

Analysis of Continuous Data project

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Protocol - Univariate part

Research question

During this research, we want to investigate how socio-economic disadvantage relates to violent crime rates. More specifically we want to explore the association between poverty and violent crime rates in the USA.

In his seminal work, Becker (1968) stated that the decision to commit crime is a rational choice where people weigh the benefits and costs against each other. It could then be argued that the incentive to commit crime is higher for people who have a lower income, as the benefits are larger for this group. Following this, we would then also expect that in communities with a higher poverty rate, there will also be higher crime rates. Depending on the results of our analysis, these results could be used to inform relevant policies. It would, for example, give another argument for the implementation of redistributive policies: if an effect is found, policymakers should take this reduction in violent crime into account, next to an economic benefit. Our analysis hopes to shed further light on this issue.

For the purpose of our research question, the following predictor variables have been selected:

- **PctPopUnderPov**: percentage of people under the poverty level (main predictor).
- **perCapInc**: per capita income. While similar to pctunderpoverty, this takes the whole income distribution into account and not just the lower end. If this average is lower, then we expect more crime to happen.
- **PctEmploy**: percentage of people 16 and over who are employed. We could argue that if more people are employed less people have an incentive to commit crime.
- **PctLess9thGrade**: percentage of people 25 and over with less than a 9th grade education. Education leads to a higher socio-economic standing, which would suggest that people have less reason to commit crime. We choose this variable for now, but as an alternative we could later use one of the following two variables if we would find them better suited as predictors: **PctNotHSGrad** (percentage of people 25 or over, that have not graduated highschool) or **PctBSorMore** (percentage of people 25 or over, with at least a bachelor's degree).
- **NumImmig**: total number of people known to be foreign born. Immigrants committing more crimes is a commonly used right-wing argument against migration, and relevant as immigrants are often from a 'lower' socio-economic background.
- **racepctblac**: percentage of population that is african american. It is a common right wing argument as well that black people commit crime, because they are from a 'lower' socio-economic background.
- **agePct12t29**: percentage of population that is 12-29 in age. We include this because young people have had less time to build up their socio-economic status, as well as their brain being less developed, and might thus commit more crime.

Design of the study

Descriptive analysis

To get a first impression of the data, a descriptive analysis will be performed for the candidate predictor variables (all continuous). The datasets are checked for missing values. The most common univariate statistics are calculated: the mean, the standard deviation, the minimum, the first quartile, the median, the third quartile and the maximum.

The distributions of the variables are visualized by boxplots, QQ plots and histograms. Outliers are identified using Tukey's 1.5 x IQR rule. For the univariate descriptive statistics also the population size of the communities is considered. The population size can influence the reliability of the data points: small communities can have a higher probability to have more extreme values of the predictor and response variables by the fact that the denominator in the response variable (total number of violent crimes per 100K population) is smaller. In the regression phase this will be used to investigate the outlier values.

To find what the relationship is between the main predictor variable and the potential extra predictor variables, scatter plots with smoothers are made for the bivariate relationships and correlations are checked.

Linear regression

Before performing linear regression and building models, the dataset is split into a training set (80% of the data) and a test set (20% of the data).

To investigate the association between the main predictor variable and the response variable, a linear regression is fitted and the output is evaluated. The various statistics are calculated and discussed: estimate regression coefficients, the F-statistics (/t-statistics), the R squared, the MSE, the p-value, the confidence interval and standard error of the slope. We first present the general formula here, before we fill in the specific variables.

$$Y_i = \beta_0 + \beta_1 X_i + \epsilon_i$$

$$ViolentCrimesPerPop_i = \beta_0 + \beta_1 pctpopUnderPov_i + \epsilon_i$$

Confidence intervals are constructed. Based on this, outliers can be identified. Subsequently, the outliers are further evaluated, e.g. are outliers linked to communities with a small population size.

Assumption checks

For linear regression, multiple assumptions, such as linearity, independence of errors, homoscedasticity (constant variance of errors), and normality of errors, are made. During this research these assumptions have to be checked by: Plotting residuals vs. fitted values for the linearity and independence of errors, squared residuals vs. fitted values for homoscedasticity checks, normality checks by qq-plot of the residuals. To also take leverage into account, the studentized residuals will be plotted.

Protocol - Multivariate part

Model building

Forward stepwise regression Evaluation of adjusted R-squared, AIC, SBC Partial regression plots? In which functional form we let a variable enter the model?

Model fit and outliers

PRESS, studentized residual plots (transformations needed?), bijv. QQ-plots to predicted value of y/log(y),
Also DFFITS, Cook's Distance, DFBETAS -> welke outliers hebben een grote invloed? Deleted residuals?

Interpretation of the parameters

Table 1: Project Schedule Overview

Deadline	Subject	Final_responsibility
3/11	Data extraction	Thomas
10/11	Descriptive analyses	Ilja
17/11	Model building	Bart
24/11	Model interpretation	Lieselot
24/11	Prediction with linear model	Ilja
1/12	Statistical discussion linear model	Bart
1/12	Fitting GLM	Lieselot
1/12	Fitting the final model	Thomas
8/12	Prediction with GLM	Ilja
8/12	Statistical discussion GLM	Thomas
8/12	Final conclusion and discussion	Lieselot

