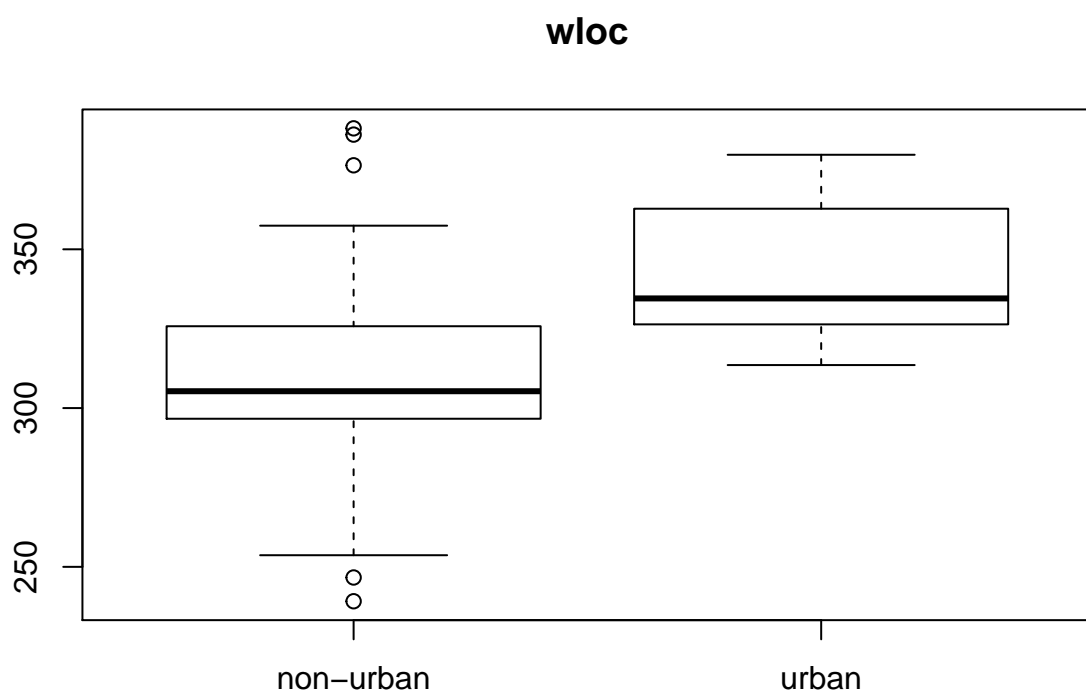


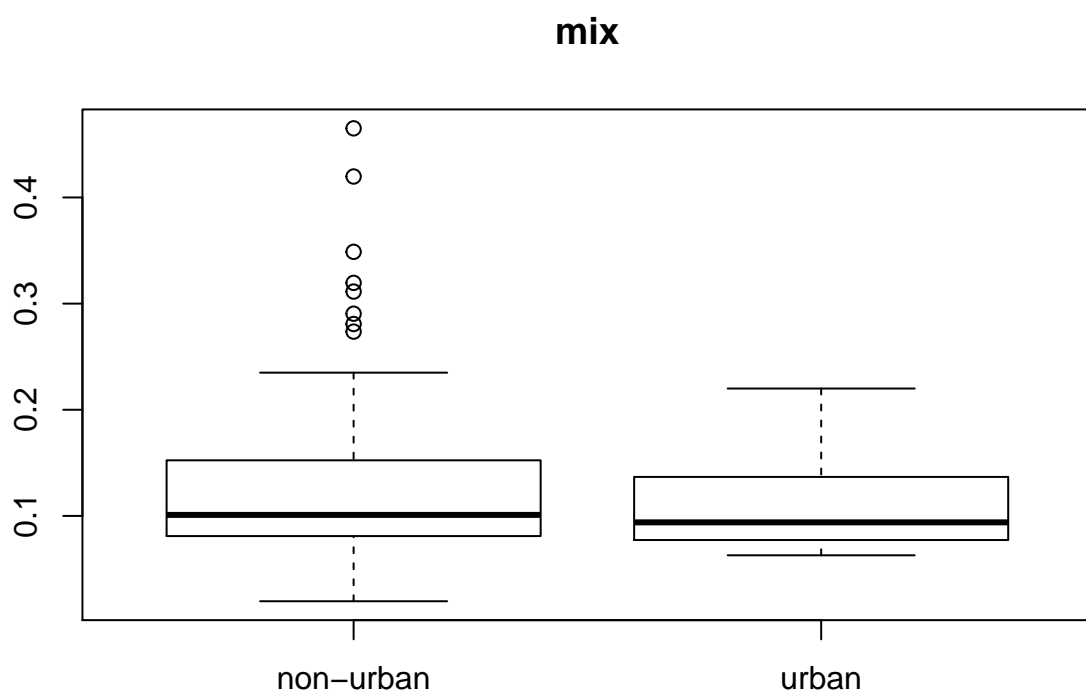




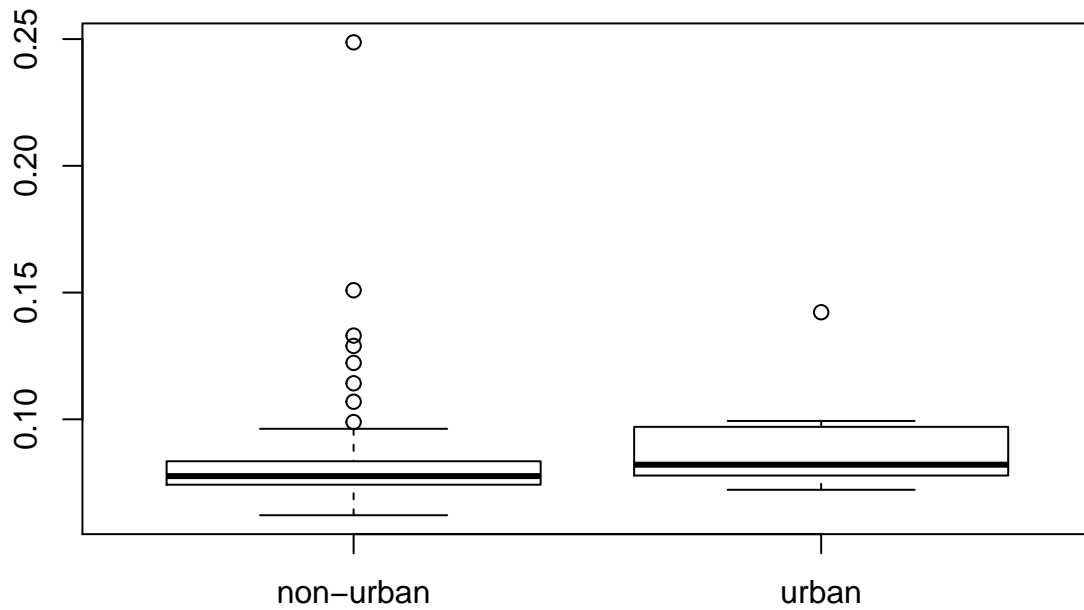








pctymle



```
##      crmrte      prbarr      prbconv      prbpris      avgsen
## stats Numeric,10 Numeric,10 Numeric,10 Numeric,10 Numeric,10
## n      Numeric,2  Numeric,2  Numeric,2  Numeric,2  Numeric,2
## conf   Numeric,4  Numeric,4  Numeric,4  Numeric,4  Numeric,4
## out    Numeric,4  Numeric,2  Numeric,11  0.15      Numeric,2
## group  Numeric,4  Numeric,2  Numeric,11  1          Numeric,2
## names  Character,2 Character,2 Character,2 Character,2 Character,2
##      polpc      density      taxpc      pctmin80      wcon
## stats Numeric,10 Numeric,10 Numeric,10 Numeric,10 Numeric,10
## n      Numeric,2  Numeric,2  Numeric,2  Numeric,2  Numeric,2
## conf   Numeric,4  Numeric,4  Numeric,4  Numeric,4  Numeric,4
## out    Numeric,5  4.834734  Numeric,6  Numeric,0  Numeric,0
## group  Numeric,5  1          Numeric,6  Numeric,0  Numeric,0
## names  Character,2 Character,2 Character,2 Character,2 Character,2
##      wtuc      wtrd      wfir      wser      wmfg
## stats Numeric,10 Numeric,10 Numeric,10 Numeric,10 Numeric,10
## n      Numeric,2  Numeric,2  Numeric,2  Numeric,2  Numeric,2
## conf   Numeric,4  Numeric,4  Numeric,4  Numeric,4  Numeric,4
## out    Numeric,9  Numeric,3  Numeric,4  Numeric,2  Numeric,7
## group  Numeric,9  Numeric,3  Numeric,4  Numeric,2  Numeric,7
## names  Character,2 Character,2 Character,2 Character,2 Character,2
##      wfed      wsta      wloc      mix      pctymle
## stats Numeric,10 Numeric,10 Numeric,10 Numeric,10 Numeric,10
## n      Numeric,2  Numeric,2  Numeric,2  Numeric,2  Numeric,2
## conf   Numeric,4  Numeric,4  Numeric,4  Numeric,4  Numeric,4
## out    Numeric,0  Numeric,2  Numeric,5  Numeric,8  Numeric,9
```

```
## group Numeric,0 Numeric,2 Numeric,5 Numeric,8 Numeric,9
## names Character,2 Character,2 Character,2 Character,2 Character,2
```

In urban areas, there is higher

- crime rates
- average sentence
- police per capita
- density
- tax revenue per capita
- percentage of minority (slightly higher but not much)
- percentage of young male (slightly higher but not much)
- wages tend to be higher overall

In urban areas, there is lower

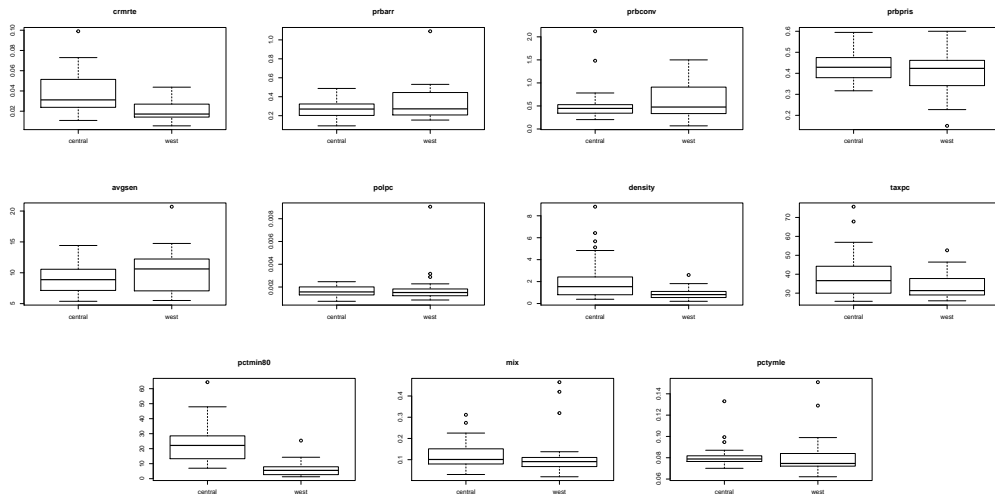
- probability of arrest
- probability of conviction
- face to face confrontation (slightly lower but not much)

In urban areas, there is about the same

- probability of prison sentence

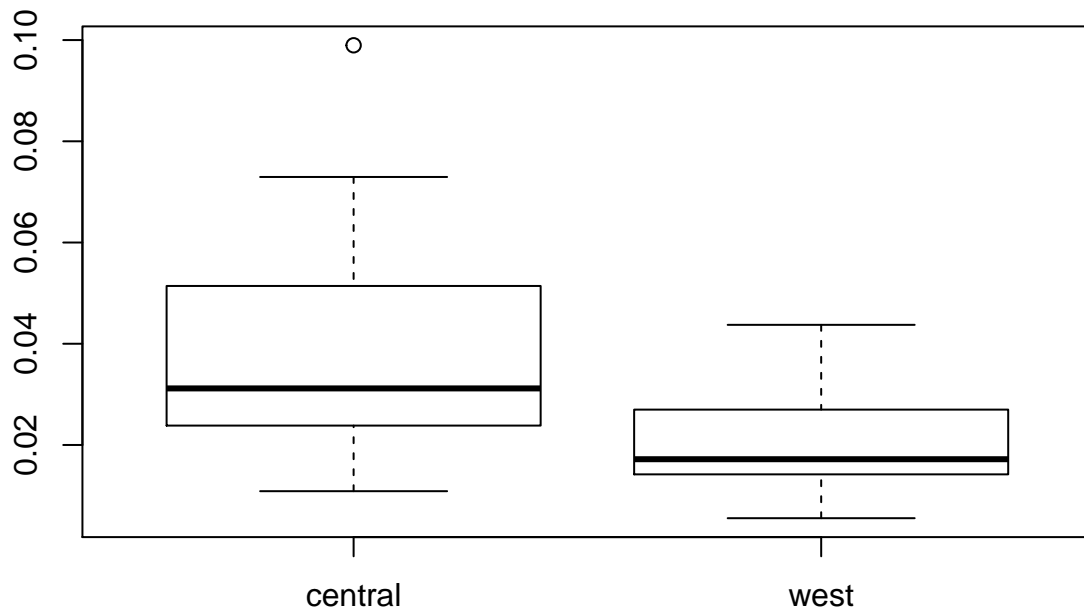
```
# comparing central and west, is there higher ...?
central.data <- data[data$central == 1, ]
subset.central.data <- subset(central.data, select=independents)
west.data <- data[data$west == 1, ]
subset.west.data <- subset(west.data, select=independents)
mapply(function(central.var, west.var, var.name) {
  boxplot(central.var, west.var, names=c("central", "west"), main=var.name)
}, subset.central.data, subset.west.data, independents)
```

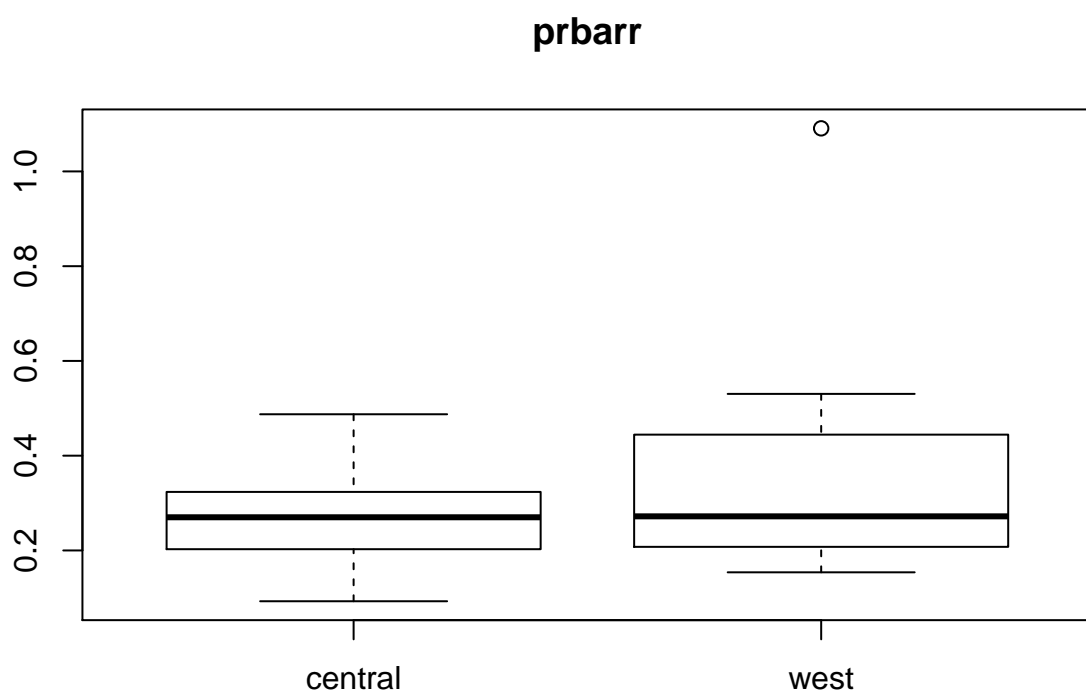
```
##      crmrte      prbarr      prbconv      prbpris      avgsen
## stats Numeric,10 Numeric,10 Numeric,10 Numeric,10 Numeric,10
## n      Numeric,2  Numeric,2  Numeric,2  Numeric,2  Numeric,2
## conf   Numeric,4  Numeric,4  Numeric,4  Numeric,4  Numeric,4
## out    0.0989659  1.09091   Numeric,2  0.15       20.7
## group  1         2         Numeric,2  2         2
## names  Character,2 Character,2 Character,2 Character,2 Character,2
##      polpc      density      taxpc      pctmin80      mix
## stats Numeric,10 Numeric,10 Numeric,10 Numeric,10 Numeric,10
## n      Numeric,2  Numeric,2  Numeric,2  Numeric,2  Numeric,2
## conf   Numeric,4  Numeric,4  Numeric,4  Numeric,4  Numeric,4
## out    Numeric,3  Numeric,5  Numeric,3  Numeric,2  Numeric,5
## group  Numeric,3  Numeric,5  Numeric,3  Numeric,2  Numeric,5
## names  Character,2 Character,2 Character,2 Character,2 Character,2
##      pctymle
## stats Numeric,10
## n      Numeric,2
## conf   Numeric,4
## out    Numeric,5
## group  Numeric,5
## names  Character,2
```



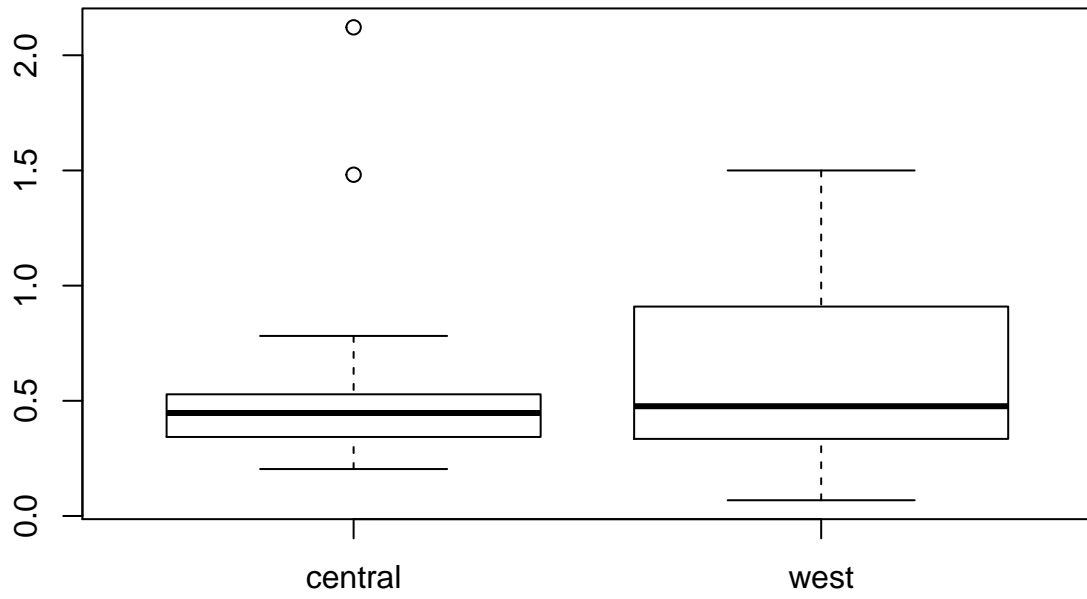
```
# repeat of above but checking out all the wages
central.data <- data[data$central == 1, ]
subset.central.data <- subset(central.data, select=names(subset.data))
west.data <- data[data$west == 1, ]
subset.west.data <- subset(west.data, select=names(subset.data))
mapapply(function(central.var, west.var, var.name) {
  boxplot(central.var, west.var, names=c("central", "west"), main=var.name)
}, subset.central.data, subset.west.data, names(subset.data))
```

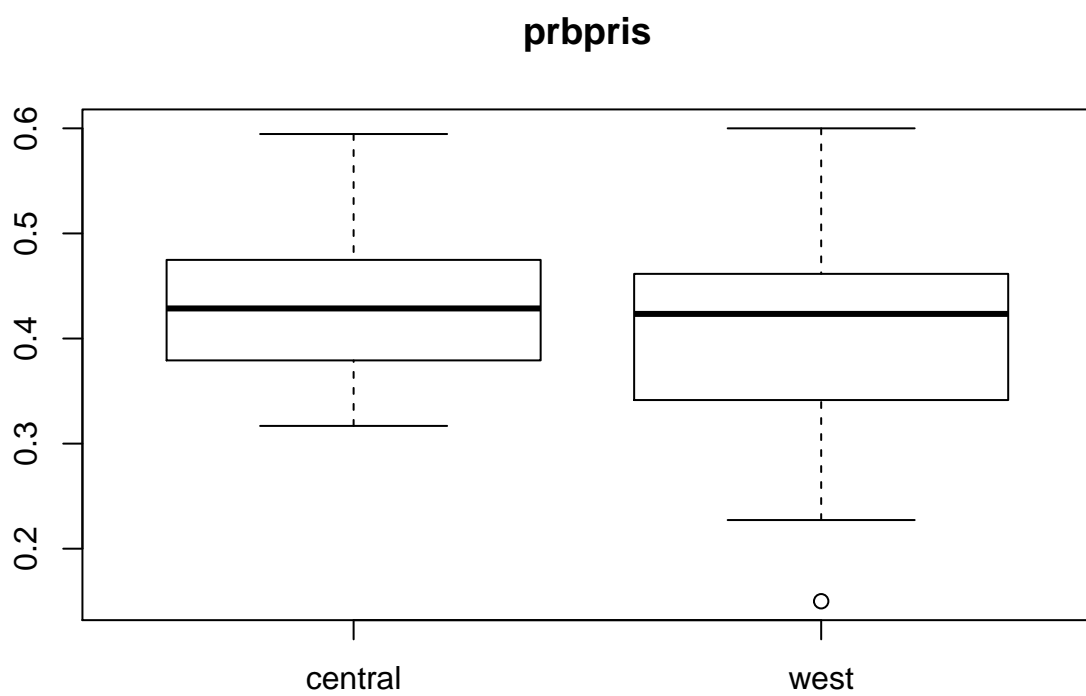
crmrte

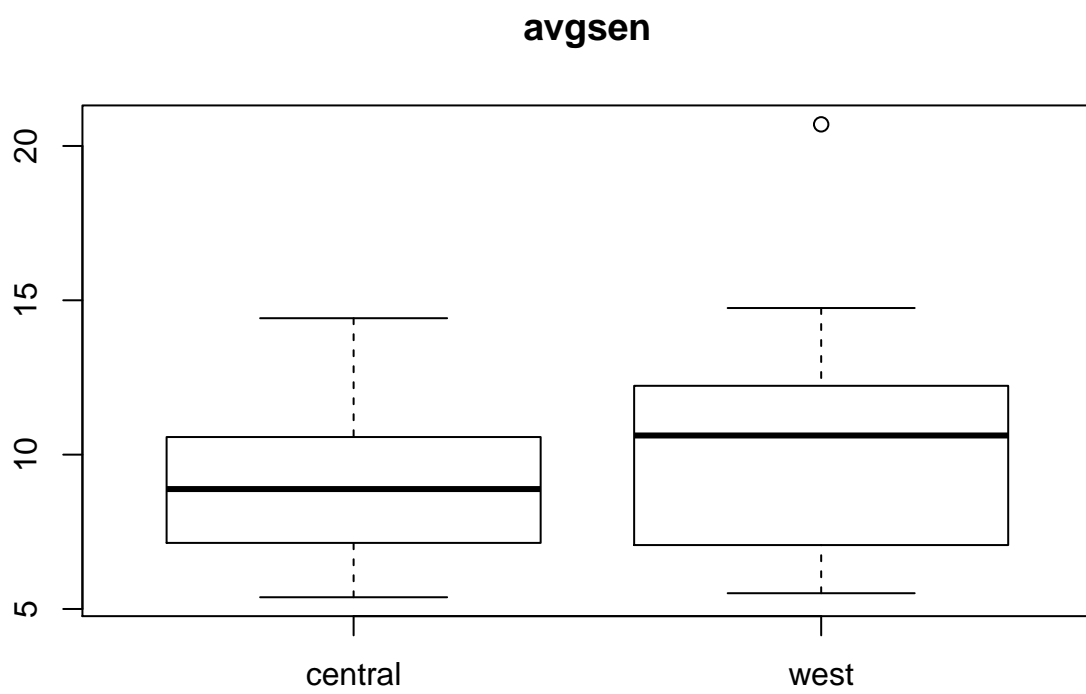


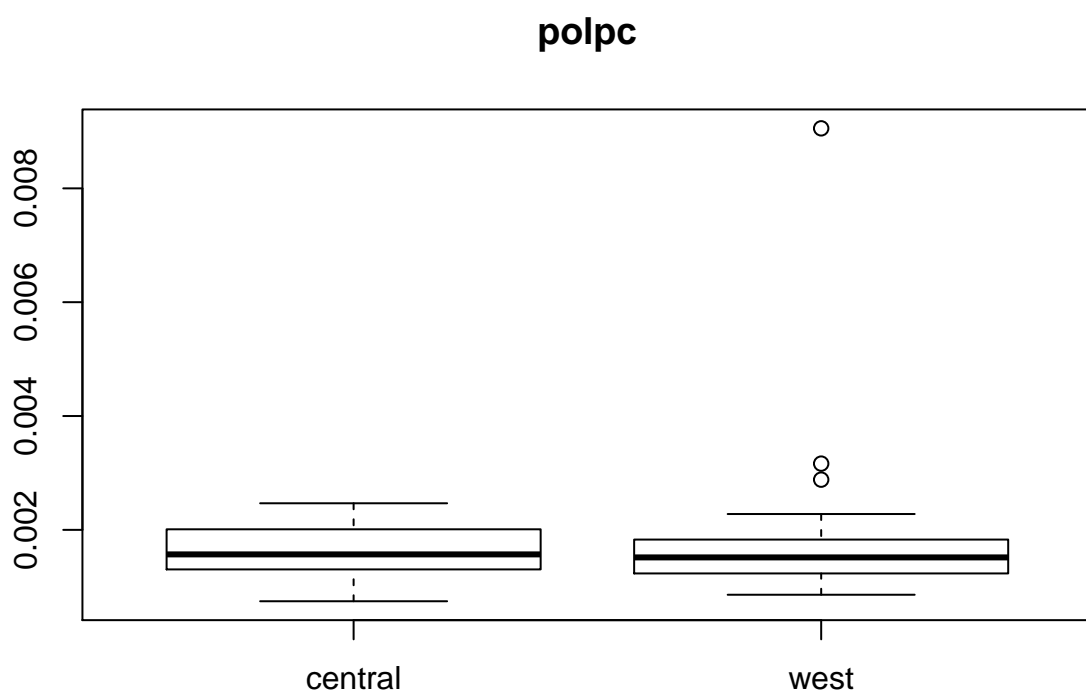


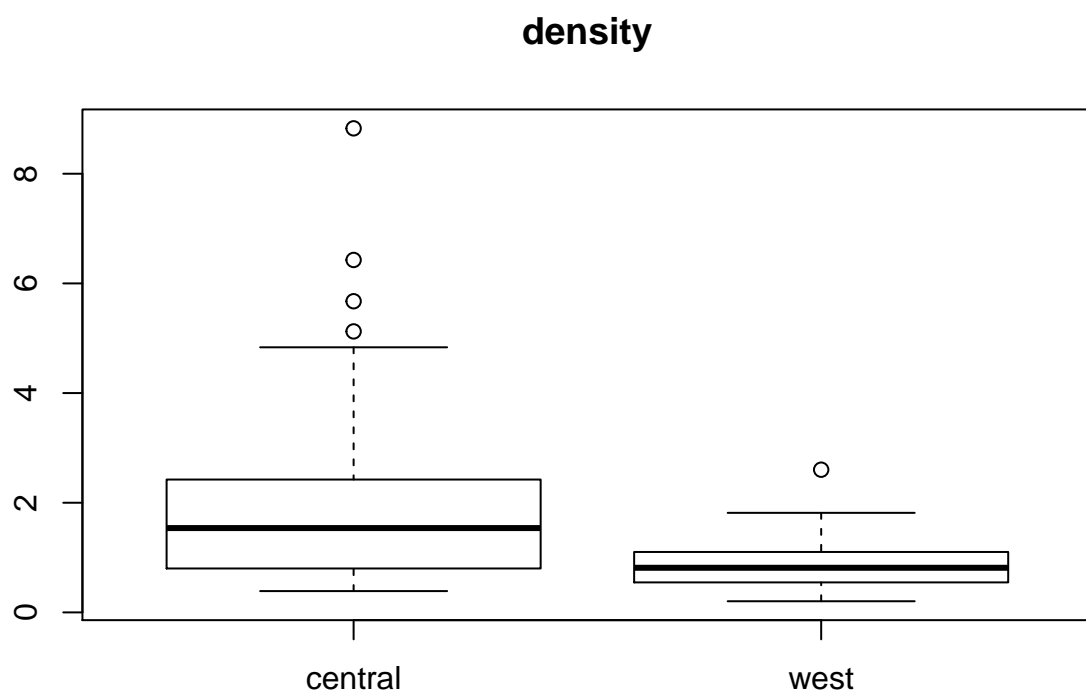
prbconv

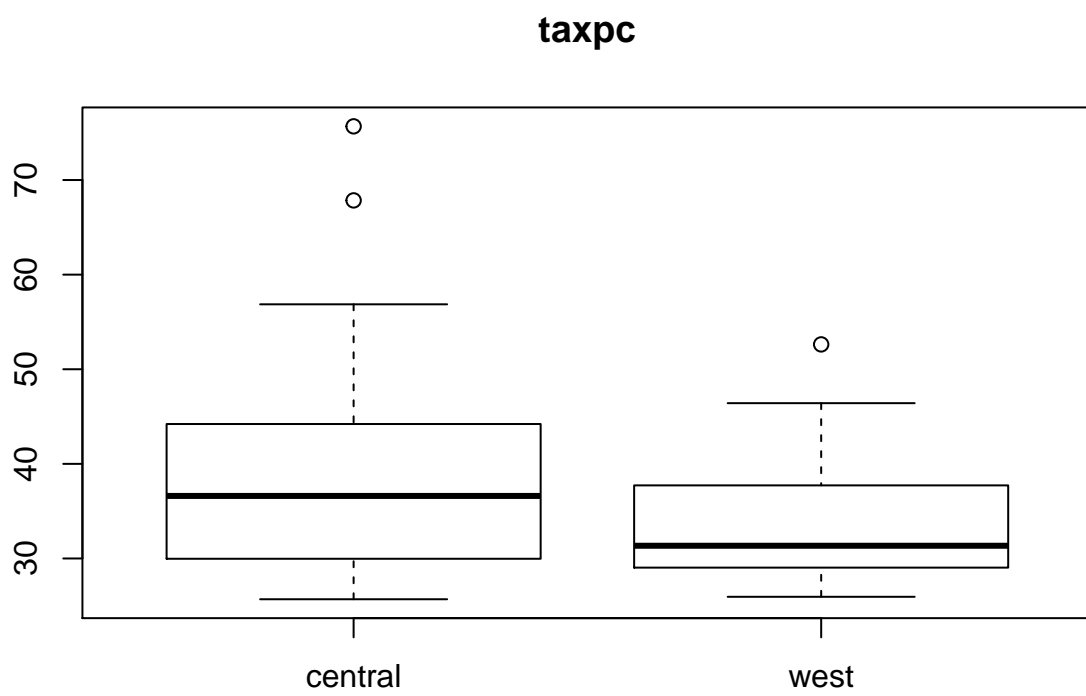




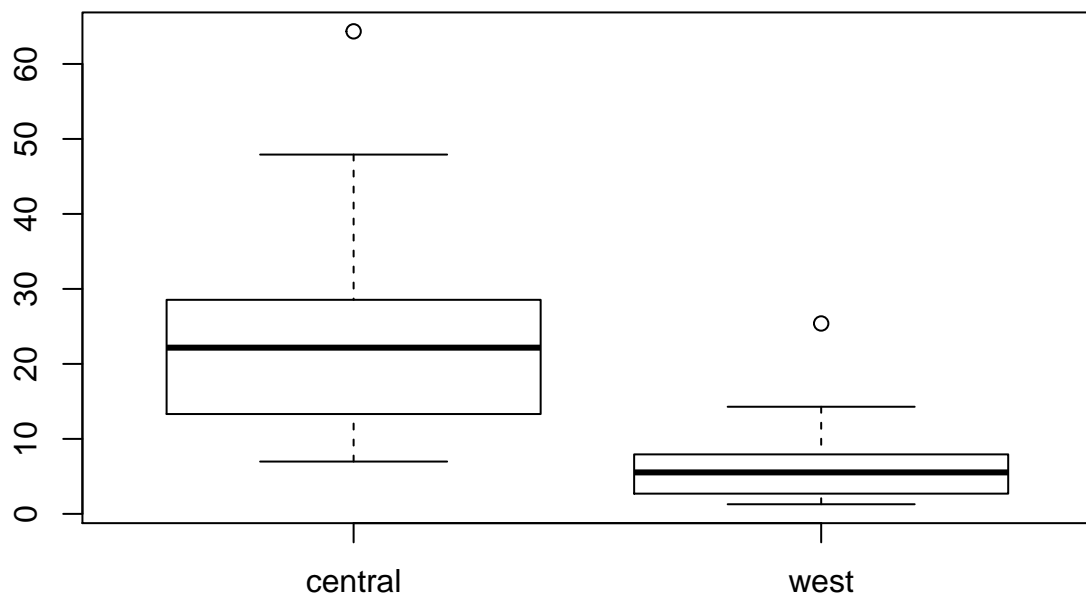


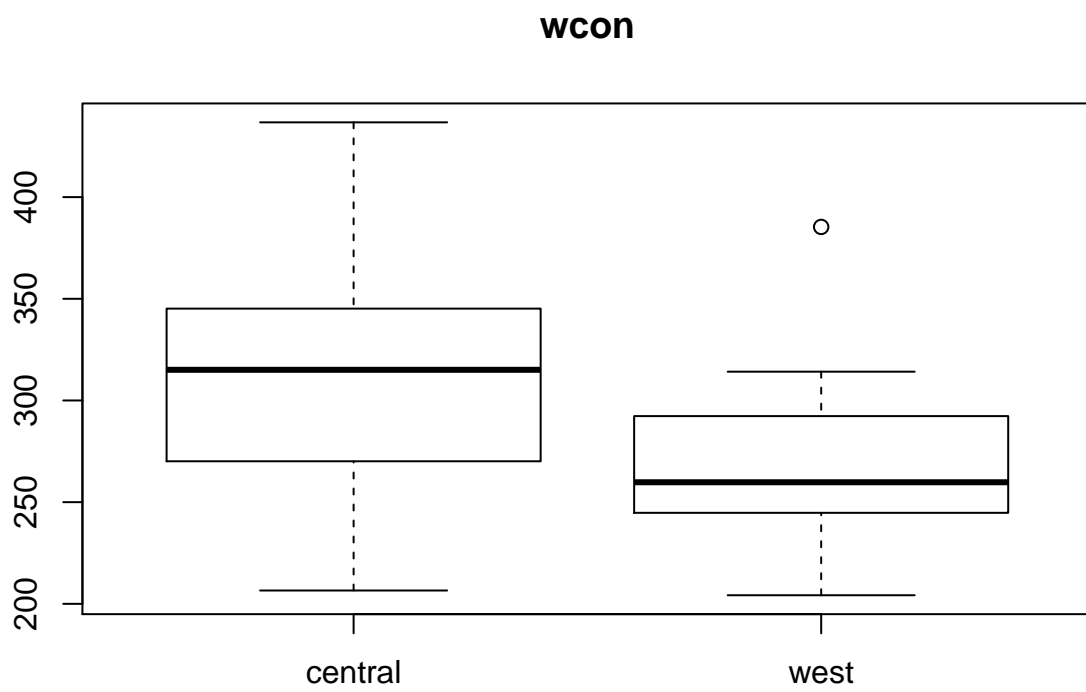


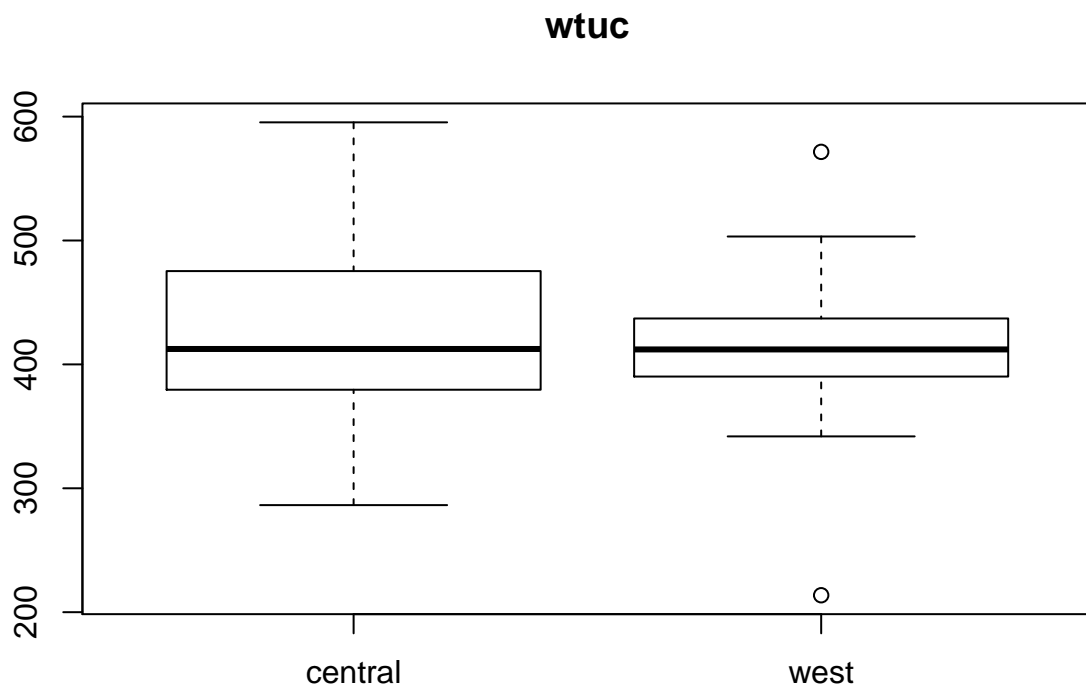


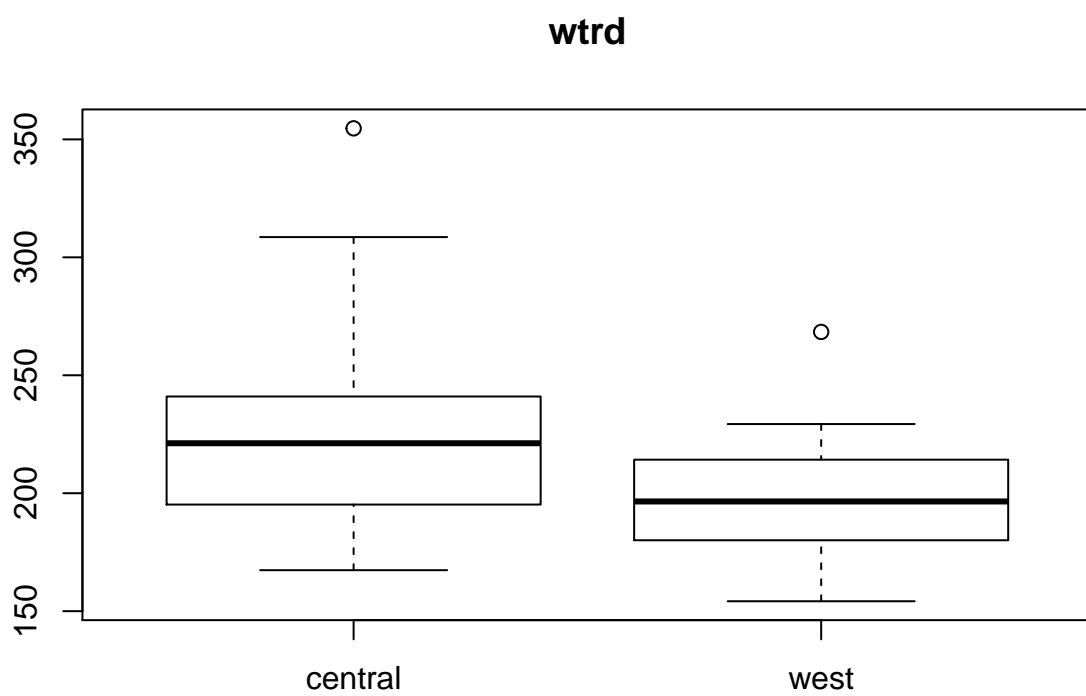


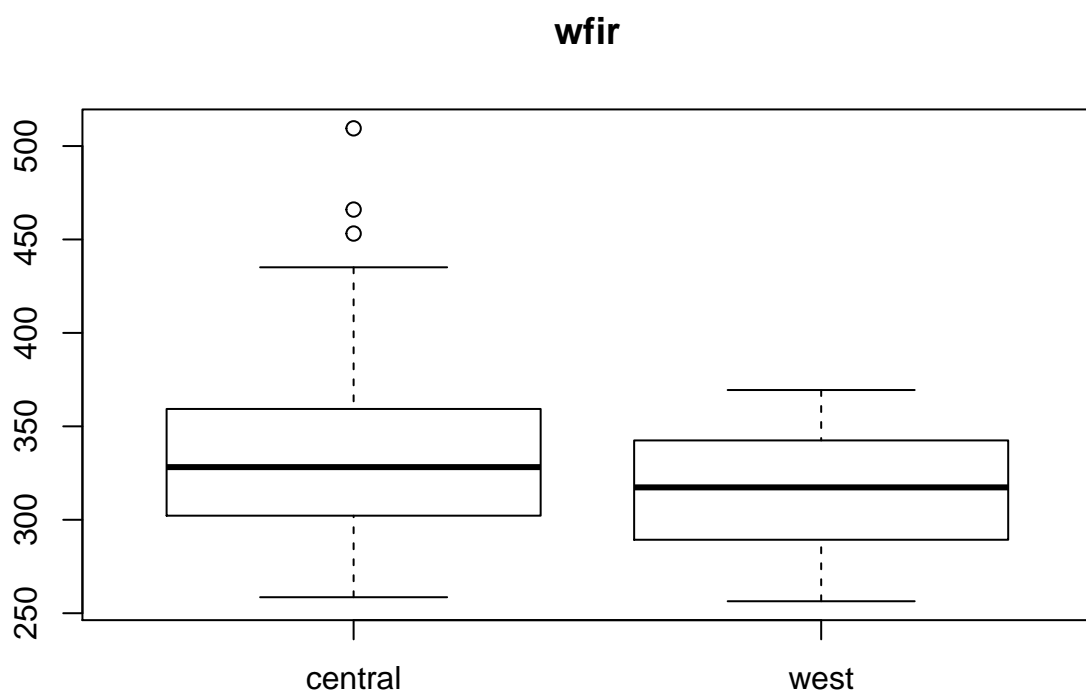
pctmin80

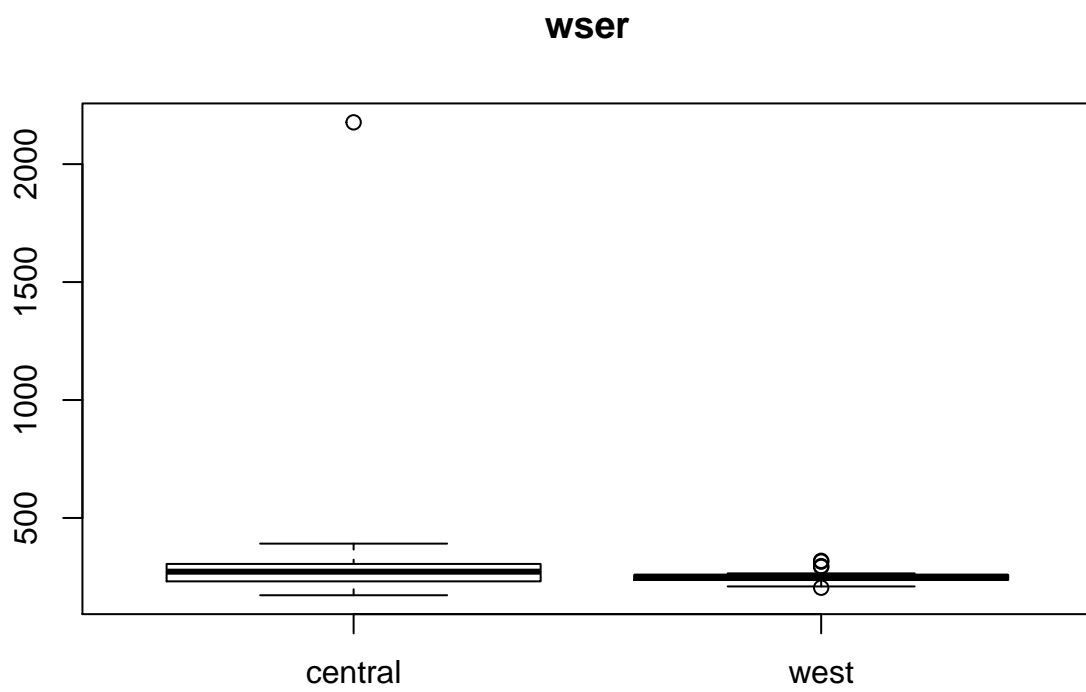


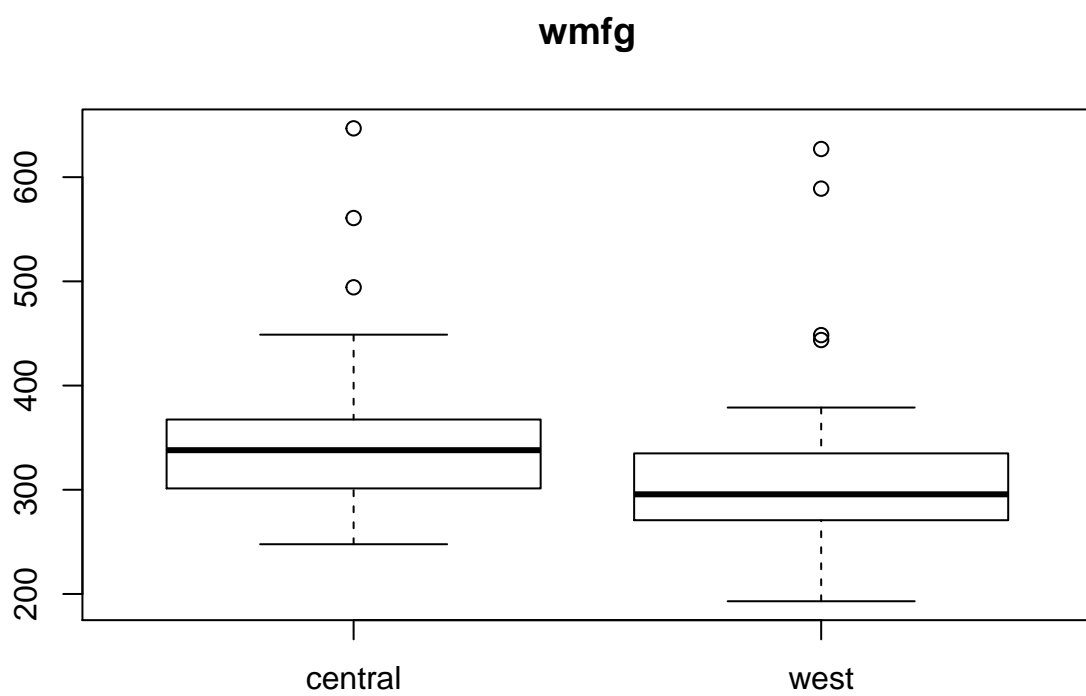


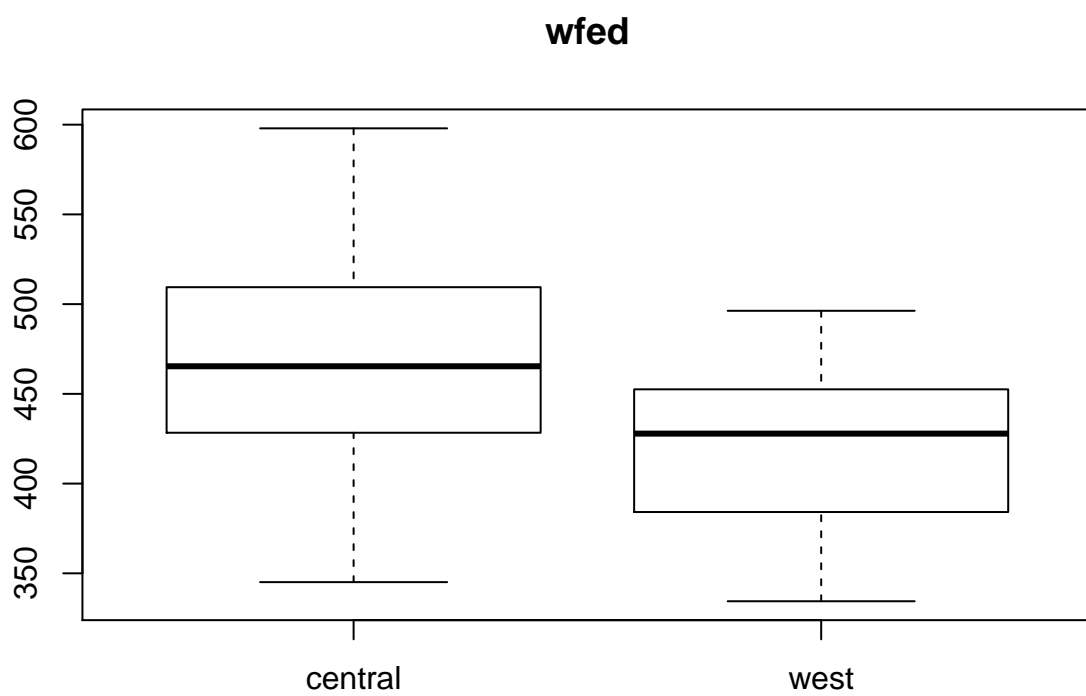


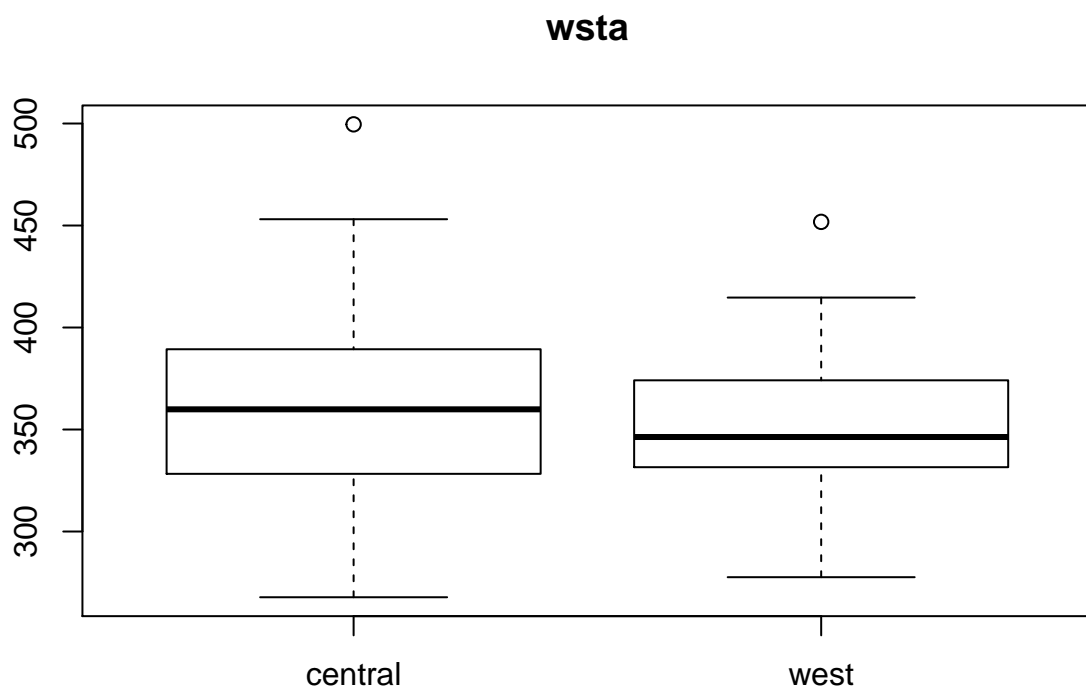


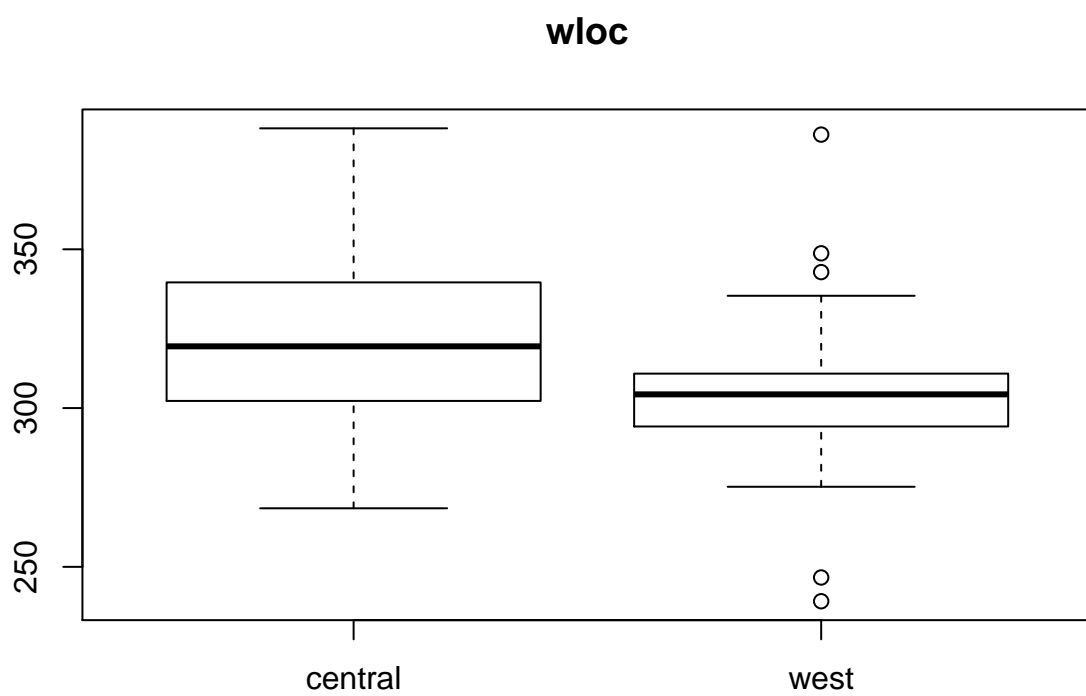


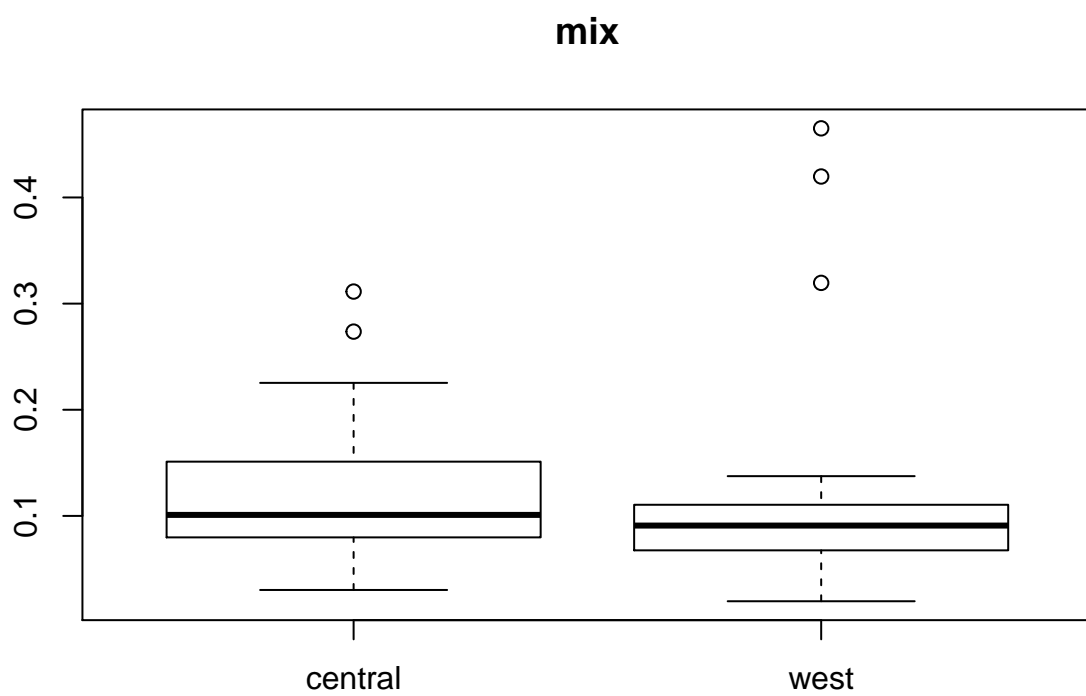




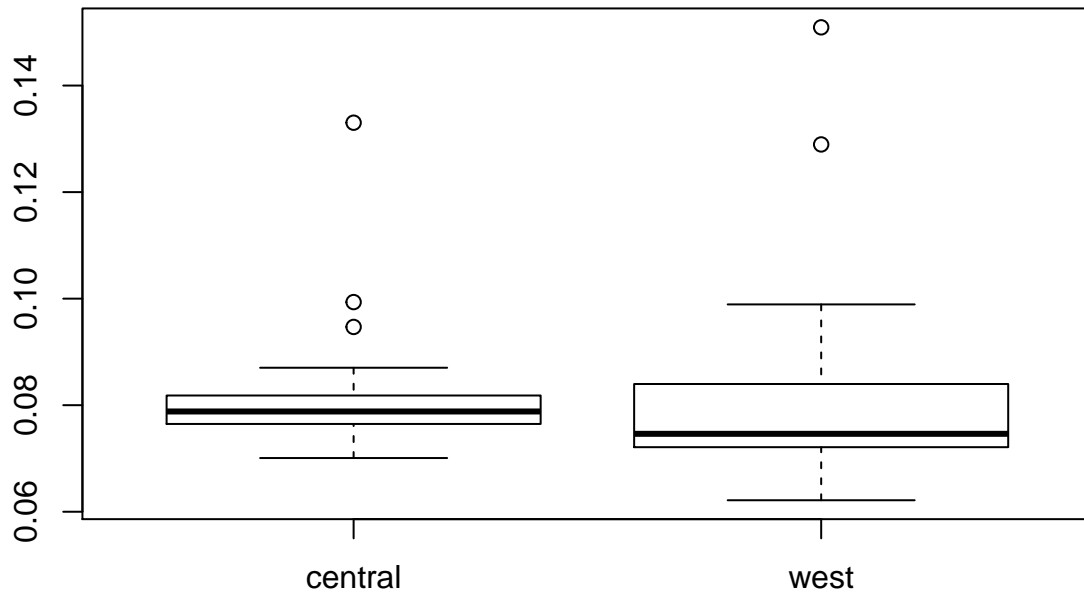








pctymle



```
##      crmrte      prbarr      prbconv      prbpris      avgsen
## stats Numeric,10 Numeric,10 Numeric,10 Numeric,10 Numeric,10
## n      Numeric,2  Numeric,2  Numeric,2  Numeric,2  Numeric,2
## conf   Numeric,4  Numeric,4  Numeric,4  Numeric,4  Numeric,4
## out    0.0989659  1.09091   Numeric,2  0.15       20.7
## group  1         2         Numeric,2  2         2
## names  Character,2 Character,2 Character,2 Character,2 Character,2
##      polpc      density      taxpc      pctmin80      wcon
## stats Numeric,10 Numeric,10 Numeric,10 Numeric,10 Numeric,10
## n      Numeric,2  Numeric,2  Numeric,2  Numeric,2  Numeric,2
## conf   Numeric,4  Numeric,4  Numeric,4  Numeric,4  Numeric,4
## out    Numeric,3  Numeric,5  Numeric,3  