Data extraction:

```

library(readr)
library(dplyr)
library(stringr)

# The .arff header is usually inside the .names file:
url <- "https://archive.ics.uci.edu/ml/machine-learning-databases/communities/communities.names"

# Read the text lines
lines <- read_lines(url)
start <- grep("Additional Variable Information", lines, ignore.case = TRUE)
end   <- grep("Summary Statistics:", lines, ignore.case = TRUE)

var_lines <- lines[str_starts(str_trim(lines), "--")]
var_names <- str_extract(var_lines, "(?<-- )[^\n]+")
var_names <- c("communityname", setdiff(var_names, "communityname"))

variable_names <- list("communityname", "state", "countyCode", "communityCode", "fold", "population", "households", "racePctBlack", "racePctWhite", "racePctAsian", "racePctHisp", "agePct12t21", "agePct65up", "numUrban", "pctUrban", "medIncome", "pctWWage", "pctWFarmSelf", "pctWPubAsst", "pctWRetire", "medFamInc", "perCapInc", "whitePerCap", "blackPerCap", "OtherPerCap", "HisPctPerCap", "NumUnderPov", "PctPopUnderPov", "PctLess9thGrade", "PctUnemployed", "PctEmploy", "PctEmplManu", "PctEmplProfServ", "PctOccupManu", "MalePctNevMarr", "FemalePctDiv", "TotalPctDiv", "PersPerFam", "PctFam2Par", "PctTeen2Par", "PctWorkMomYoungKids", "PctWorkMom", "NumKidsBornNeverMar", "PctKidsBornRecent", "PctImmigRecent", "PctImmigRec5", "PctImmigRec8", "PctImmigRec10", "PctRecentImmig")
  
```

```

"PctRecImmig10", "PctSpeakEnglOnly", "PctNotSpeakEnglWell", "PctLargHouseFam", "PctOwnOccup", "PctPerOwnOccHous", "PersPerRentOccHous", "PctPersOwnOccup", "PctPersDenseHous", "HousVacant", "PctHousOccup", "PctHousOwnOcc", "PctVacantBoarded", "PctVacMore6M", "PctHousNoPhone", "PctWOFullPlumb", "OwnOccLowQuart", "OwnOccMedVal", "OwnOccHiQ", "RentMedian", "RentHighQ", "RentQrange", "MedRent", "MedRentPctHousInc", "MedOwnOccup", "NumInShelters", "NumStreet", "PctForeignBorn", "PctBornSameState", "PctSameHouse", "PctSameState85", "LemasSwornFT", "LemasSwFTPerPop", "LemasSwFTFieldOps", "LemasPctOffic", "LemasTotReqPerPop", "PolicReqPerOffic", "PolicPerPop", "RacialMatchCommPol", "PctPolicHisp", "PctPolicAsian", "PctPolicMinor", "OfficAssgnDrugUnits", "NumKinds", "LandArea", "PopDens", "PctUsePubTrans", "PolicCars", "PolicOperBudg", "LemasPctOffic", "LemasPctOfficDrugUn", "PolicBudgPerPop", "murders", "murdPerPop", "rapes", "rape", "assaults", "assaultPerPop", "burglaries", "burglPerPop", "larcenies", "larcPerPop", "arsons", "arsonsPerPop", "ViolentCrimesPerPop", "nonViolPerPop")

```

```

library(data.table)
violent_crimes_table <- fread("curl https://archive.ics.uci.edu/static/public/211/communities+and+crimes.csv")

```

```

colnames(violent_crimes_table) <- unlist(variable_names)

```

```

crimes_table_subset <- violent_crimes_table %>%
  select(communityname, state, countyCode, communityCode, fold, population,
         PctPopUnderPov, perCapInc, PctEmploy, PctLess9thGrade, PctNotHSGrad, PctBSorMore,
         NumImmig, racepctblack, agePct12t29, ViolentCrimesPerPop
  )

```

Design (tekst gekopieerd van website)

The source datasets needed to be combined via programming. Many variables are included so that algorithms that select or learn weights for attributes could be tested. However, clearly unrelated attributes were not included; attributes were picked if there was any plausible connection to crime (N=125), plus the crime variables which are potential dependent variables. The variables included in the dataset involve the community, such as the percent of the population considered urban, and the median family income, and involving law enforcement, such as per capita number of police officers, and percent of officers assigned to drug units. The crime attributes (N=18) that could be predicted are the 8 crimes considered ‘Index Crimes’ by the FBI)(Murders, Rape, Robbery, . . .), per capita (actually per 100,000 population) versions of each, and Per Capita Violent Crimes and Per Capita Nonviolent Crimes).

A limitation was that the LEMAS survey was of the police departments with at least 100 officers, plus a random sample of smaller departments. For our purposes, communities not found in both census and crime datasets were omitted. Many communities are missing LEMAS data.

The per capita crimes variables were calculated using population values included in the 1995 FBI data (which differ from the 1990 Census values).

The per capita violent crimes variable was calculated using population and the sum of crime variables considered violent crimes in the United States: murder, rape, robbery, and assault. There was apparently some controversy in some states concerning the counting of rapes. These resulted in missing values for rape, which resulted in missing values for per capita violent crime. Many of these omitted communities were from the midwestern USA (Minnesota, Illinois, and Michigan have many of these).

The per capita nonviolent crime variable was calculated using the sum of crime variables considered non-violent crimes in the United States: burglaries, larcenies, auto thefts and arsons. (There are many other types of crimes, these only include FBI ‘Index Crimes’)

Some further pre-processing of the dataset must be done. Choose the desirable dependent variable from among the 18 possible. It would not be interesting or appropriate to predict total crime (e.g. violent crime) while including subtotals (e.g. murders) as independent variables. There are also identifying variables (community name, county code, community code) that are not predictive, and would get in the way of some algorithms. Weka's Unsupervised Attribute Remove Filter can be used to remove unwanted attributes.