Numeric,2  385.3424
## group  Numeric,3  Numeric,5  Numeric,3  Numeric,2  2
## names  Character,2 Character,2 Character,2 Character,2 Character,2
##      wtuc      wtrd      wfir      wser      wmfg
## stats Numeric,10 Numeric,10 Numeric,10 Numeric,10 Numeric,10
## n      Numeric,2  Numeric,2  Numeric,2  Numeric,2  Numeric,2
## conf   Numeric,4  Numeric,4  Numeric,4  Numeric,4  Numeric,4
## out    Numeric,2  Numeric,2  Numeric,3  Numeric,6  Numeric,7
## group  Numeric,2  Numeric,2  Numeric,3  Numeric,6  Numeric,7
## names  Character,2 Character,2 Character,2 Character,2 Character,2
##      wfed      wsta      wloc      mix      pctymle
## stats Numeric,10 Numeric,10 Numeric,10 Numeric,10 Numeric,10
## n      Numeric,2  Numeric,2  Numeric,2  Numeric,2  Numeric,2
## conf   Numeric,4  Numeric,4  Numeric,4  Numeric,4  Numeric,4
## out    Numeric,0  Numeric,2  Numeric,5  Numeric,5  Numeric,5
```



```
## group Numeric,0 Numeric,2 Numeric,5 Numeric,5 Numeric,5
## names Character,2 Character,2 Character,2 Character,2 Character,2
```

Comparing central vs west,

central has higher

- crime rate
- density
- tax revenue per capita
- percentage of minority
- percentage of young male
- wages in general are higher except for wtuc

west has higher

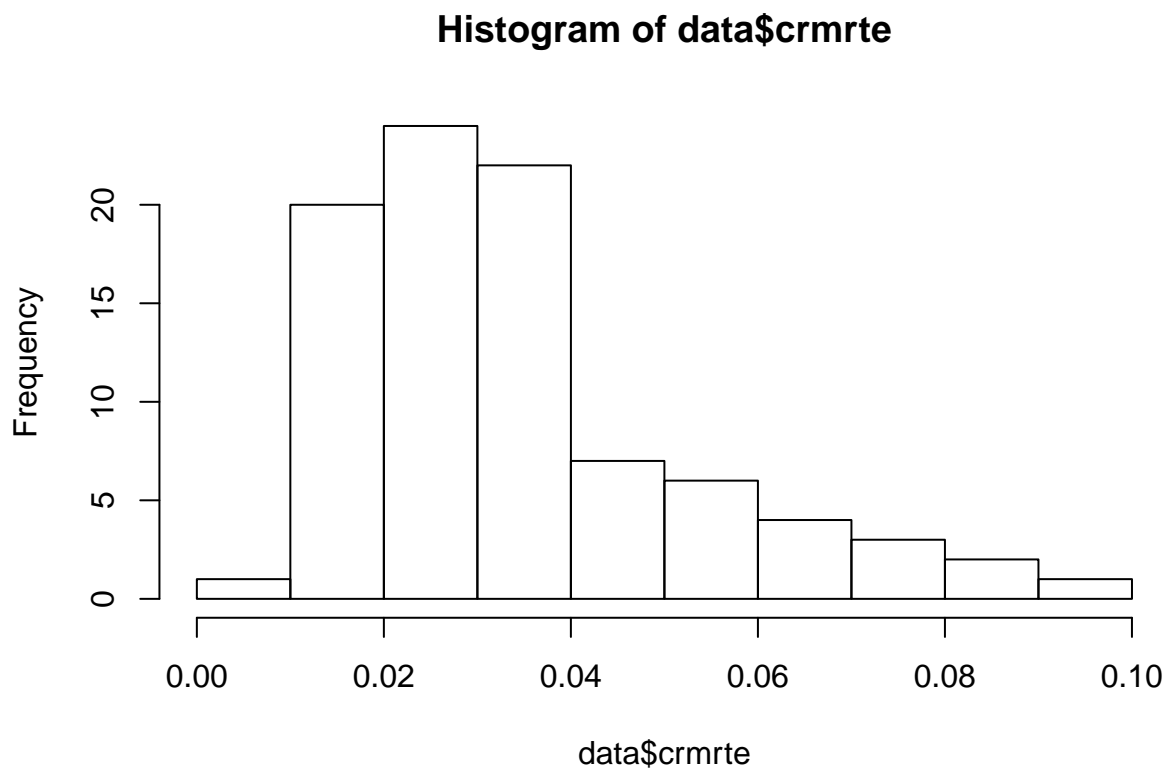
- average sentence

central and west are about the same

- probability of arrest
- probability of conviction
- probability of prison sentence
- police per capita
- face to face confrontation (central has slightly higher)