The FBI notes that use of this data to evaluate communities is over-simplistic, as many relevant factors are not included. For one example, communities with large numbers of visitors will have higher per capita crime (measured by residents) than communities with fewer visitors, other things being equal.

Data preparation and descriptive analysis

Since the outcome variable *ViolentCrimesPerPop* (total number of violent crimes per 100K population) is expressed relative to the population size, the variable *NumImmig* is converted (by dividing it by the population size and multiplying by 100%).

```
crimes_table_subset$ViolentCrimesPerPop <- as.numeric(crimes_table_subset$ViolentCrimesPerPop)
crimes_table_subset$PctImmig <- crimes_table_subset$NumImmig / crimes_table_subset$population * 100
crimes_table_subset = crimes_table_subset[, -c('NumImmig', 'fold')]

sjlabelled::set_label(crimes_table_subset) <- c("communityname", "state", "countyCode", "communityCode"
or over, that have not graduated highschool (%)", "percentage of people 25 or over, with
at least a bachelor's degree (%)", "percentage of population that is african american (%)", "percentage
```

It is examined how many NA values are present in the database.

```
crimes_table_subset %>%
  pivot_longer(cols = where(is.numeric), names_to = "variable", values_to = "value") %>%
  group_by(variable) %>%
  summarise(
    NAs = sum(is.na(value))
  )

## # A tibble: 11 x 2
##   variable           NAs
##   <chr>              <int>
## 1 PctBSorMore        0
## 2 PctEmploy          0
## 3 PctImmig           0
## 4 PctLess9thGrade    0
## 5 PctNotHSGrad       0
## 6 PctPopUnderPov     0
## 7 ViolentCrimesPerPop 221
## 8 agePct12t29        0
## 9 perCapInc          0
## 10 population         0
## 11 racepctblack      0

na_subset <- crimes_table_subset %>%
  filter(is.na(ViolentCrimesPerPop)
  )
na_subset <- na_subset[, -'ViolentCrimesPerPop']
na_subset
```

```

##      communityname state countyCode communityCode population PctPopUnderPov
##                <char> <char>     <char>       <char>    <int>      <num>
## 1:   Bemidjicity    MN        7      5068    11245    29.99
## 2:   NewUlmcity     MN       15     46042    13132     6.84
## 3:  Maplewoodcity   MN      123     40382    30954     6.22
## 4:  Plymouthcity    MN       53     51730    50889     3.36
## 5:  Pontiaccity     MI      125     65440    71166    26.67
## ---
## 217:  Bristoltown   CT        3      8490    60640     4.35
## 218: Wilmettevillage IL       ?       ?    26690     2.14
## 219: EastLansingcity MI       65     24120    50677    33.77
## 220: CrystalLakecity IL       ?       ?    24512     2.15
## 221: Burtoncity     MI       49     12060    27617    14.28
##      perCapInc PctEmploy PctLess9thGrade PctNotHSGrad PctBSorMore racepctblack
##                <int>    <num>       <num>       <num>      <num>      <num>
## 1:     8483    52.44     12.15     23.06    25.28    0.53
## 2:    11907    65.62     16.28     25.41    15.31    0.06
## 3:    16459    68.12      4.40     14.64    20.28    2.52
## 4:    21908    78.05      1.57      5.56    41.39    1.61
## 5:    9847     51.07     12.04     37.61     7.95    42.20
## ---
## 217:   16909    68.24     10.04     24.97    15.39    2.08
## 218:   38465    63.90      2.78      4.88    63.69    0.49
## 219:   11212    57.45      0.93      3.38    71.23    6.93
## 220:   17681    71.89      3.53     11.00    28.50    0.20
## 221:   12940    53.67      7.56     27.32     6.68    2.57
##      agePct12t29 PctImmig
##                <num>    <num>
## 1:     40.53  1.7429969
## 2:     25.03  0.9061834
## 3:     25.43  2.6232474
## 4:     26.94  2.6135314
## 5:     32.21  2.3058764
## ---
## 217:   27.32  7.0052770
## 218:   18.30 13.0760584
## 219:   67.80 10.6872940
## 220:   25.81  4.2142624
## 221:   27.14  1.8177210

```

The only variable for which NA values are found is the outcome variable *ViolentCrimesPerPop*. The rows where the outcome variable has an NA value are removed, as these rows are not useful for the regression. It can be noted that the variables *countyCode* and *communityCode* are also frequently unknown.

```

crimes_table_subset = na.omit(crimes_table_subset)
colSums(crimes_table_subset == "?", na.rm = TRUE)

```

	communityname	state	countyCode	communityCode
##	0	0	1174	1177
##	population	PctPopUnderPov	perCapInc	PctEmploy
##	0	0	0	0
##	PctLess9thGrade	PctNotHSGrad	PctBSorMore	racepctblack
##	0	0	0	0

```

##      agePct12t29 ViolentCrimesPerPop          PctImmig
##            0                      0

```

After removing NA values from the database univariate descriptives are calculated.

```
str(crimes_table_subset)
```

```

## Classes 'data.table' and 'data.frame': 1994 obs. of 15 variables:
##   $ communityname : chr "BerkeleyHeightstownship" "Marpletownship" "Tigardcity" "Gloversvilleci"
##   ..- attr(*, "label")= Named chr "communityname"
##   ... ..- attr(*, "names")= chr "communityname"
##   $ state         : chr "NJ" "PA" "OR" "NY" ...
##   ..- attr(*, "label")= Named chr "state"
##   ... ..- attr(*, "names")= chr "state"
##   $ countyCode    : chr "39" "45" "?" "35" ...
##   ..- attr(*, "label")= Named chr "countyCode"
##   ... ..- attr(*, "names")= chr "countyCode"
##   $ communityCode : chr "5320" "47616" "?" "29443" ...
##   ..- attr(*, "label")= Named chr "communityCode"
##   ... ..- attr(*, "names")= chr "communityCode"
##   $ population    : int 11980 23123 29344 16656 140494 28700 59459 74111 103590 31601 ...
##   ..- attr(*, "label")= Named chr "population"
##   ... ..- attr(*, "names")= chr "population"
##   $ PctPopUnderPov : num 1.96 3.98 4.75 17.23 17.78 ...
##   ..- attr(*, "label")= Named chr "people under the poverty level (%)"
##   ... ..- attr(*, "names")= chr "PctPopUnderPov"
##   $ perCapInc     : int 29711 20148 16946 10810 11878 18193 12161 13554 10195 12929 ...
##   ..- attr(*, "label")= Named chr "per capita income ($)"
##   ... ..- attr(*, "names")= chr "perCapInc"
##   $ PctEmploy     : num 64.5 62 69.8 54.7 59 ...
##   ..- attr(*, "label")= Named chr "percentage of people 16 and over who are employed (%)"
##   ... ..- attr(*, "names")= chr "PctEmploy"
##   $ PctLess9thGrade : num 5.81 5.61 2.8 11.05 8.76 ...
##   ..- attr(*, "label")= Named chr "percentage of people 25 and over with less than a 9th grade education"
##   ... ..- attr(*, "names")= chr "PctLess9thGrade"
##   $ PctNotHSGrad  : num 9.9 13.72 9.09 33.68 23.03 ...
##   ..- attr(*, "label")= Named chr "percentage of people 25\nor over, that have not graduated highschool"
##   ... ..- attr(*, "names")= chr "PctNotHSGrad"
##   $ PctBSorMore   : num 48.2 29.9 30.1 10.8 20.7 ...
##   ..- attr(*, "label")= Named chr "percentage of people 25 or over, with\nat least a bachelor's degree"
##   ... ..- attr(*, "names")= chr "PctBSorMore"
##   $ racepctblack  : num 1.37 0.8 0.74 1.7 2.51 ...
##   ..- attr(*, "label")= Named chr "percentage of population that is african american (%)"
##   ... ..- attr(*, "names")= chr "racepctblack"
##   $ agePct12t29   : num 21.4 21.3 25.9 25.2 32.9 ...
##   ..- attr(*, "label")= Named chr "percentage of population that is 12-29 in age (%)"
##   ... ..- attr(*, "names")= chr "agePct12t29"
##   $ ViolentCrimesPerPop: num 41 128 219 307 443 ...
##   ..- attr(*, "label")= Named chr "total number of violent crimes per 100K population"
##   ... ..- attr(*, "names")= chr "ViolentCrimesPerPop"
##   $ PctImmig      : num 10.66 8.3 5 2.04 1.49 ...
##   ..- attr(*, "label")= Named chr "percentage of immigrants (%)"
##   ... ..- attr(*, "names")= chr "PctImmig"
##   - attr(*, ".internal.selfref")=<externalptr>

```

```

summary(crimes_table_subset)

##   communityname      state    countyCode  communityCode
##   Length:1994      Length:1994    Length:1994    Length:1994
##   Class :character Class :character  Class :character  Class :character
##   Mode  :character  Mode :character   Mode :character  Mode :character
##
## 
## 
##   population  PctPopUnderPov  perCapInc  PctEmploy
##   Min.    : 10005  Min.    : 0.640  Min.    : 5237  Min.    :24.82
##   1st Qu.: 14359  1st Qu.: 4.692  1st Qu.:11548  1st Qu.:56.35
##   Median  : 22681  Median  : 9.650  Median  :13977  Median  :62.27
##   Mean    : 52251  Mean    :11.796  Mean    :15522  Mean    :61.78
##   3rd Qu.: 43154  3rd Qu.:17.078  3rd Qu.:17775  3rd Qu.:67.50
##   Max.    :7322564  Max.    :48.820  Max.    :63302  Max.    :84.67
##   PctLess9thGrade  PctNotHSGrad  PctBSorMore  racepctblack
##   Min.    : 0.200  Min.    : 2.09  Min.    : 1.63  Min.    : 0.00
##   1st Qu.: 4.770  1st Qu.:14.20  1st Qu.:14.09  1st Qu.: 0.94
##   Median  : 7.920  Median  :21.66  Median  :19.62  Median  : 3.15
##   Mean    : 9.444  Mean    :22.70  Mean    :22.99  Mean    : 9.51
##   3rd Qu.:12.245  3rd Qu.:29.66  3rd Qu.:28.93  3rd Qu.:11.96
##   Max.    :49.890  Max.    :73.66  Max.    :73.63  Max.    :96.67
##   agePct12t29  ViolentCrimesPerPop  PctImmig
##   Min.    : 9.38  Min.    : 0.0  Min.    : 0.1778
##   1st Qu.:24.38  1st Qu.:161.7  1st Qu.: 2.0753
##   Median  :26.77  Median  :374.1  Median  : 4.4935
##   Mean    :27.62  Mean    :589.1  Mean    : 7.6062
##   3rd Qu.:29.18  3rd Qu.:794.4  3rd Qu.: 9.5848
##   Max.    :70.51  Max.    :4877.1  Max.    :60.4013

crimes_table_subset %>%
  pivot_longer(cols = where(is.numeric), names_to = "variable", values_to = "value") %>%
  group_by(variable) %>%
  summarise(
    min = min(value, na.rm = TRUE),
    q25 = quantile(value, 0.25, na.rm = TRUE),
    mean = mean(value, na.rm = TRUE),
    sd = sd(value, na.rm = TRUE),
    q75 = quantile(value, 0.75, na.rm = TRUE),
    max = max(value, na.rm = TRUE),
    n = n(),
    NAs = sum(is.na(value))
  )

## # A tibble: 11 x 9
##   variable      min     q25     mean     sd     q75     max     n   NAs
##   <chr>      <dbl>   <dbl>   <dbl>   <dbl>   <dbl>   <dbl> <int> <int>
## 1 PctBSorMore  1.63  1.41e+1  2.30e1  1.25e1  2.89e1  7.36e1  1994  0
## 2 PctEmploy    24.8   5.64e+1  6.18e1  8.11e0  6.75e1  8.47e1  1994  0
## 3 PctImmig     0.178  2.08e+0  7.61e0  8.70e0  9.58e0  6.04e1  1994  0
## 4 PctLess9thGrade  0.2   4.77e+0  9.44e0  6.84e0  1.22e1  4.99e1  1994  0
## 5 PctNotHSGrad  2.09  1.42e+1  2.27e1  1.11e1  2.97e1  7.37e1  1994  0