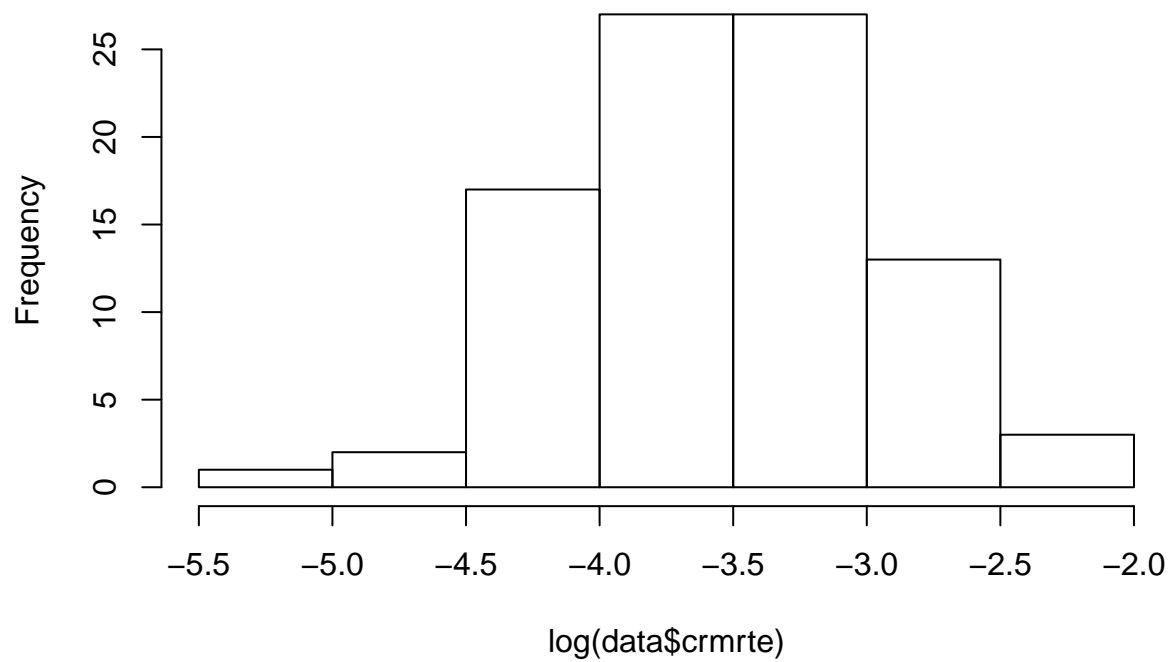
potential dependent variable: crime rate

```
hist(data$crmrte)
```



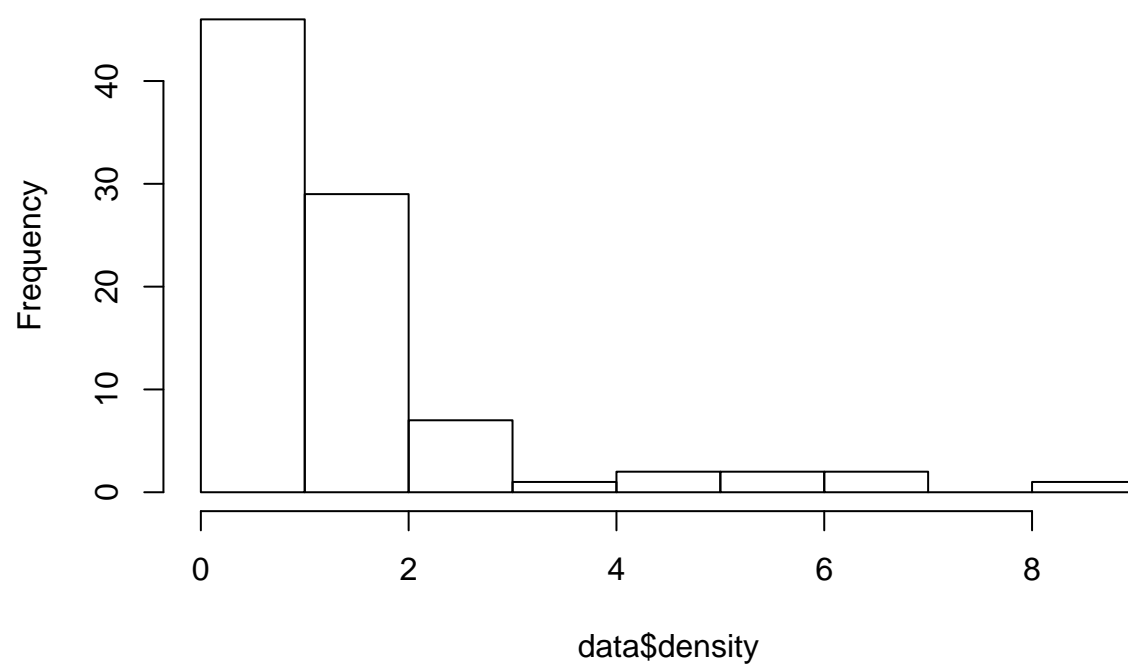
```
hist(log(data$crmrte))
```

Histogram of log(data\$crmrte)



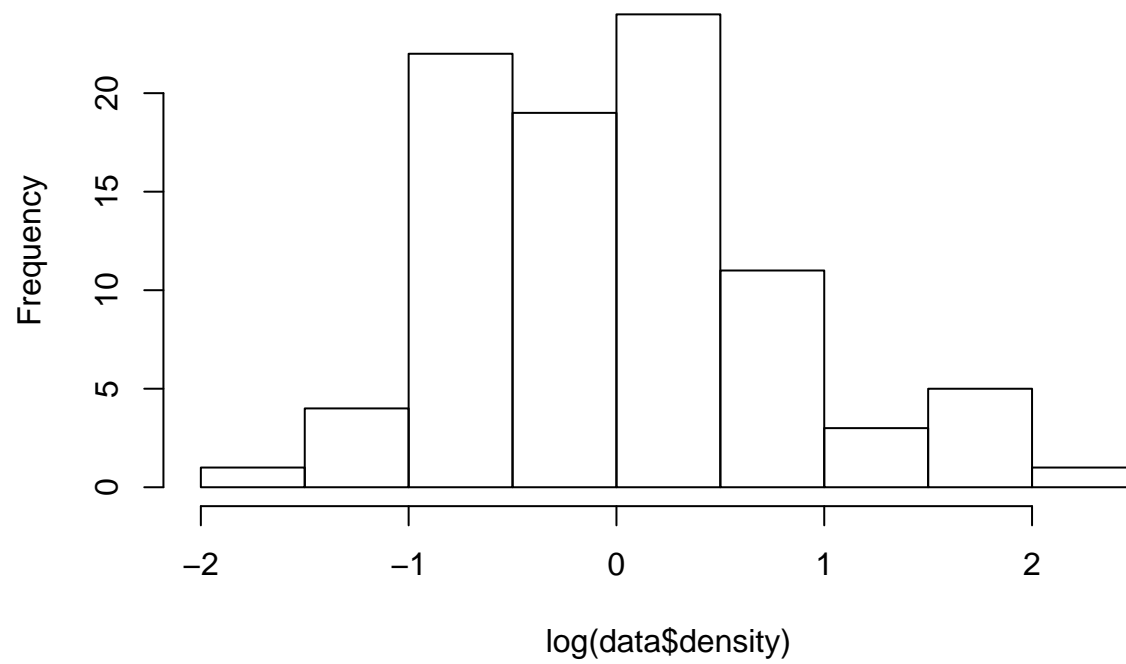
```
# potential independent variables  
# 1. density  
hist(data$density)
```

Histogram of data\$density



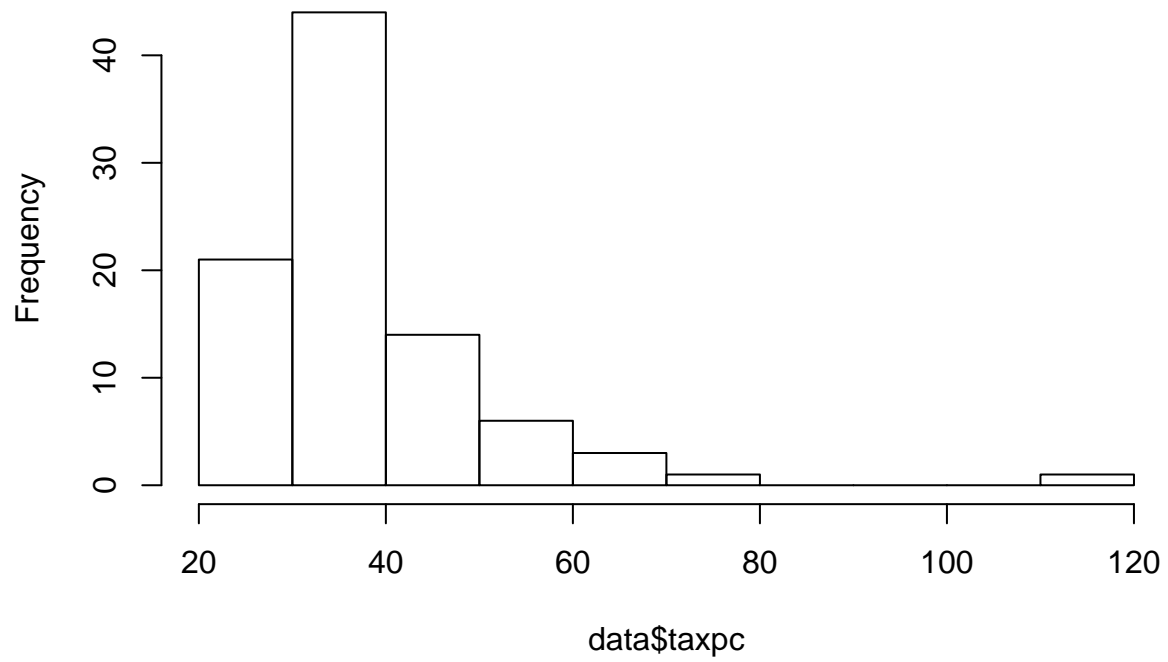
```
hist(log(data$density))
```

Histogram of $\log(\text{data\$density})$

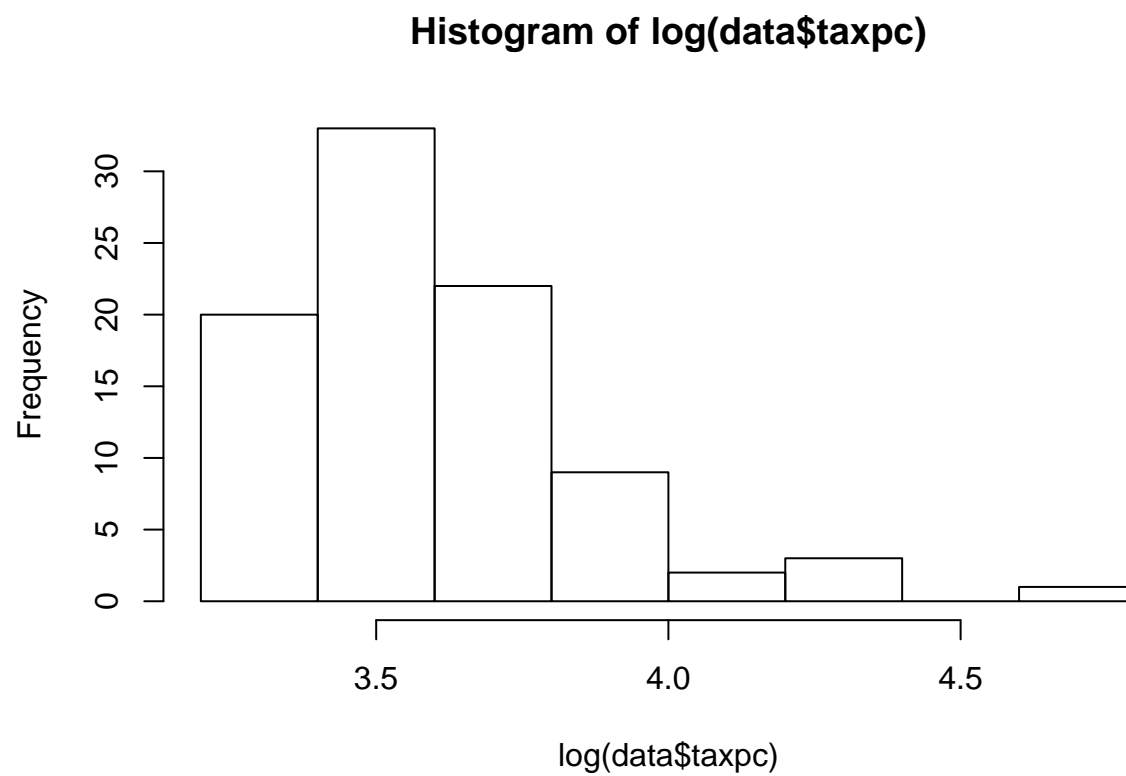


```
# 2. taxpc  
hist(data$taxpc)
```

Histogram of data\$taxpc

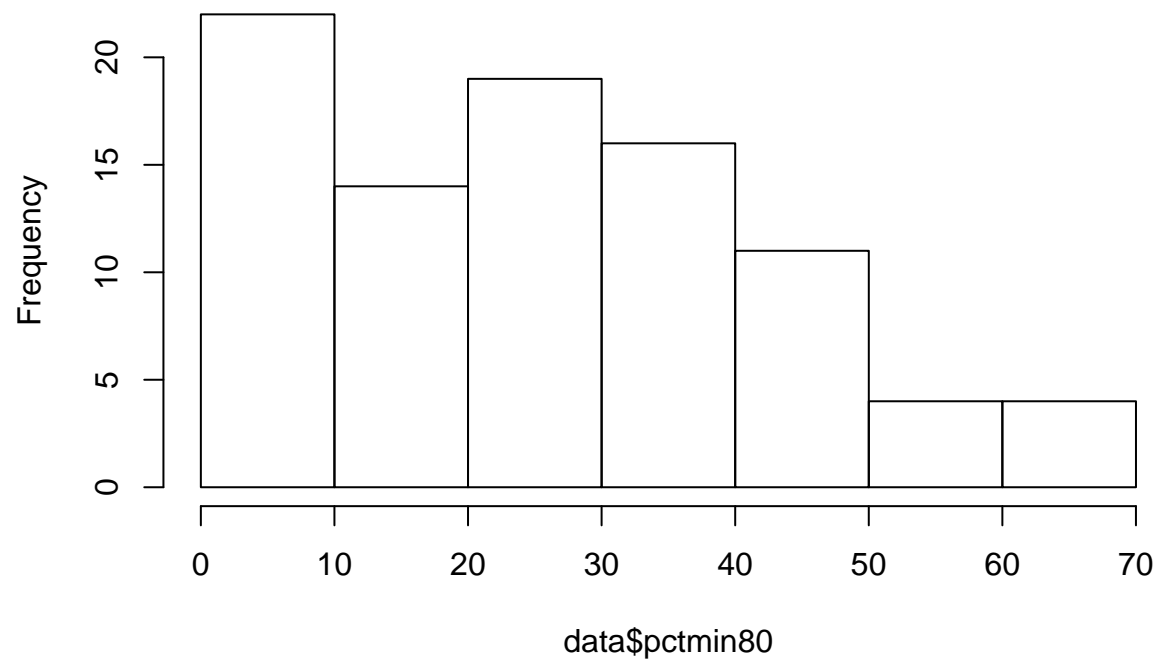


```
hist(log(data$taxpc))
```



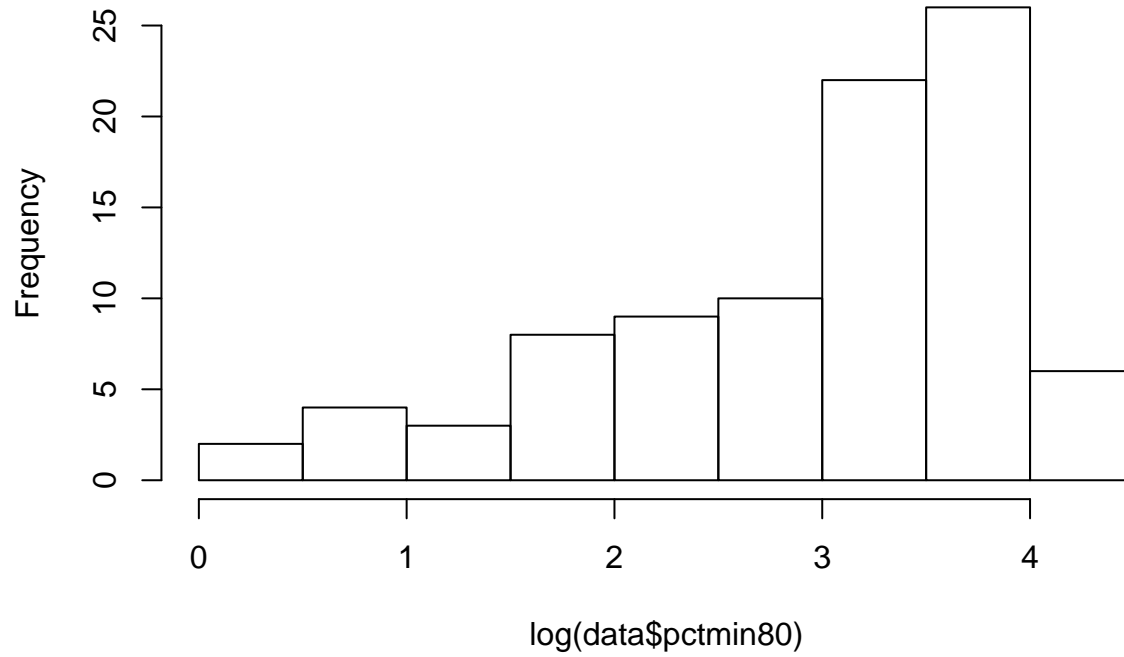
```
# log didn't quite change much because of one outlier  
# 3. pctmin80  
hist(data$pctmin80)
```

Histogram of data\$pctmin80



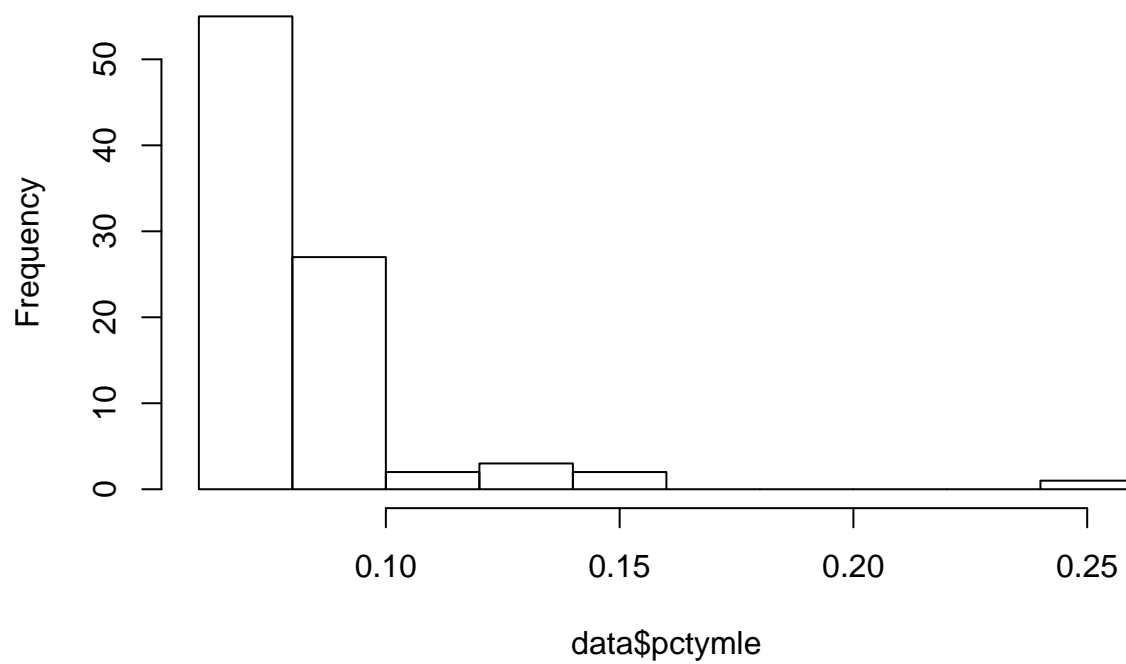
```
hist(log(data$pctmin80))
```

Histogram of $\log(\text{data}\$pctmin80)$



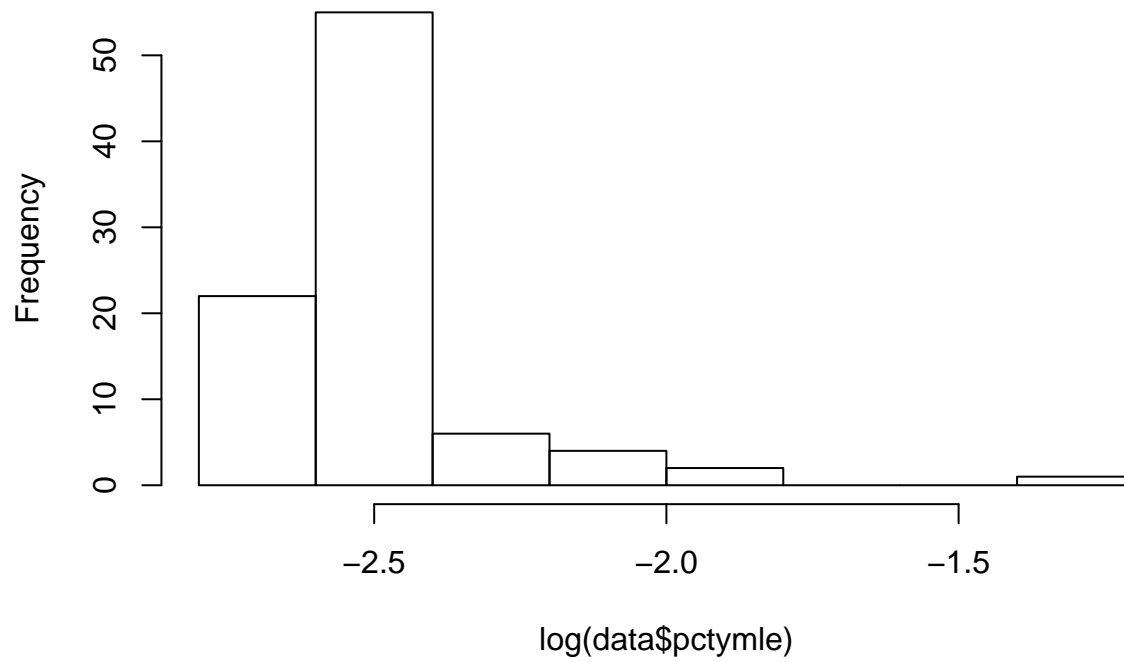
```
# better leave variable as is because applying a log skewed the distribution  
# 4. pctymle  
hist(data$pctymle)
```


Histogram of data\$pctymle



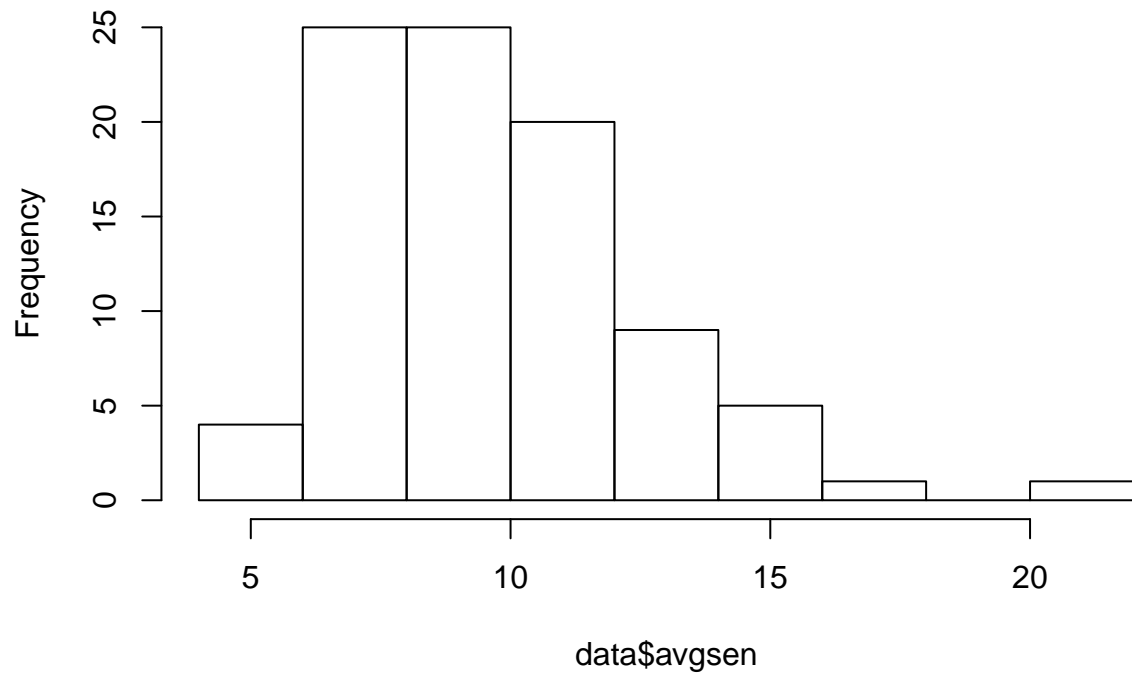
```
hist(log(data$pctymle))
```

Histogram of $\log(\text{data\$pctymle})$

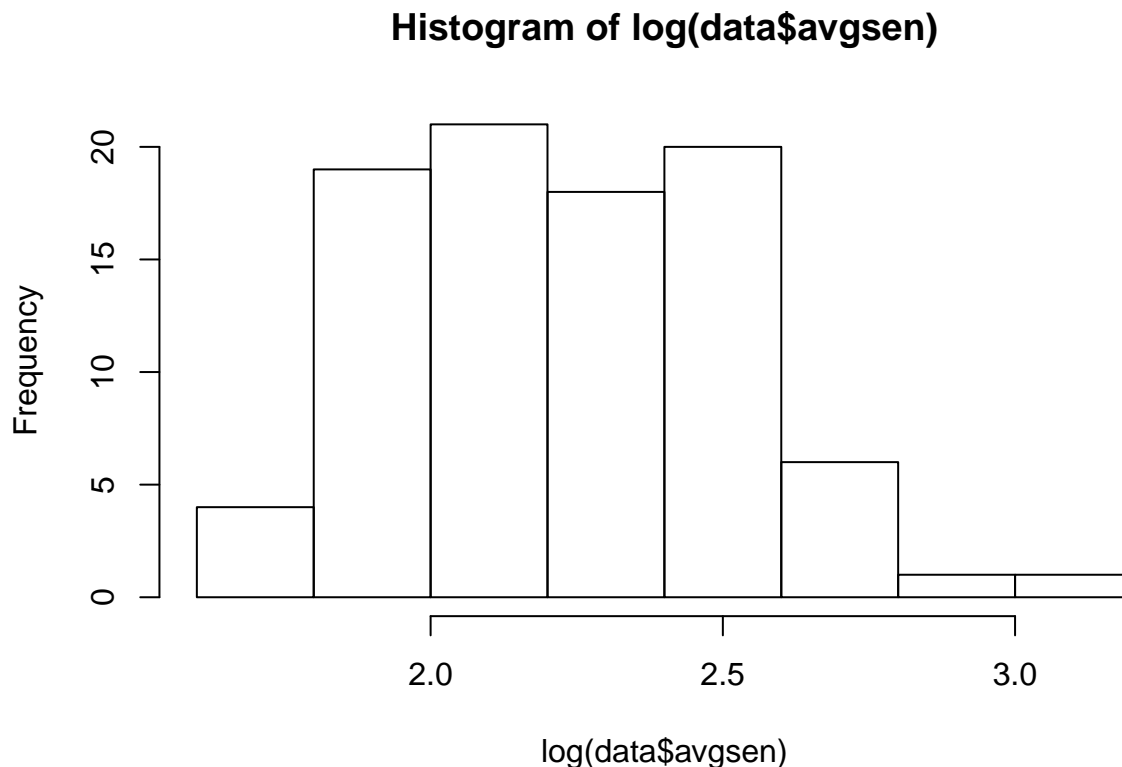


```
# log didn't quite change much because of one outlier  
# 5. avgse  
hist(data$avgse)
```

Histogram of data\$avgsen



```
hist(log(data$avgsen))
```



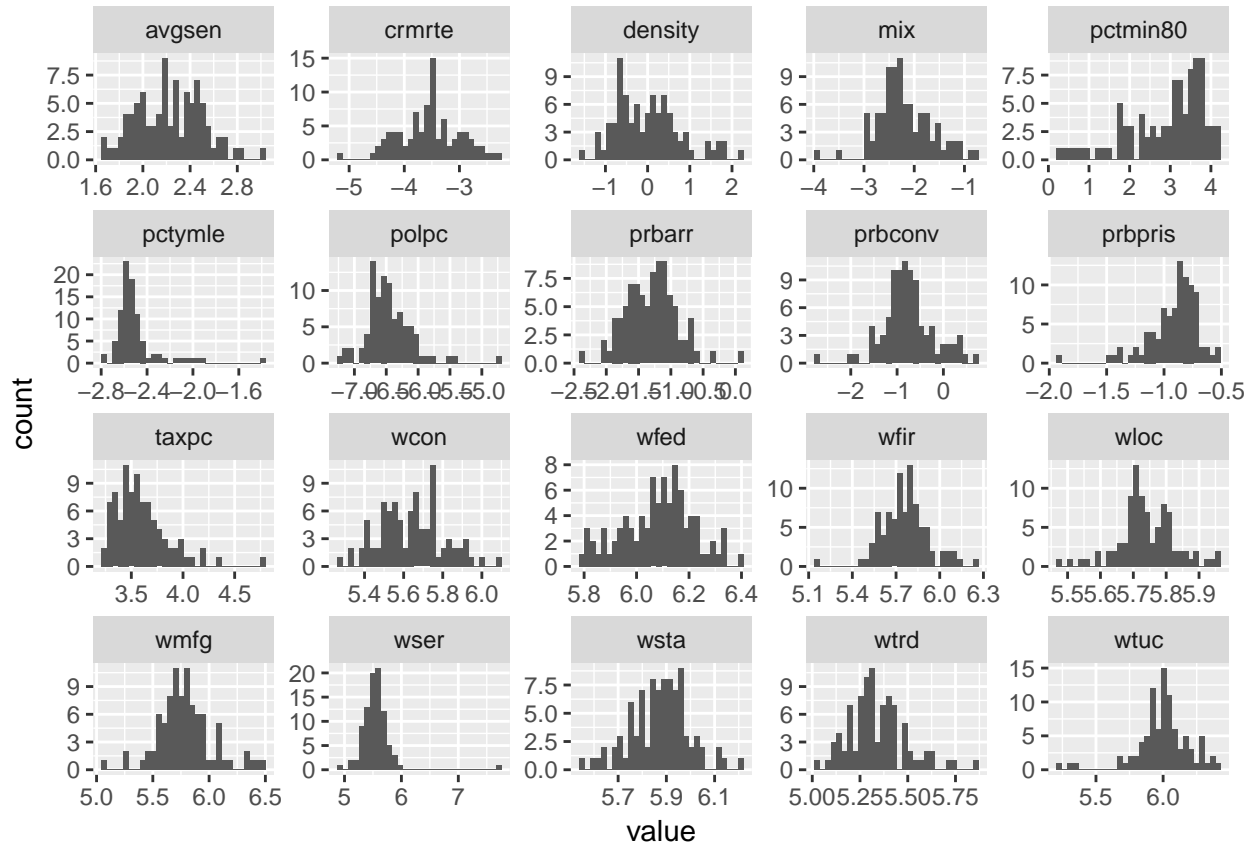
already verified that these variables are not correlated to each other

Observations

- All observations were recorded for the year of 1987.
- *urban*, *central*, *west* are categorical variables.
- Most wages variables (*wcon*, *wfed*, *wfir*, *wmfg*, *wser*, *wsta*, *wtrd*) have a positively skewed distribution which may be due to few number people getting paid above the average.
- *wtuc*, *wloc* appear to be normally distributed.
- *prbarr*, *prbconv* appear to be positively skewed while *prbpris* is more negatively skewed.
- *avgsen* appears to be positively skewed.
- *crrmrte*, *density*, *polpc*, *prbconv*: The histogram indicates that these variables have positive skews. Given the variables have a meaningful zero-point, we can take the log for a more normal distribution.

```
# taking the log of the data excluding x, county, year and categorical variables
plot.log.data <- log(data[!(names(data) %in% c("X", "county", "year", "urban", "central", "west"))])
ggplot(gather(plot.log.data, aes(value))) +
  facet_wrap(~key, scales="free") +
  geom_histogram()
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```



Observations after taking logs: It looks like the log transformation of variables crmrte, mix, prbconv, wfed, wfir, wmfg, and wsta have made the distributions quite normal. This will help ensure the errors of the model are normal.

Proposed Model Based on the exploratory data analysis and general intuition about the potential determinants of crime, an initial proposed model specification and coefficient expectation are:

- Sum of all wages: Summing the wage variables across sectors may identify whether income inequality between counties may explain the difference in crime rates. Initial thoughts are that sum of wages (higher incomes) lead to lower crime rates, so a linear relationship between sum of wages and crimes committed per person is expected with a negative coefficient.
- $\log(\text{prbarr})$ Probability of arrest: There is the possibility that the probability of arrest is positively correlated with crime rate, as higher numbers of arrest can increase the number of crimes recorded.
- tax revenue per capita (taxpc): Lack of government-funded resources can take the form of a lack of educational opportunities or employment options, thus leading to rise in crime rates.

Model 1: $\log(\text{crmrte}) = \beta_0 + \beta_1(\text{allwages}) + \beta_2 \log(\text{prbarr}) + \beta_3(\text{taxrev}) + u$

Model Building Process

TODO: A model building process, supported by exploratory analysis. Your EDA should be interspersed with, and support, your modeling decisions. In particular, you should use exploratory techniques to address *

What transformations to apply to variables and what new variables should be created. *

What variables should be included in each model *

Whether model assumptions are met

```
$density [1] "8 : 3.93455100059509, 4.38875865936279, 4.8347339630127, 5.12442398071289, 5.6744966506958, 6.28648662567139, 6.42718458175659, 8.82765197753906"
```

```
$taxpc [1] "6 : 56.8621063232422, 61.1525115966797, 67.6796340942383, 67.8479766845703, 75.6724319458008, 119.761451721191"
```

Model Specifications

Model 1

TODO: * One model with only the explanatory variables of key interest (possibly transformed, as determined by your EDA), and no other covariates.

$$\log(\text{crmrte}) = \beta_0 + \beta_1 \cdot \log(\text{density}) + \beta_2 \cdot (\text{taxpc}) + u$$

```
m1 <- lm(log(crmrte) ~ log(density) + taxpc, data=data)
```

TODO: For your first model, a detailed assessment of the 6 CLM assumptions. For additional models, you should check all assumptions, but only highlight major differences from your first model in your report.

CLM 1 - A linear model

The model is specified such that the dependent variable is a linear function of the explanatory variables. Is the assumption valid? *Yes*

Response: No response required.

CLM 2 - Random Sampling

TODO

CLM 3 - Multicollinearity

```
# show correlation
data$log.density <- log(data$density)
cor(data.matrix(subset(data, select=c("density", "log.density", "taxpc"))))
```

```
##           density log.density      taxpc
## density      1.0000000  0.8896747 0.3204737
## log.density  0.8896747  1.0000000 0.1841222
## taxpc        0.3204737  0.1841222 1.0000000
```

```
# show vif
vif(m1)
```

```
## log(density)      taxpc
##      1.035091      1.035091
```

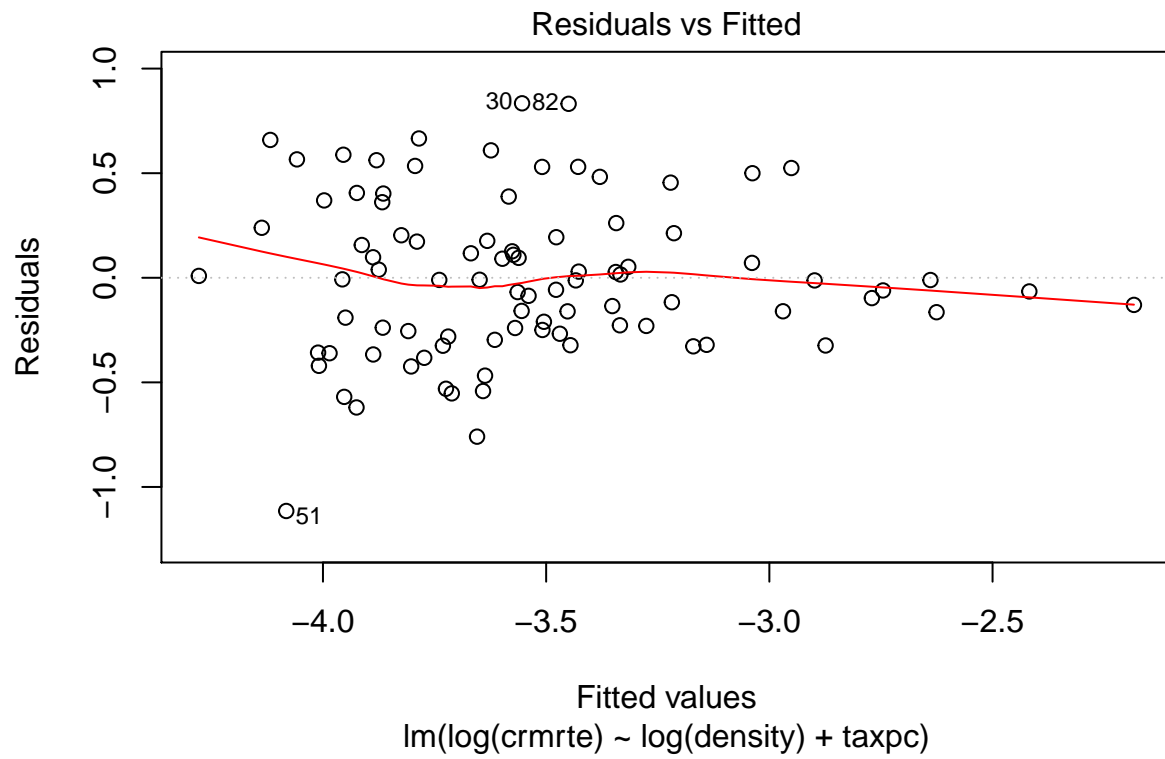
The two explanatory variables (log.density and taxpc) are not perfectly correlated and the VIFs are low (less than 4), so there is no perfect multicollinearity of the independent variables.

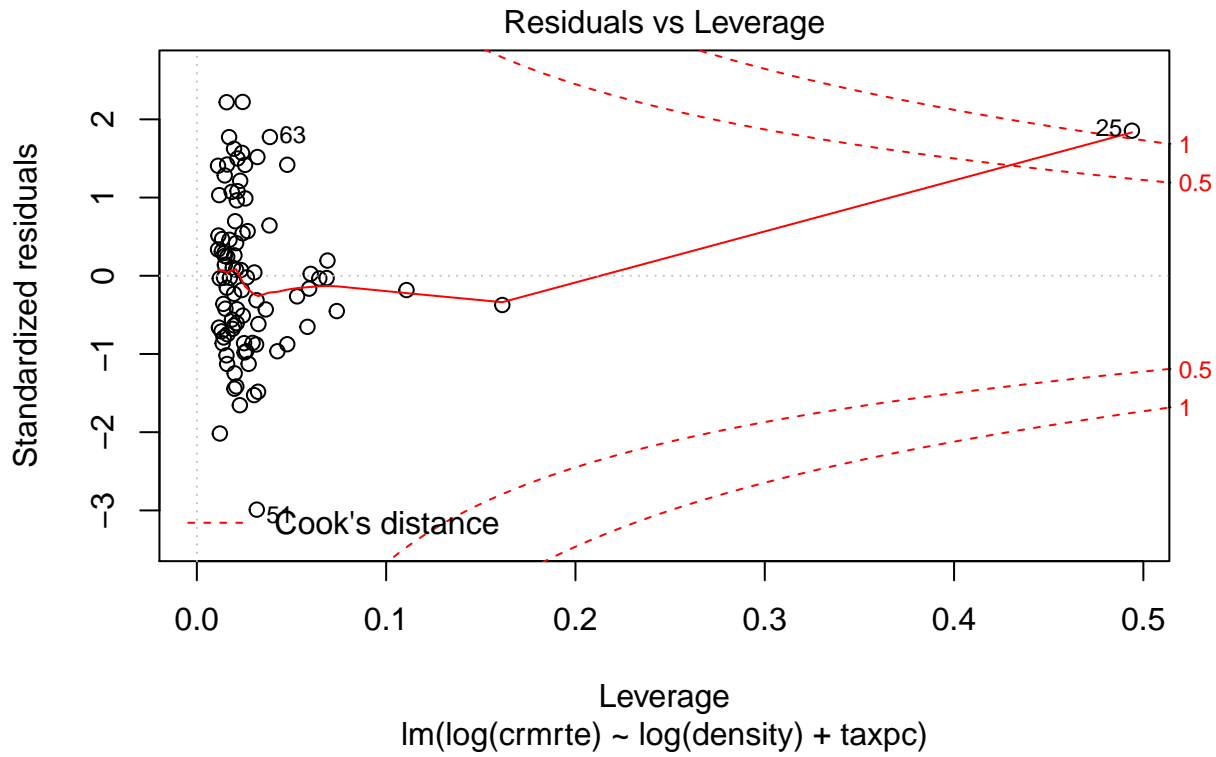
Is the assumption valid? *Yes*

Response: No response required.

CLM 4 - Zero-Conditional Mean