```

```

##   6 PctPopUnderPov      0.64  4.69e+0 1.18e1 8.51e0 1.71e1 4.88e1 1994      0
##   7 ViolentCrimesPerPop     0    1.62e+2 5.89e2 6.15e2 7.94e2 4.88e3 1994      0
##   8 agePct12t29      9.38  2.44e+1 2.76e1 6.15e0 2.92e1 7.05e1 1994      0
##   9 perCapInc        5237  1.15e+4 1.55e4 6.23e3 1.78e4 6.33e4 1994      0
##  10 population       10005  1.44e+4 5.23e4 2.02e5 4.32e4 7.32e6 1994      0
##  11 racepctblack      0    9.4 e-1 9.51e0 1.41e1 1.20e1 9.67e1 1994      0

na_subset %>%
  pivot_longer(cols = where(is.numeric), names_to = "variable", values_to = "value") %>%
  group_by(variable) %>%
  summarise(
    min = min(value, na.rm = TRUE),
    q25 = quantile(value, 0.25, na.rm = TRUE),
    mean = mean(value, na.rm = TRUE),
    sd = sd(value, na.rm = TRUE),
    q75 = quantile(value, 0.75, na.rm = TRUE),
    max = max(value, na.rm = TRUE),
    n = n(),
    NAs = sum(is.na(value))
  )

## # A tibble: 10 x 9
##   variable      min     q25     mean      sd     q75     max     n     NAs
##   <chr>     <dbl>   <dbl>   <dbl>   <dbl>   <dbl>   <dbl> <int> <int>
## 1 PctBSorMore  3.27    14.2    23.6  1.42e1  3.09e1  7.92e1  221    0
## 2 PctEmploy    33.7    57.4    64.2  9.71e0  7.04e1  8.45e1  221    0
## 3 PctImmig     0.445    1.99    4.94  4.62e0  6.27e0  2.84e1  221    0
## 4 PctLess9thGrade 0.41    3.54    6.86  4.12e0  9.64e0  2.12e1  221    0
## 5 PctNotHSGrad 1.46    11.1    18.7  9.63e0  2.54e1  5.26e1  221    0
## 6 PctPopUnderPov 1.25    3.49    10.0  9.25e0  1.33e1  5.8 e1  221    0
## 7 agePct12t29   17.4    25.0    27.9  6.48e0  2.94e1  6.78e1  221    0
## 8 perCapInc    5622   12205  16342.  6.72e3  1.81e4  6.25e4  221    0
## 9 population    10066  14903  60937.  2.26e5  4.18e4  2.78e6  221    0
## 10 racepctblack 0.03    0.49    7.76  1.54e1  6.4 e0  9.28e1  221    0

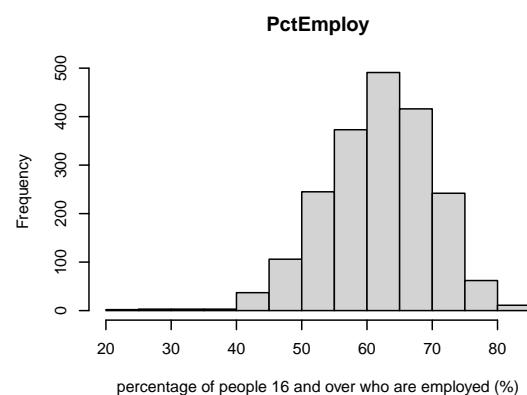
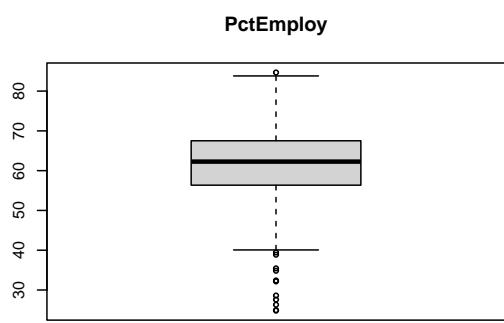
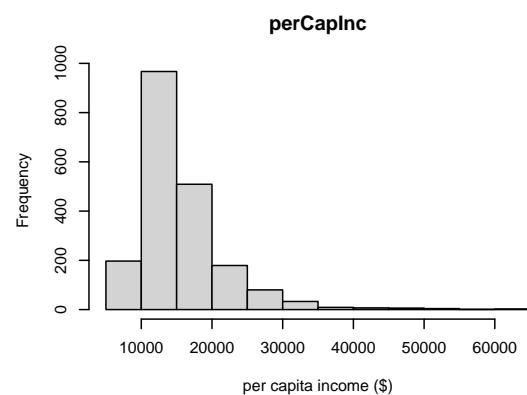
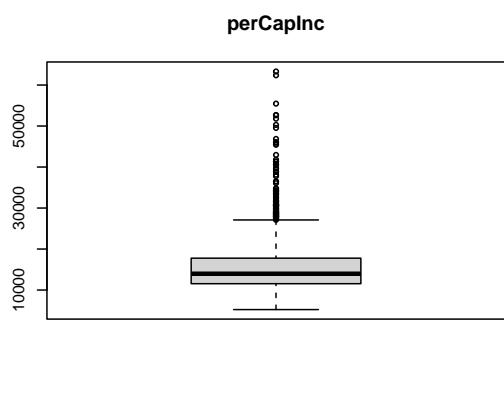
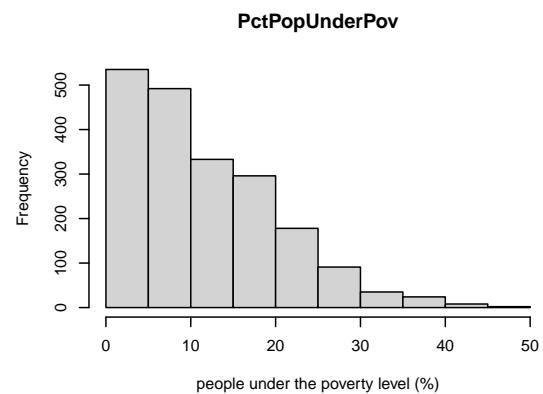
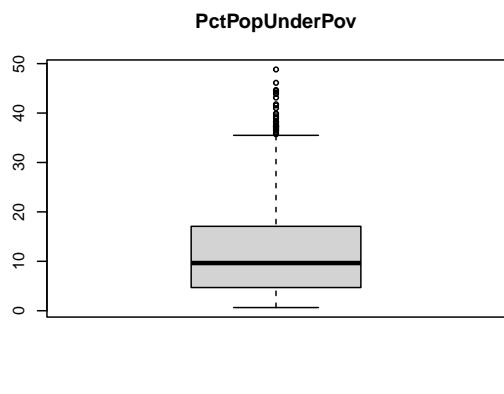
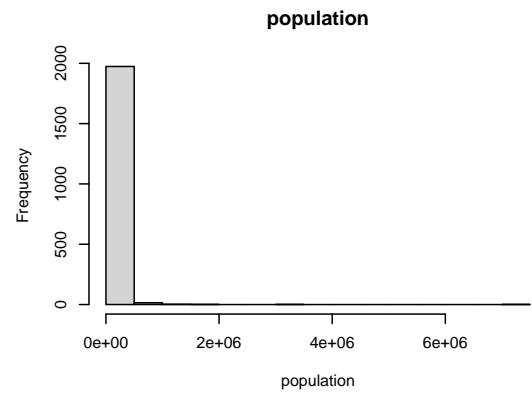
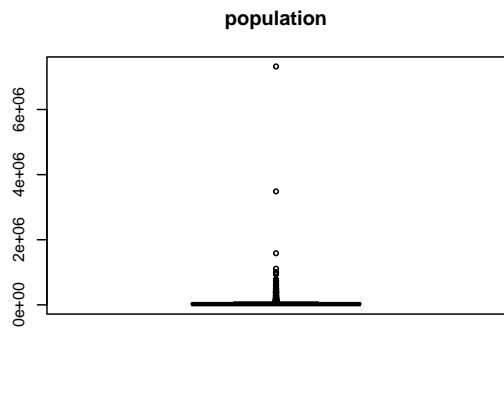
```