```
plot(m1, which=c(1, 5))
```





```
# exogenous check
cov(log(data$density), m1$residuals)
```

```
## [1] -8.896016e-18
```

```
cov(data$taxpc, m1$residuals)
```

```
## [1] -4.453513e-16
```

The covariances of the two independent variables with the residuals are very close to zero indicating they are likely exogenous.

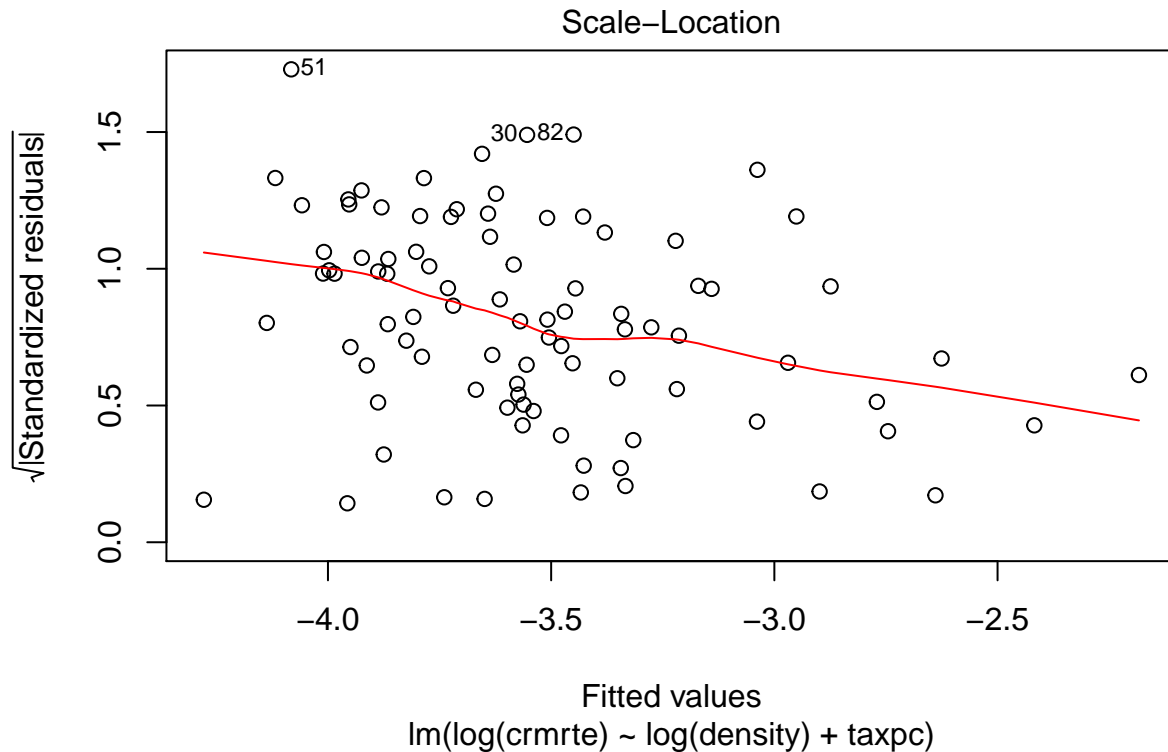
There is a data point with a large Cook's distance.

Is the assumption valid?

Response:

CLM 5 - Homoscedasticity

```
# spreadLevelPlot(m1)
plot(m1, which=3)
```

```
bptest(m1)
```

```
##
## studentized Breusch-Pagan test
##
## data: m1
## BP = 7.2256, df = 2, p-value = 0.02698
not significant p-value for Breusch-Pagan test -> assumption of homoscedasticity met
```

```
ncvTest(m1)
```

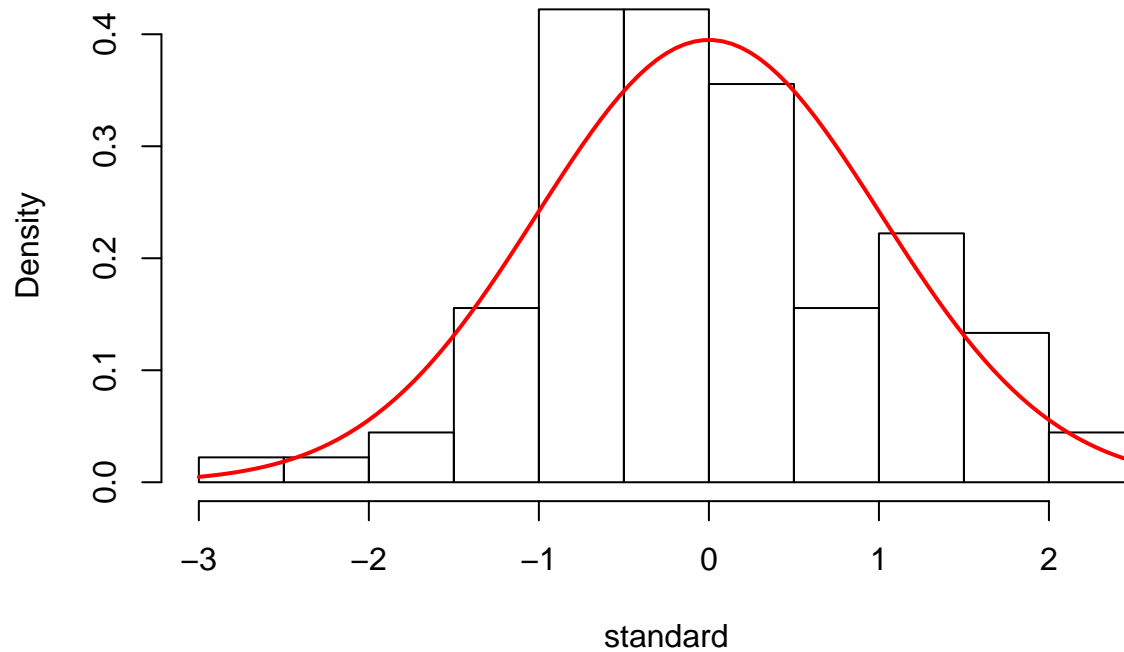
```
## Non-constant Variance Score Test
## Variance formula: ~ fitted.values
## Chisquare = 6.968285 Df = 1 p = 0.008296697
significant p-value for Score Test -> assumption of homoscedasticity met
Is the assumption valid? Yes
```

Response:

CLM 6 - Normality of residuals

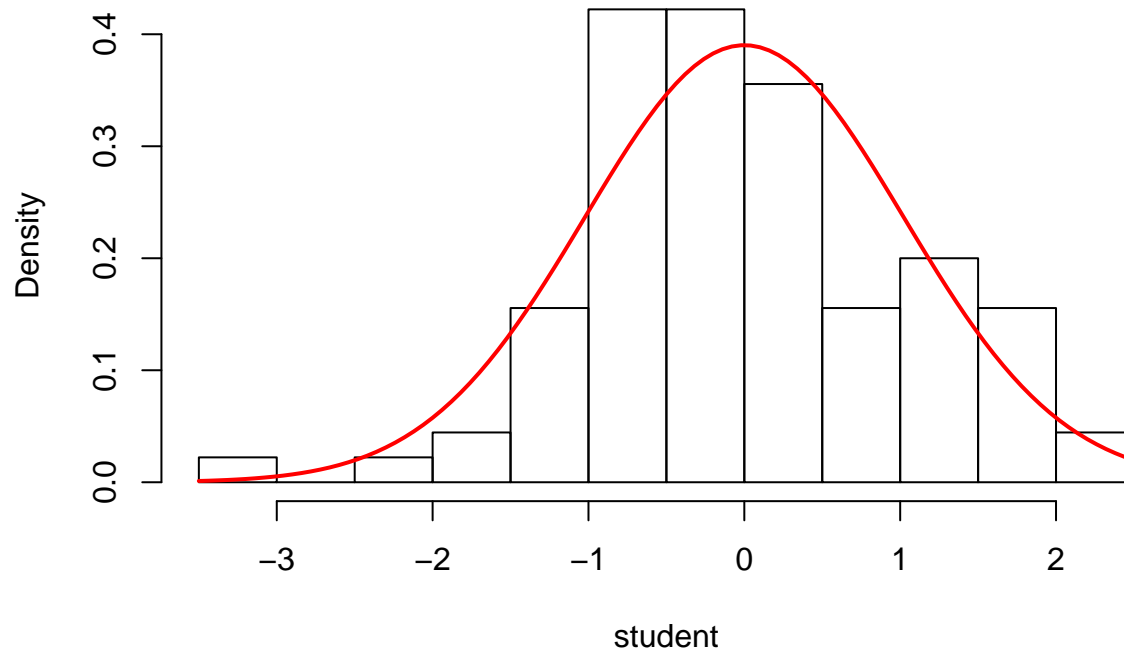
```
# normality of standard residuals
standard <- rstandard(m1)
hist(standard, main="Histogram of standard residuals", freq=FALSE)
curve(dnorm(x, mean=0, sd=sd(standard)), col="red", lwd=2, add=TRUE)
```

Histogram of standard resiudals



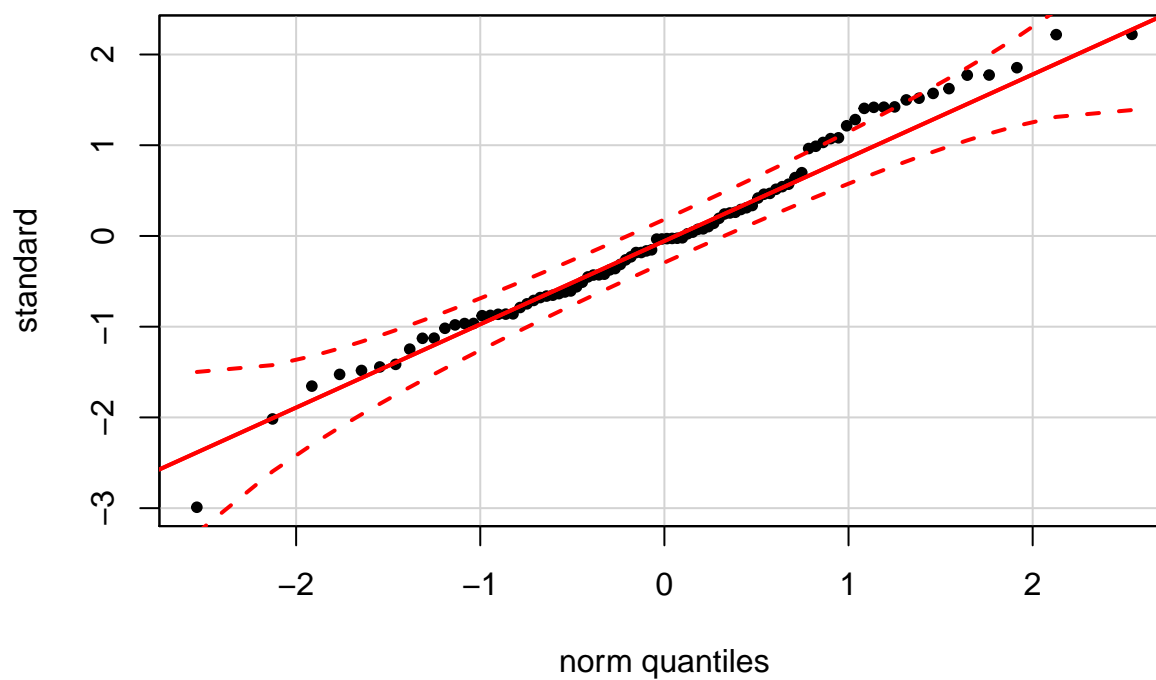
```
# normality of studentized residuals
student <- rstudent(m1)
hist(student, main="Histogram of studentized resiudals", freq=FALSE)
curve(dnorm(x, mean=0, sd=sd(student)), col="red", lwd=2, add=TRUE)
```

Histogram of studentized residuals

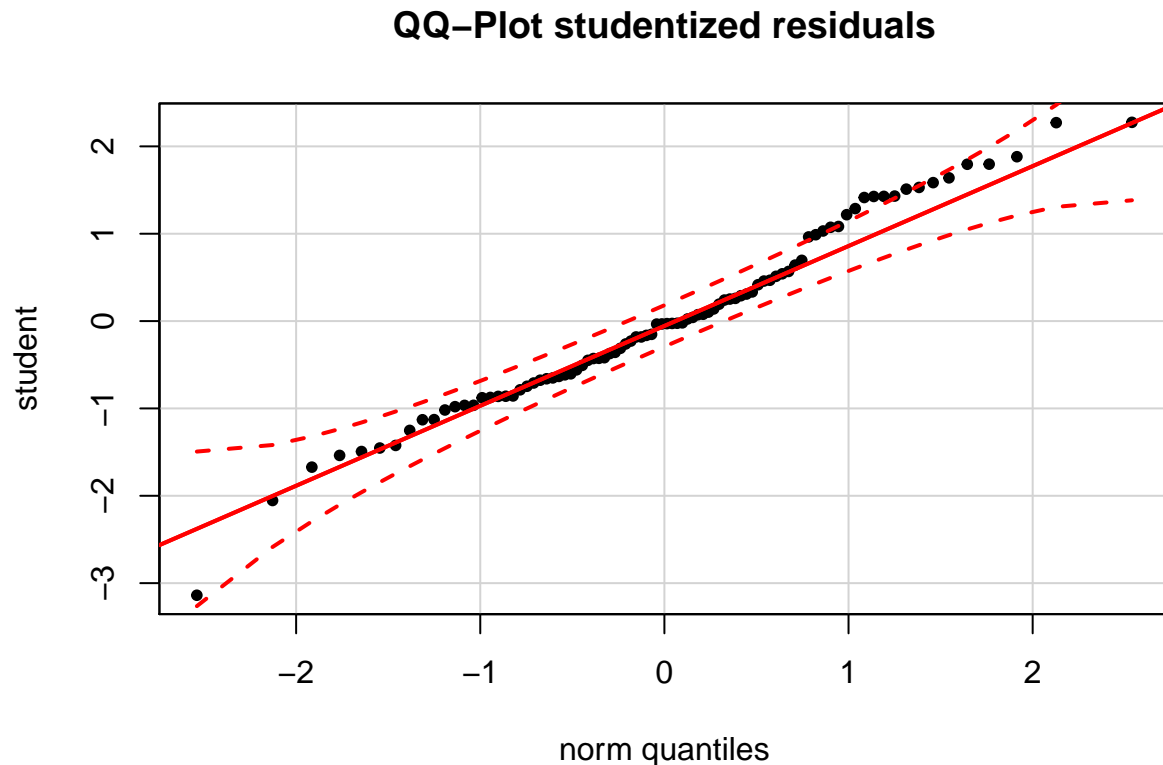


```
# QQ plot standard residuals  
qqPlot(standard, distribution="norm", pch=20, main="QQ-Plot standard residuals")  
qqline(standard, col="red", lwd=2)
```

QQ-Plot standard residuals



```
# QQ plot studentized residuals  
qqPlot(student, distribution="norm", pch=20, main="QQ-Plot studentized residuals")  
qqline(student, col="red", lwd=2)
```



The histograms in particular appear to be fairly normally distributed, while the Q-Q plot doesn't deviate significantly from normality either, so overall the residuals appear to be fairly normal. Notably the log transformation of views has helped to produce normal errors.

Is the assumption valid? *Yes*

Response: *No response required.*

TODO: * One model that includes key explanatory variables and only covariates that you believe increase the accuracy of your results without introducing bias (for example, you should not include outcome variables that will absorb some of the causal effect you are interested in). This model should strike a balance between accuracy and parsimony and reflect your best understanding of the determinants of crime.

```
m2 <- lm(log(crmrte) ~ log(density) + taxpc + urban, data=data)
```

CLM 1 - A linear model

The model is specified such that the dependent variable is a linear function of the explanatory variables. Is the assumption valid? *Yes*

Response: *No response required.*

CLM 2 - Random Sampling

TODO

CLM 3 - Multicollinearity

```
# show correlation
data$log.density <- log(data$density)
cor(data.matrix(subset(data, select=c("density", "log.density", "taxpc", "urban"))))
```

```
##           density log.density    taxpc    urban
## density      1.0000000  0.8896747 0.3204737 0.8206825
## log.density  0.8896747  1.0000000 0.1841222 0.6490078
## taxpc        0.3204737  0.1841222 1.0000000 0.3457462
## urban        0.8206825  0.6490078 0.3457462 1.0000000
```

```
# show vif
vif(m2)
```

```
## log(density)    taxpc    urban
##      1.733261    1.139396    1.901850
```

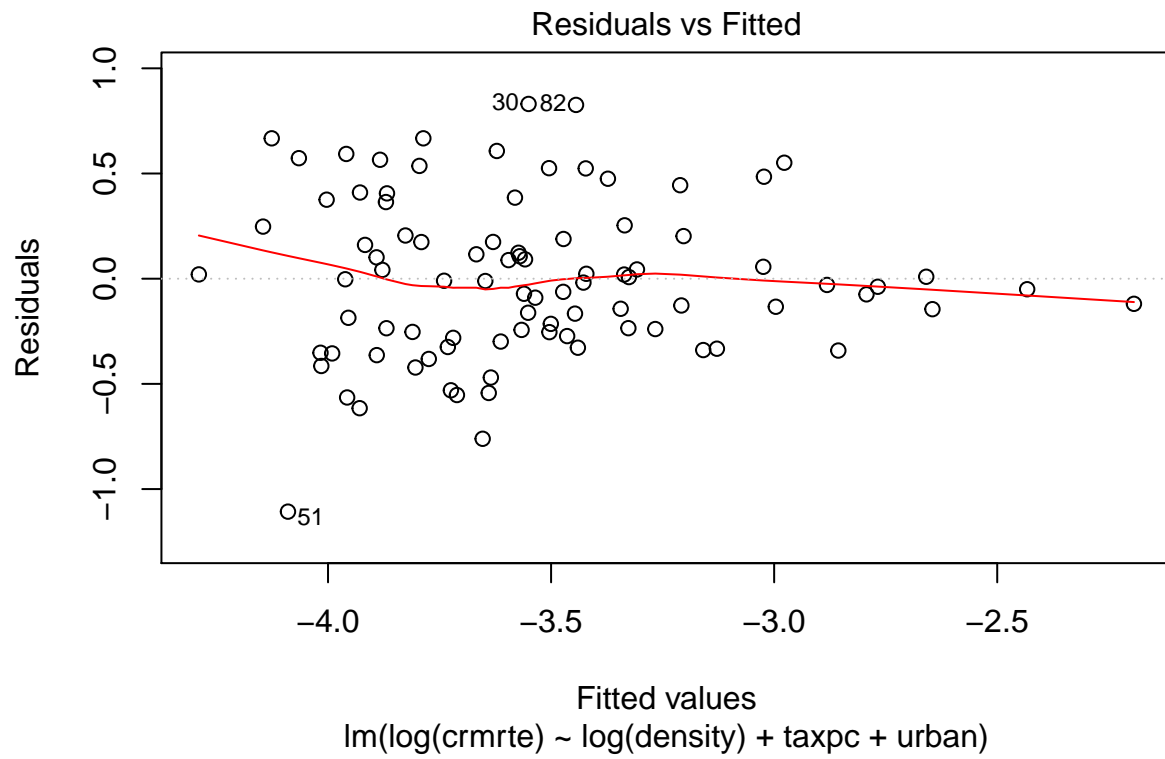
The two explanatory variables (log.density and taxpc) are not perfectly correlated and the VIFs are low (less than 4), so there is no perfect multicollinearity of the independent variables.

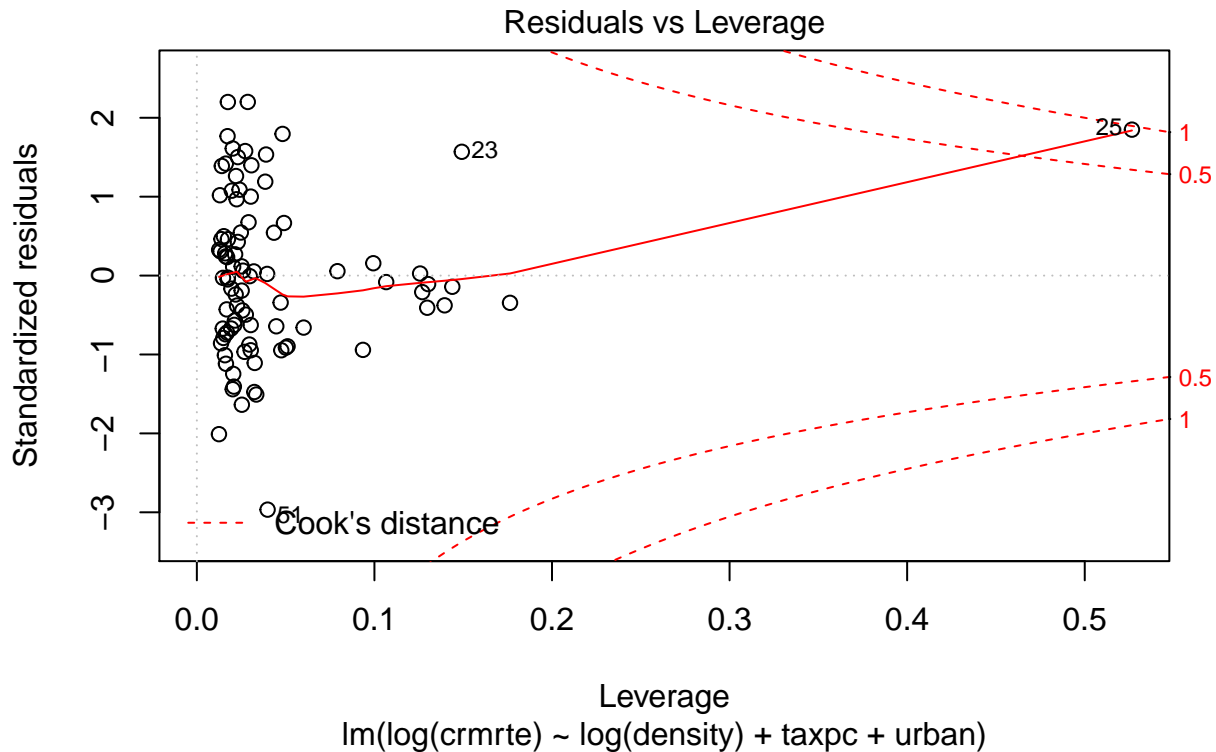
Is the assumption valid? *Yes*

Response: No response required.

CLM 4 - Zero-Conditional Mean