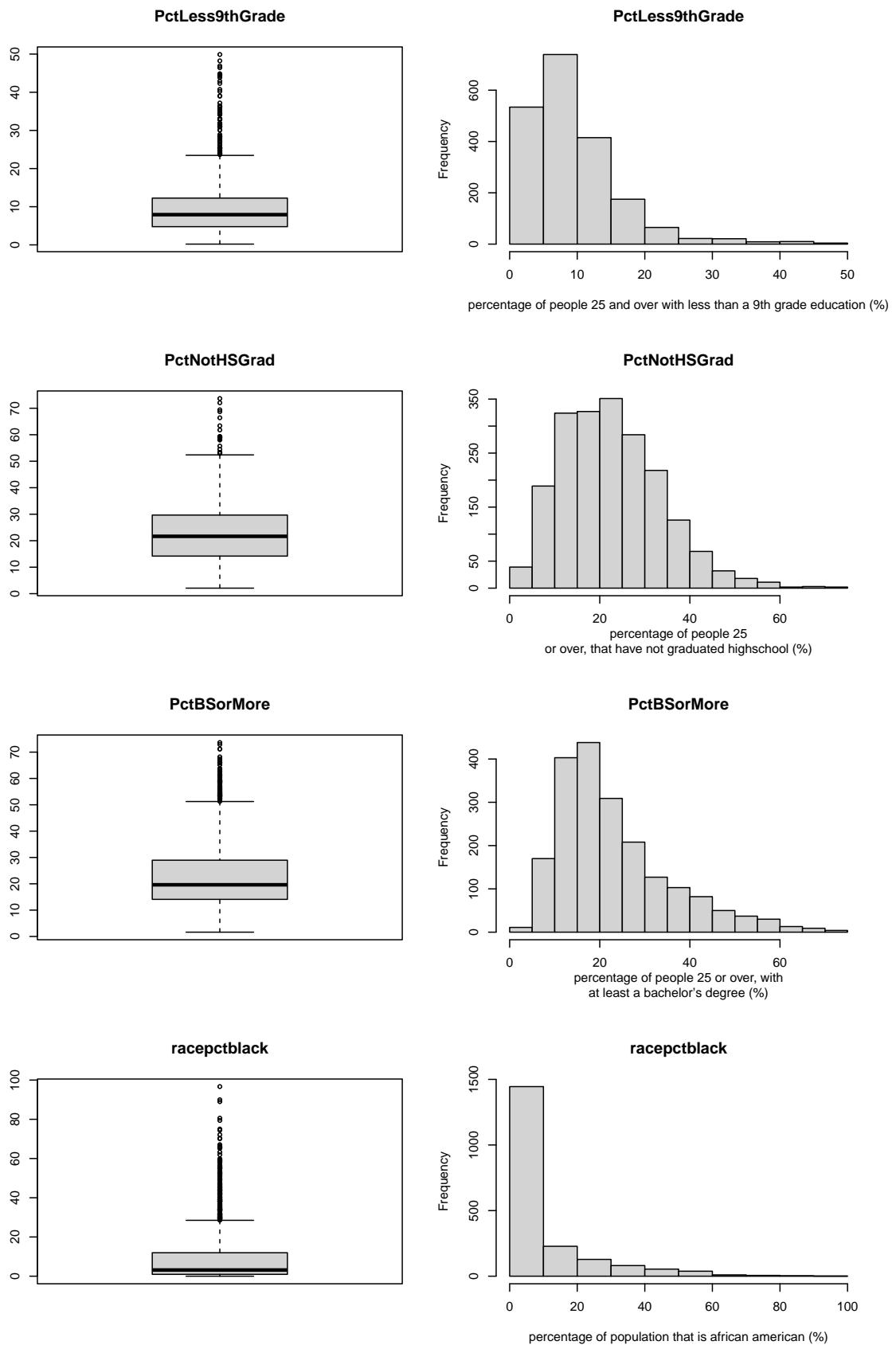
To gain insight into the univariate distributions, boxplots and histograms are generated.

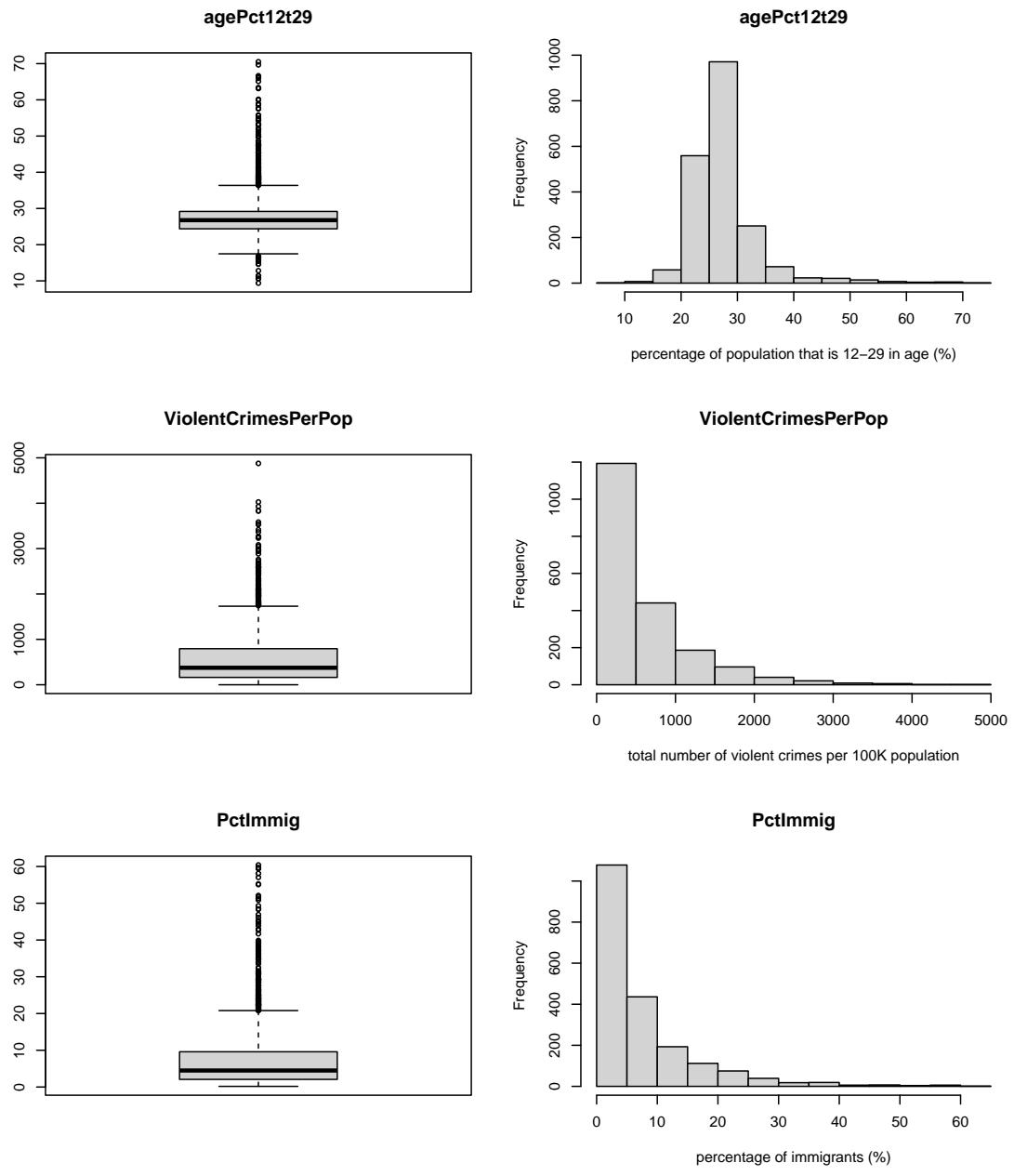
```

numeric_cols <- sapply(crimes_table_subset, is.numeric)
crimes_table_subset_num <- crimes_table_subset[, ..numeric_cols]
par(mfrow = c(4,2))
for(columnname in names(crimes_table_subset_num)){