```
plot(m2, which=c(1, 5))
```





```
# exogenous check
cov(log(data$density), m2$residuals)
```

```
## [1] -8.492333e-18
```

```
cov(data$taxpc, m2$residuals)
```

```
## [1] -3.523097e-16
```

```
# TODO: figure out what to do with covariate
cov(data$urban, m2$residuals)
```

```
## [1] 9.501997e-19
```

The covariances of the two independent variables with the residuals are very close to zero indicating they are likely exogenous.

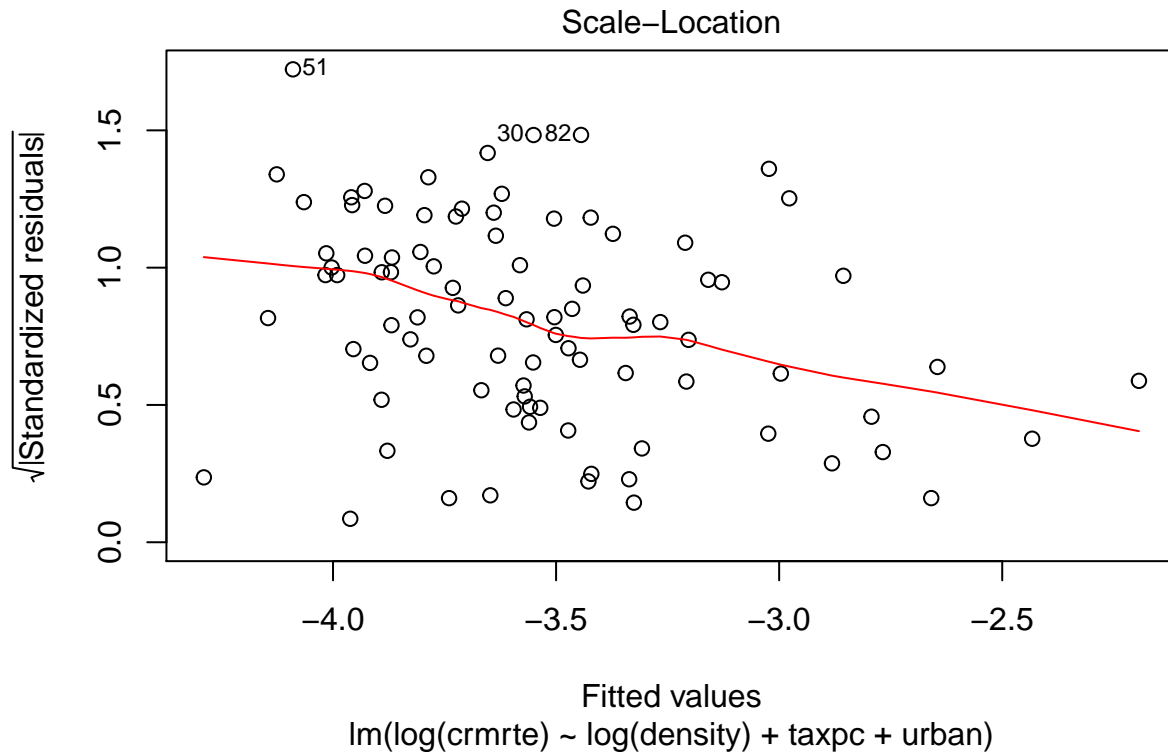
There is a data point with a large Cook's distance.

Is the assumption valid?

Response:

CLM 5 - Homoscedasticity

```
# spreadLevelPlot(m1)
plot(m2, which=3)
```

```
bptest(m2)
```

```
##
## studentized Breusch-Pagan test
##
## data: m2
## BP = 7.708, df = 3, p-value = 0.05245
not significant p-value for Breusch-Pagan test -> assumption of homoscedasticity met
```

```
ncvTest(m2)
```

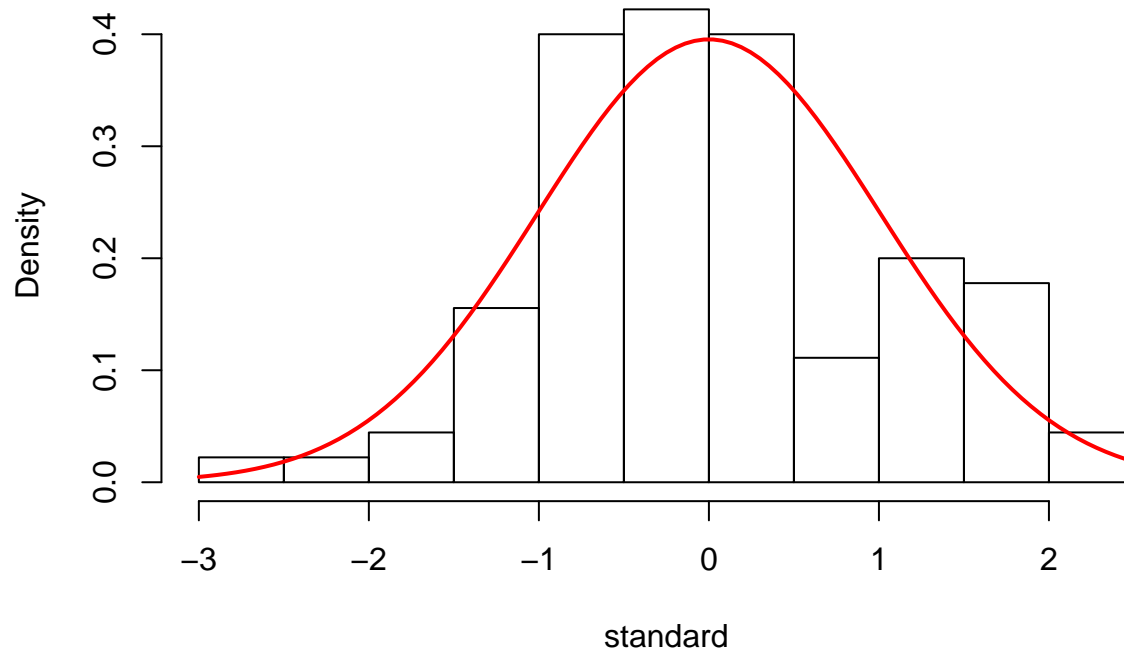
```
## Non-constant Variance Score Test
## Variance formula: ~ fitted.values
## Chisquare = 7.064728 Df = 1 p = 0.007861626
significant p-value for Score Test -> assumption of homoscedasticity met
Is the assumption valid? Yes
```

Response:

CLM 6 - Normality of residuals

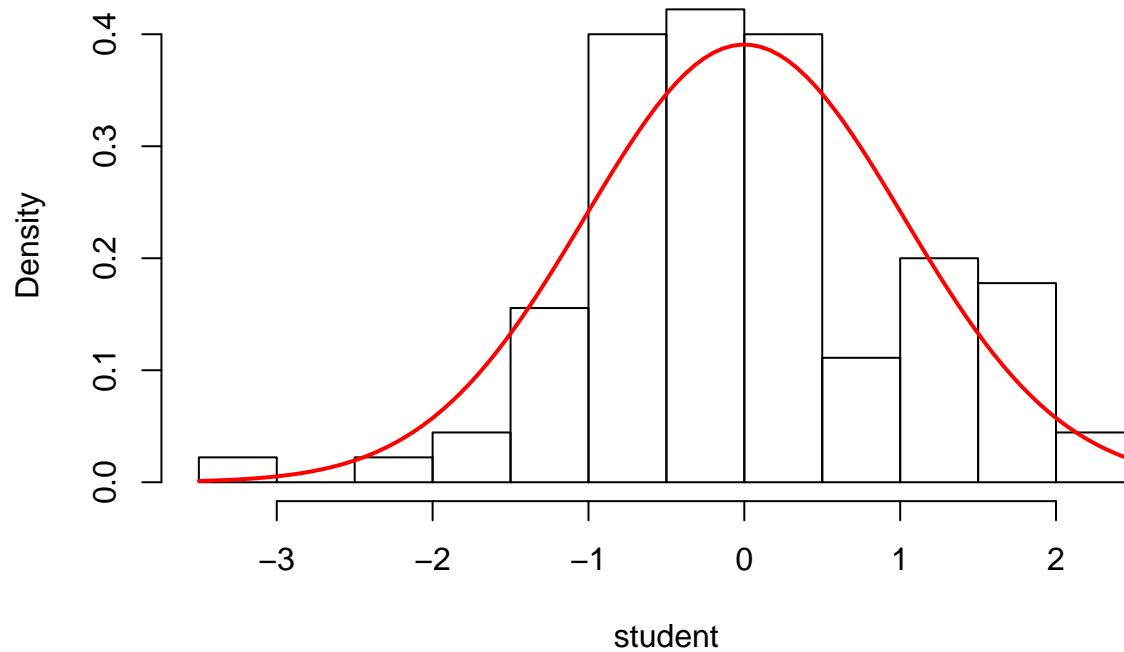
```
# normality of standard residuals
standard <- rstandard(m2)
hist(standard, main="Histogram of standard residuals", freq=FALSE)
curve(dnorm(x, mean=0, sd=sd(standard)), col="red", lwd=2, add=TRUE)
```

Histogram of standard resiudals



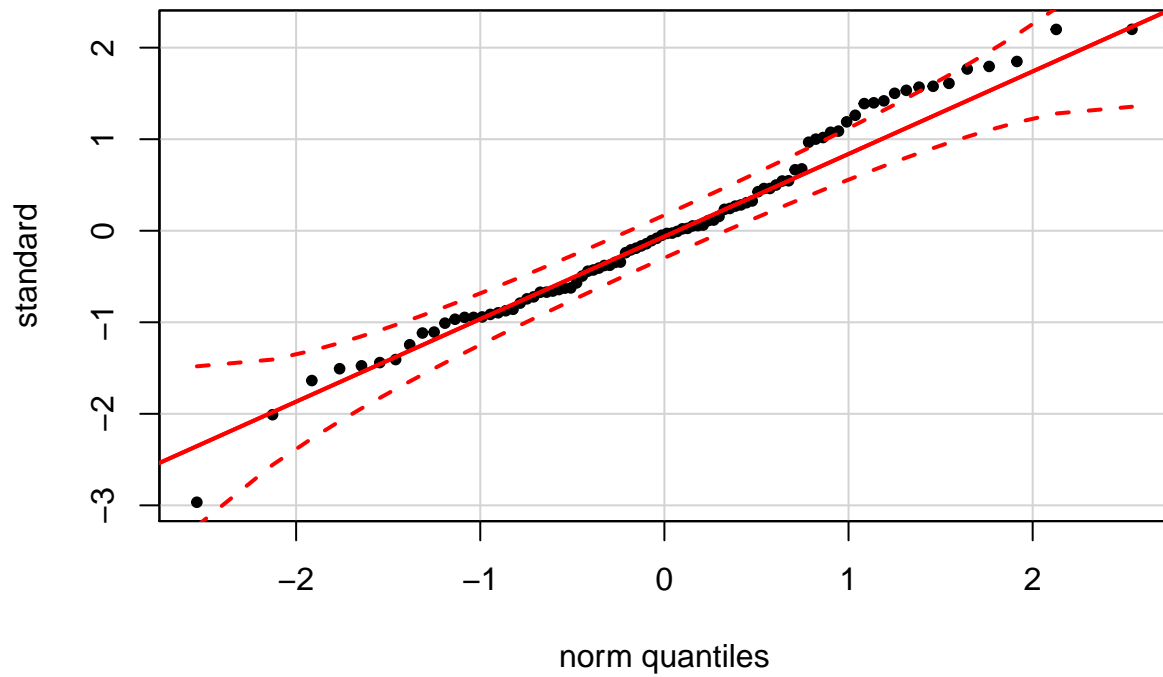
```
# normality of studentized residuals
student <- rstudent(m2)
hist(student, main="Histogram of studentized resiudals", freq=FALSE)
curve(dnorm(x, mean=0, sd=sd(student)), col="red", lwd=2, add=TRUE)
```

Histogram of studentized residuals



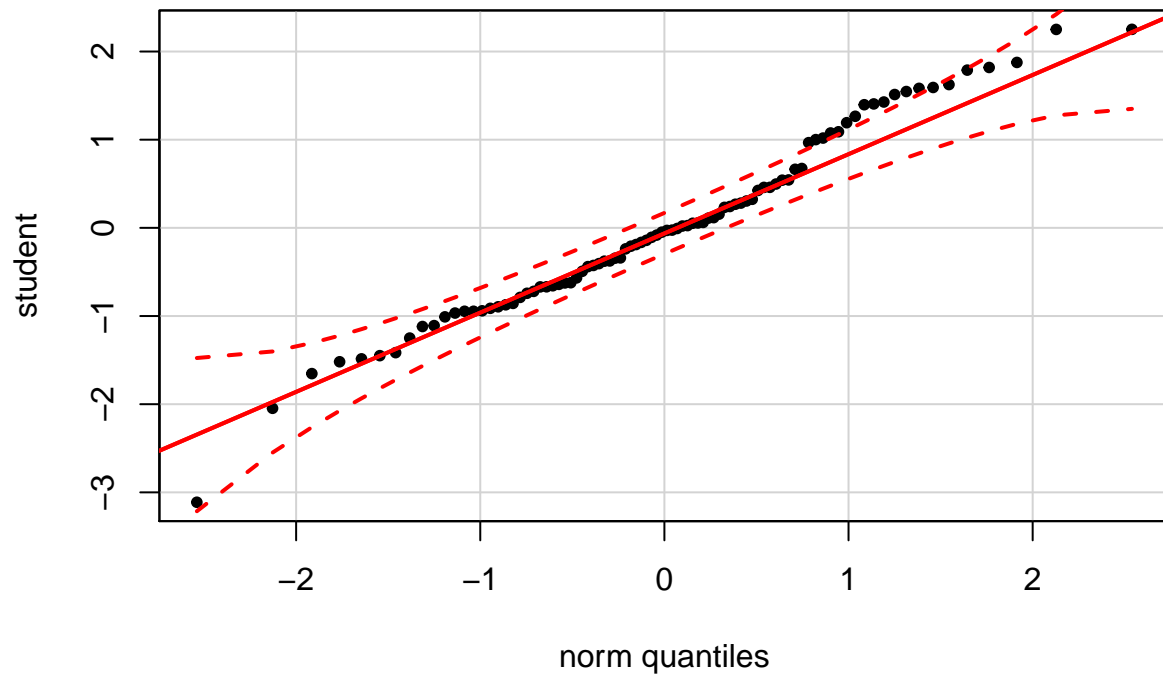
```
# QQ plot standard residuals  
qqPlot(standard, distribution="norm", pch=20, main="QQ-Plot standard residuals")  
qqline(standard, col="red", lwd=2)
```

QQ-Plot standard residuals



```
# QQ plot studentized residuals  
qqPlot(student, distribution="norm", pch=20, main="QQ-Plot studentized residuals")  
qqline(student, col="red", lwd=2)
```

QQ-Plot studentized residuals



The histograms in particular appear to be fairly normally distributed, while the Q-Q plot doesn't deviate significantly from normality either, so overall the residuals appear to be fairly normal. Notably the log transformation of views has helped to produce normal errors.

Is the assumption valid? *Yes*

Response: *No response required.*

TODO: * One model that includes the previous covariates, and most, if not all, other covariates. A key purpose of this model is to demonstrate the robustness of your results to model specification.

```
m3 <- lm(log(crmrte) ~ log(density) + taxpc + urban + central + west, data=data)
```

CLM 1 - A linear model

The model is specified such that the dependent variable is a linear function of the explanatory variables. Is the assumption valid? *Yes*

Response: *No response required.*

CLM 2 - Random Sampling

TODO

CLM 3 - Multicollinearity

```
# show correlation
data$log.density <- log(data$density)
cor(data.matrix(subset(data, select=c("density", "log.density", "taxpc", "urban"))))
```

```
##           density log.density    taxpc    urban
## density      1.0000000  0.8896747 0.3204737 0.8206825
## log.density  0.8896747  1.0000000 0.1841222 0.6490078
## taxpc        0.3204737  0.1841222 1.0000000 0.3457462
## urban        0.8206825  0.6490078 0.3457462 1.0000000
```

```
# show vif
vif(m2)
```

```
## log(density)    taxpc    urban
##      1.733261    1.139396    1.901850
```

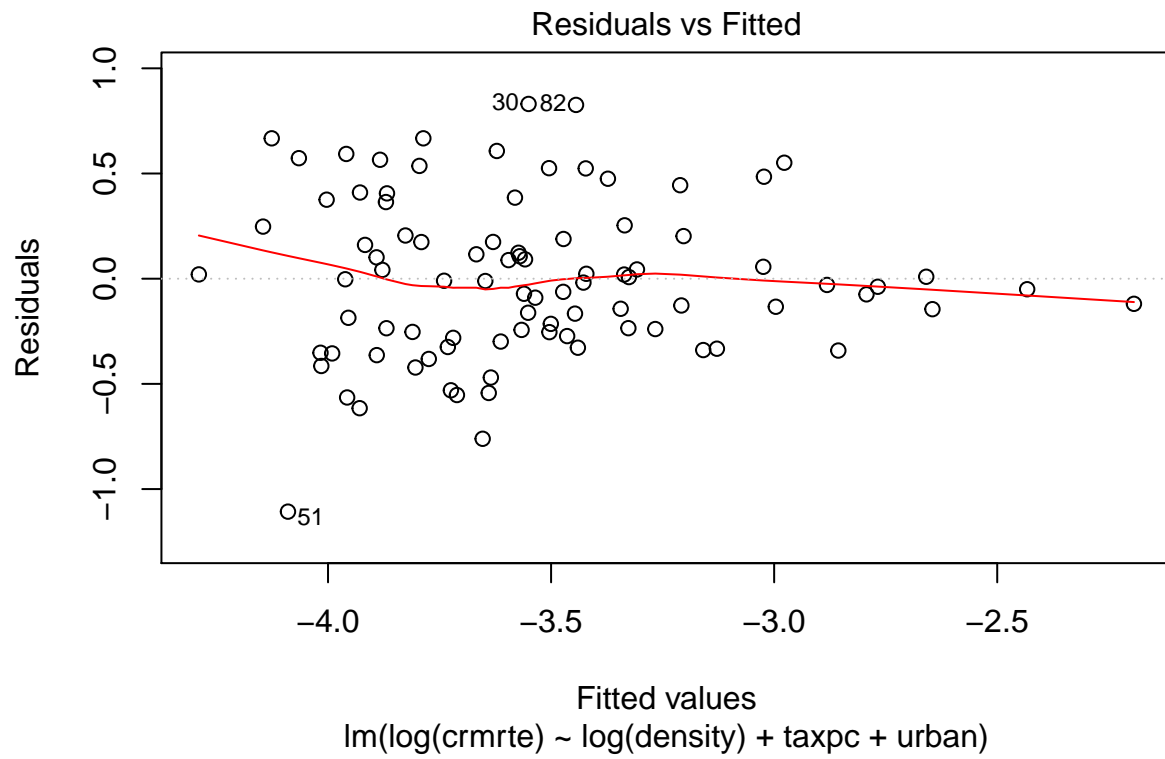
The two explanatory variables (log.density and taxpc) are not perfectly correlated and the VIFs are low (less than 4), so there is no perfect multicollinearity of the independent variables.

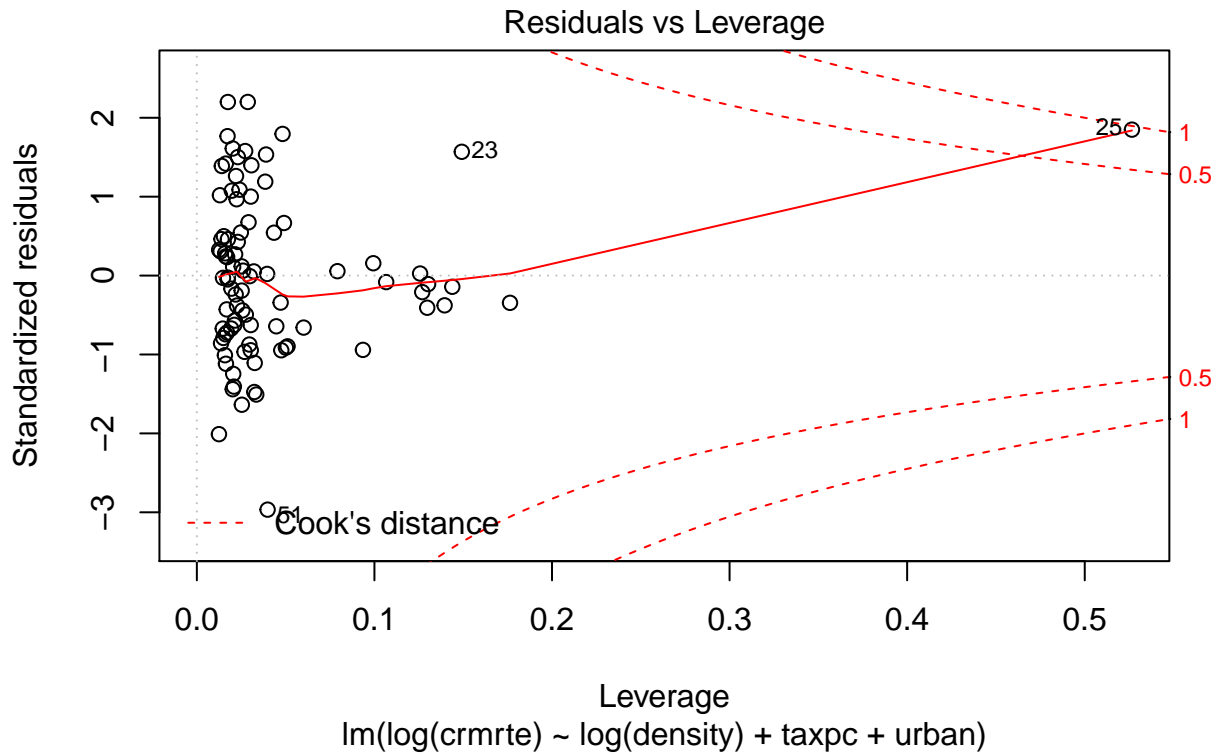
Is the assumption valid? *Yes*

Response: No response required.

CLM 4 - Zero-Conditional Mean

```
plot(m2, which=c(1, 5))
```





```
# exogenous check
cov(log(data$density), m2$residuals)
```

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## [1] 9.501997e-19
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The covariances of the two independent variables with the residuals are very close to zero indicating they are likely exogenous.

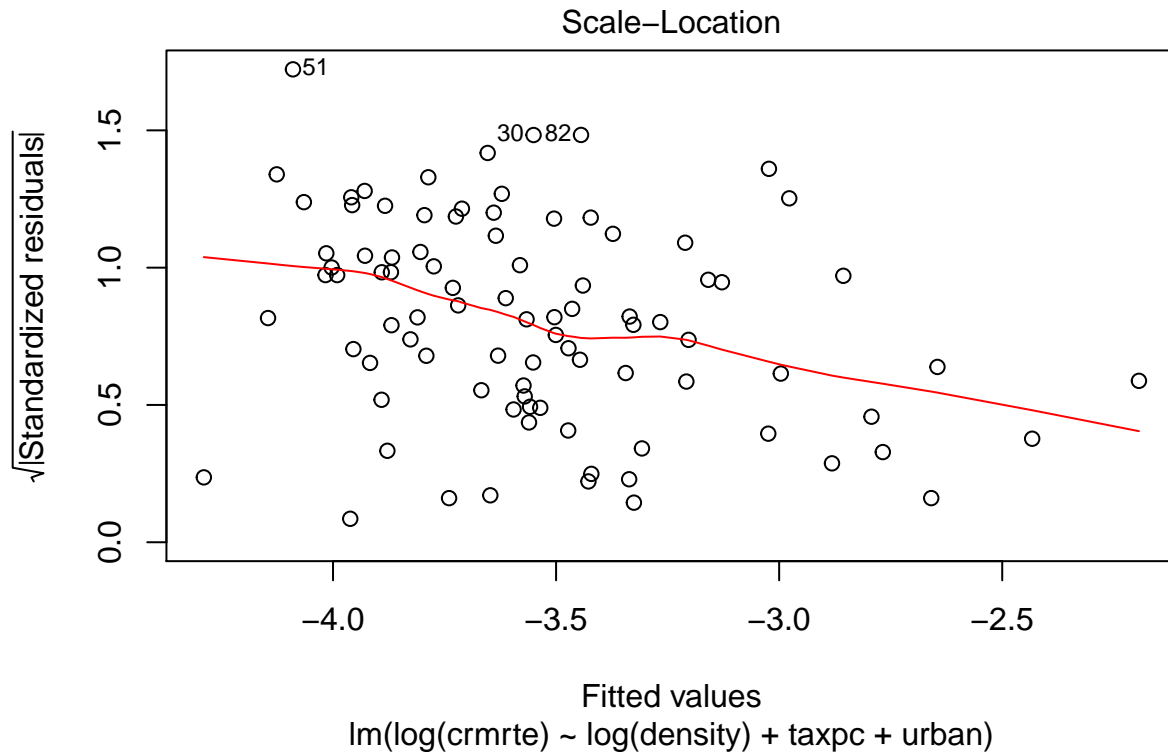
There is a data point with a large Cook's distance.

Is the assumption valid?

Response:

CLM 5 - Homoscedasticity

```
# spreadLevelPlot(m1)
plot(m2, which=3)
```

```
bptest(m2)
```

```
##
## studentized Breusch-Pagan test
##
## data: m2
## BP = 7.708, df = 3, p-value = 0.05245
not significant p-value for Breusch-Pagan test -> assumption of homoscedasticity met
```

```
ncvTest(m2)
```

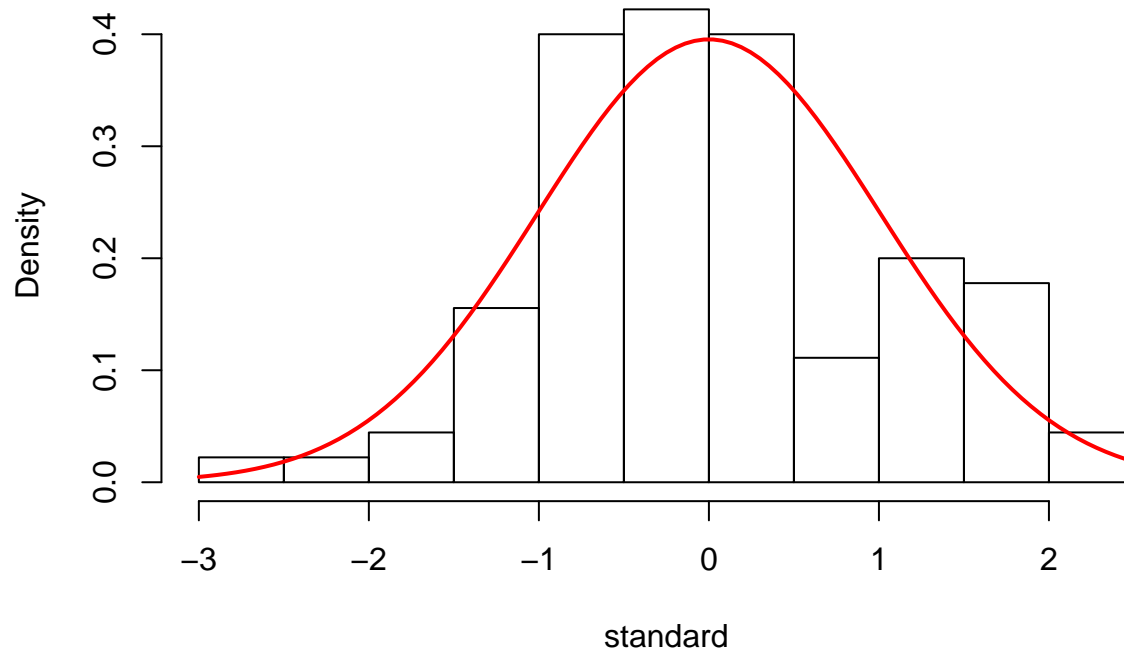
```
## Non-constant Variance Score Test
## Variance formula: ~ fitted.values
## Chisquare = 7.064728 Df = 1 p = 0.007861626
significant p-value for Score Test -> assumption of homoscedasticity met
Is the assumption valid? Yes
```

Response:

CLM 6 - Normality of residuals

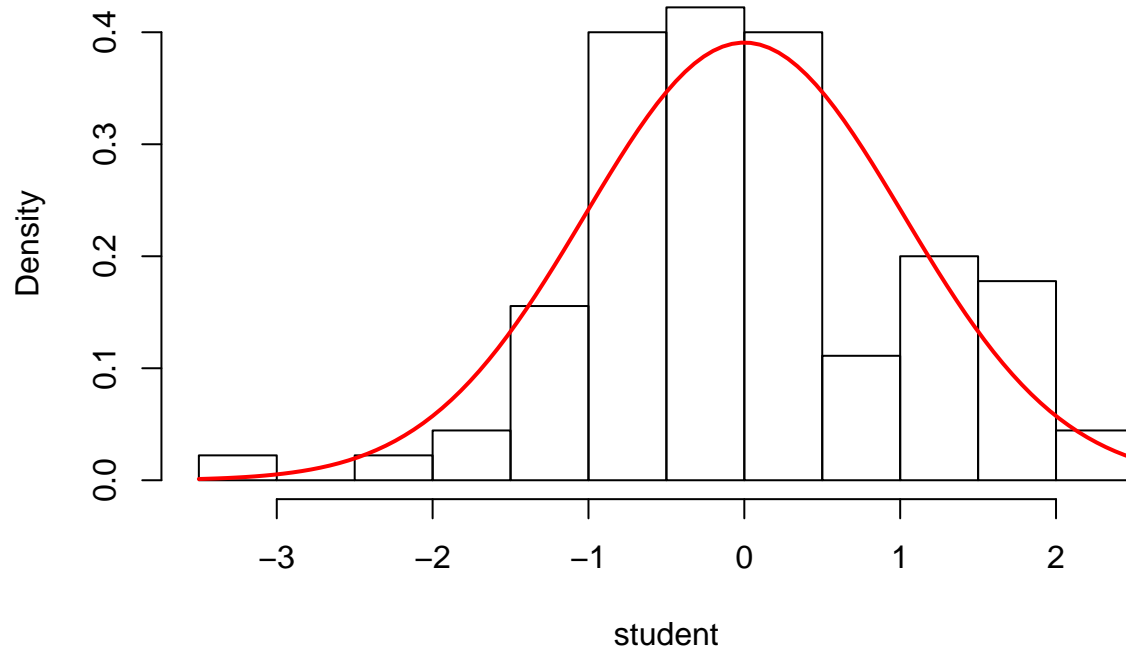
```
# normality of standard residuals
standard <- rstandard(m2)
hist(standard, main="Histogram of standard residuals", freq=FALSE)
curve(dnorm(x, mean=0, sd=sd(standard)), col="red", lwd=2, add=TRUE)
```

Histogram of standard resiudals



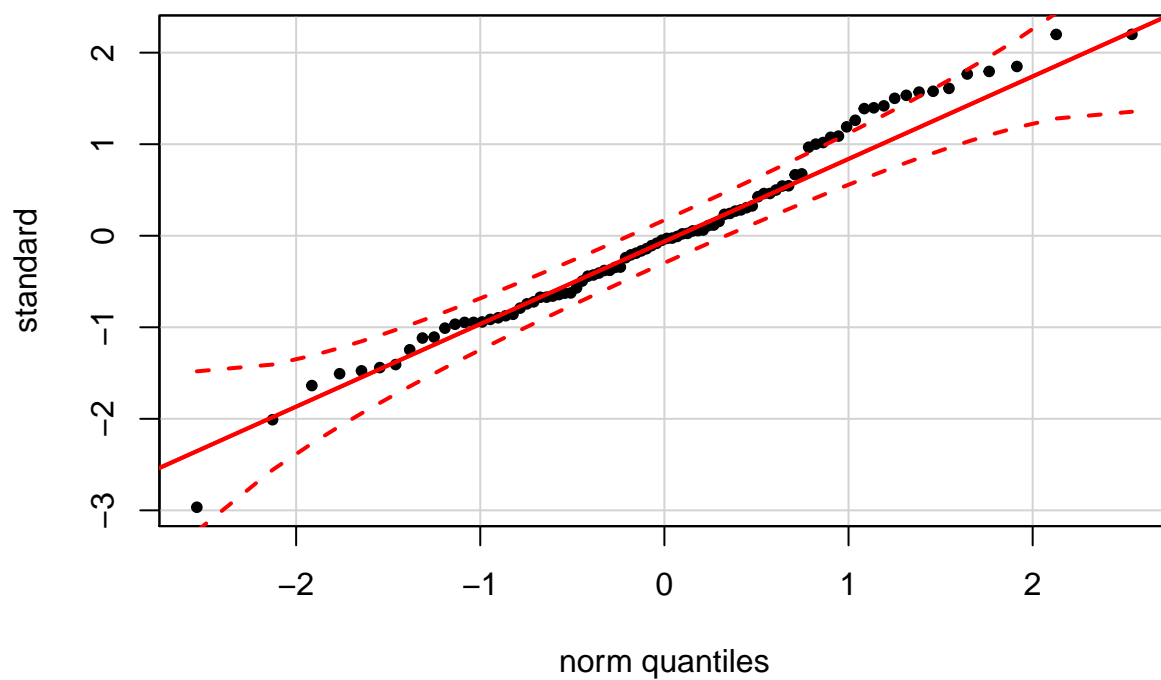
```
# normality of studentized residuals
student <- rstudent(m2)
hist(student, main="Histogram of studentized resiudals", freq=FALSE)
curve(dnorm(x, mean=0, sd=sd(student)), col="red", lwd=2, add=TRUE)
```

Histogram of studentized residuals



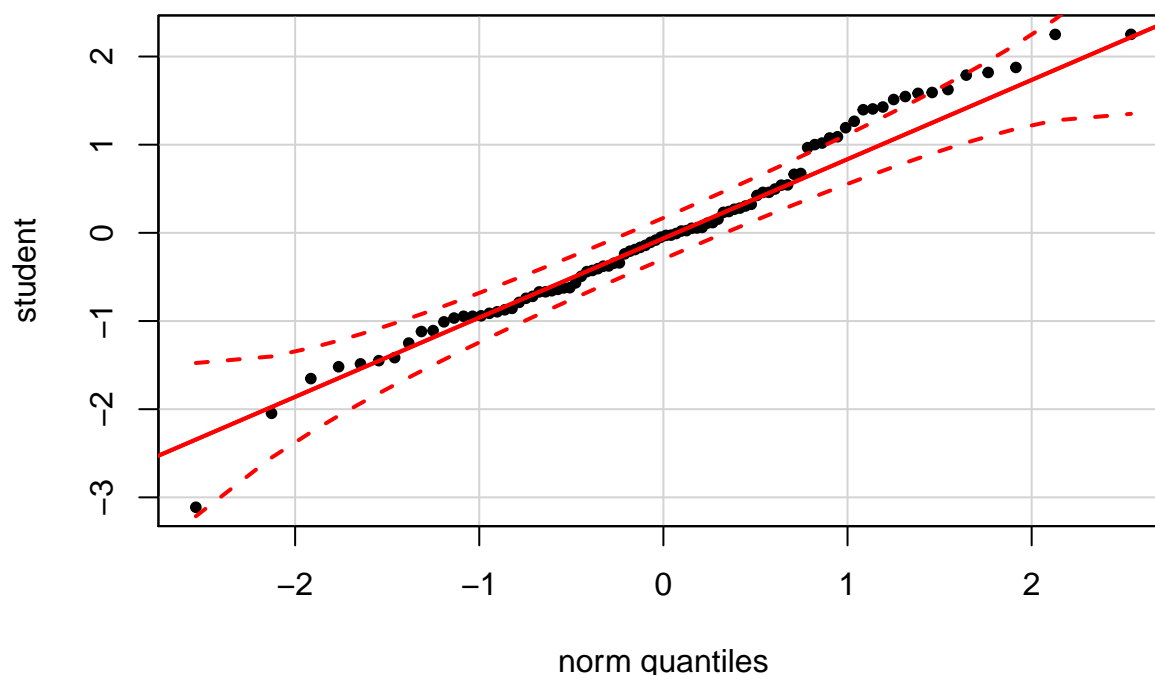
```
# QQ plot standard residuals  
qqPlot(standard, distribution="norm", pch=20, main="QQ-Plot standard residuals")  
qqline(standard, col="red", lwd=2)
```

QQ-Plot standard residuals



```
# QQ plot studentized residuals  
qqPlot(student, distribution="norm", pch=20, main="QQ-Plot studentized residuals")  
qqline(student, col="red", lwd=2)
```

QQ-Plot studentized residuals



The histograms in particular appear to be fairly normally distributed, while the Q-Q plot doesn't deviate significantly from normality either, so overall the residuals appear to be fairly normal. Notably the log transformation of views has helped to produce normal errors.

Is the assumption valid? *Yes*

Response: *No response required.*

Model Results

TODO: A well-formatted regression table summarizing your model results. Make sure that standard errors presented in this table are valid. Also, be sure to comment on both statistical and practical significance.

```
se.m1 <- coef(summary(m1))[, "Std. Error"]
se.m2 <- coef(summary(m2))[, "Std. Error"]
se.m3 <- coef(summary(m3))[, "Std. Error"]
stargazer(m1, m2, m3, se=list(se.m1, se.m2, se.m3), omit.stat=c("f", "ser"),
          star.cutoffs=c(0.05, 0.01, 0.001),
          type="latex", title="Linear Models Predicting Crime Rates")
```

```
##
## % Table created by stargazer v.5.2 by Marek Hlavac, Harvard University. E-mail: hlavac at fas.harvard.edu
## % Date and time: Wed, Dec 06, 2017 - 03:07:54
## \begin{table}[!htbp] \centering
##   \caption{Linear Models Predicting Crime Rates}
##   \label{}
```

```

## \begin{tabular}{@{\extracolsep{5pt}}lccc}
## \l[-1.8ex]\hline
## \hline \l[-1.8ex]
## & \multicolumn{3}{c}{\textit{Dependent variable:}} \\
## \cline{2-4}
## \l[-1.8ex] & \multicolumn{3}{c}{\log(crmrte)} \\
## \l[-1.8ex] & (1) & (2) & (3) \\
## \hline \l[-1.8ex]
## \log(density) & 0.455^{***} & 0.464^{***} & 0.488^{***} \\
## & (0.052) & (0.068) & (0.063) \\
## & & & \\
## taxpc & 0.010^{**} & 0.010^{**} & 0.007^{*} \\
## & (0.003) & (0.003) & (0.003) \\
## & & & \\
## urban & & -\$0.043 & -\$0.015 \\
## & & (0.195) & (0.168) \\
## & & & \\
## central & & & -\$0.306^{***} \\
## & & & (0.085) \\
## & & & \\
## west & & & -\$0.515^{***} \\
## & & & (0.091) \\
## & & & \\
## Constant & -\$3.931^{***} & -\$3.936^{***} & -\$3.589^{***} \\
## & (0.125) & (0.128) & (0.125) \\
## & & & \\
## \hline \l[-1.8ex]
## Observations & 90 & 90 & 90 \\
## R^{2} & 0.534 & 0.534 & 0.669 \\
## Adjusted R^{2} & 0.523 & 0.518 & 0.649 \\
## \hline
## \hline \l[-1.8ex]
## \textit{Note:} & \multicolumn{3}{r}{^{*}p<0.05; ^{**}p<0.01; ^{***}p<0.001} \\
## \end{tabular}
## \end{table}

```

Causality

TODO: A detailed discussion of causality. In particular, include a discussion of what variables are not included in your analysis and the likely direction of omitted variable bias. Highlight any coefficients you find that appear to have the wrong sign from a causal perspective, and explain why this is the case.

Conclusion

TODO: A brief conclusion with a few high-level takeaways.