  column <- crimes_table_subset_num[[columnname]]
  boxplot(column,
           main = columnname
           )
  hist(column,
        main = columnname,
        xlab = get_label(column)
        )
}

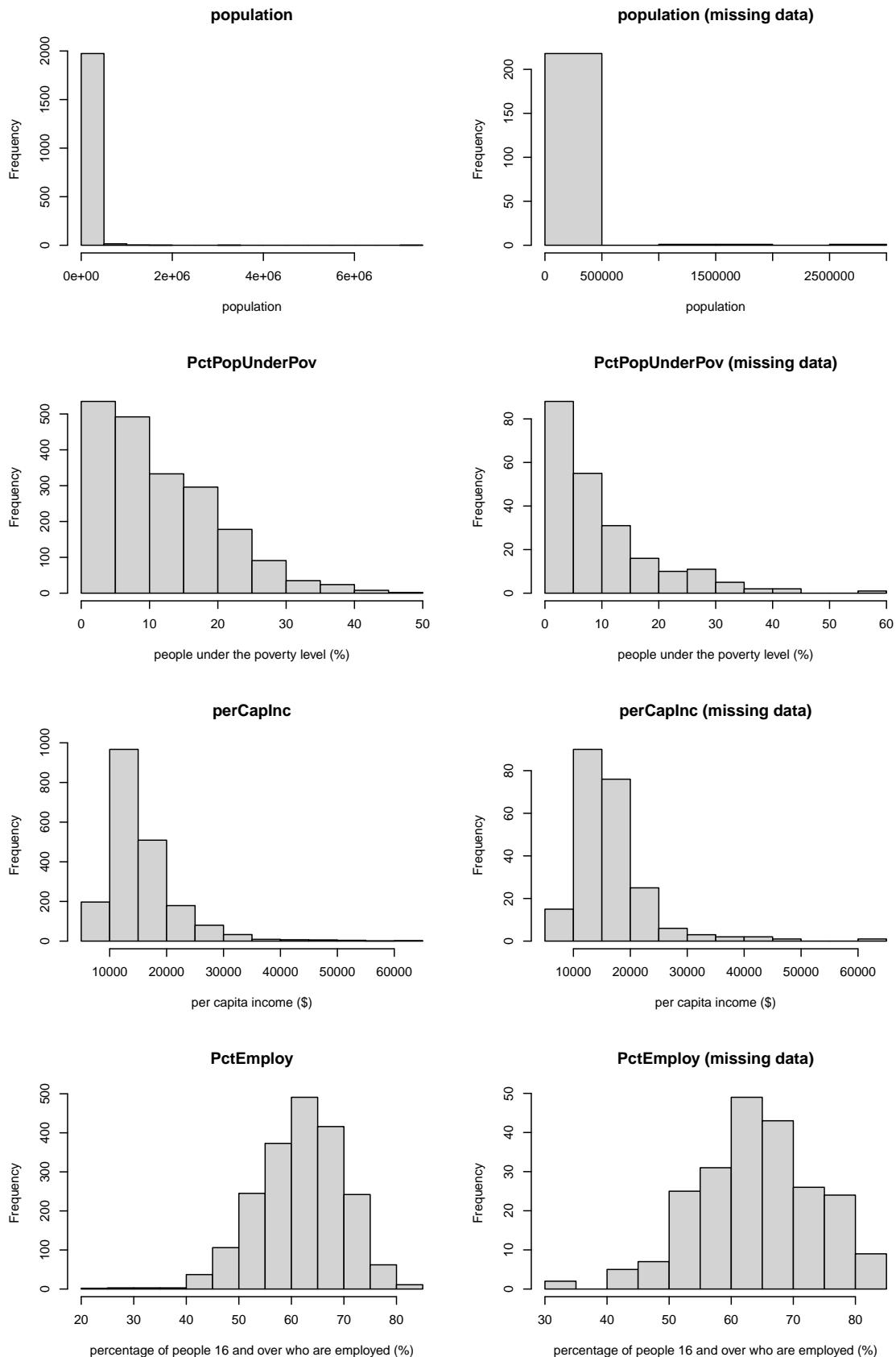
```

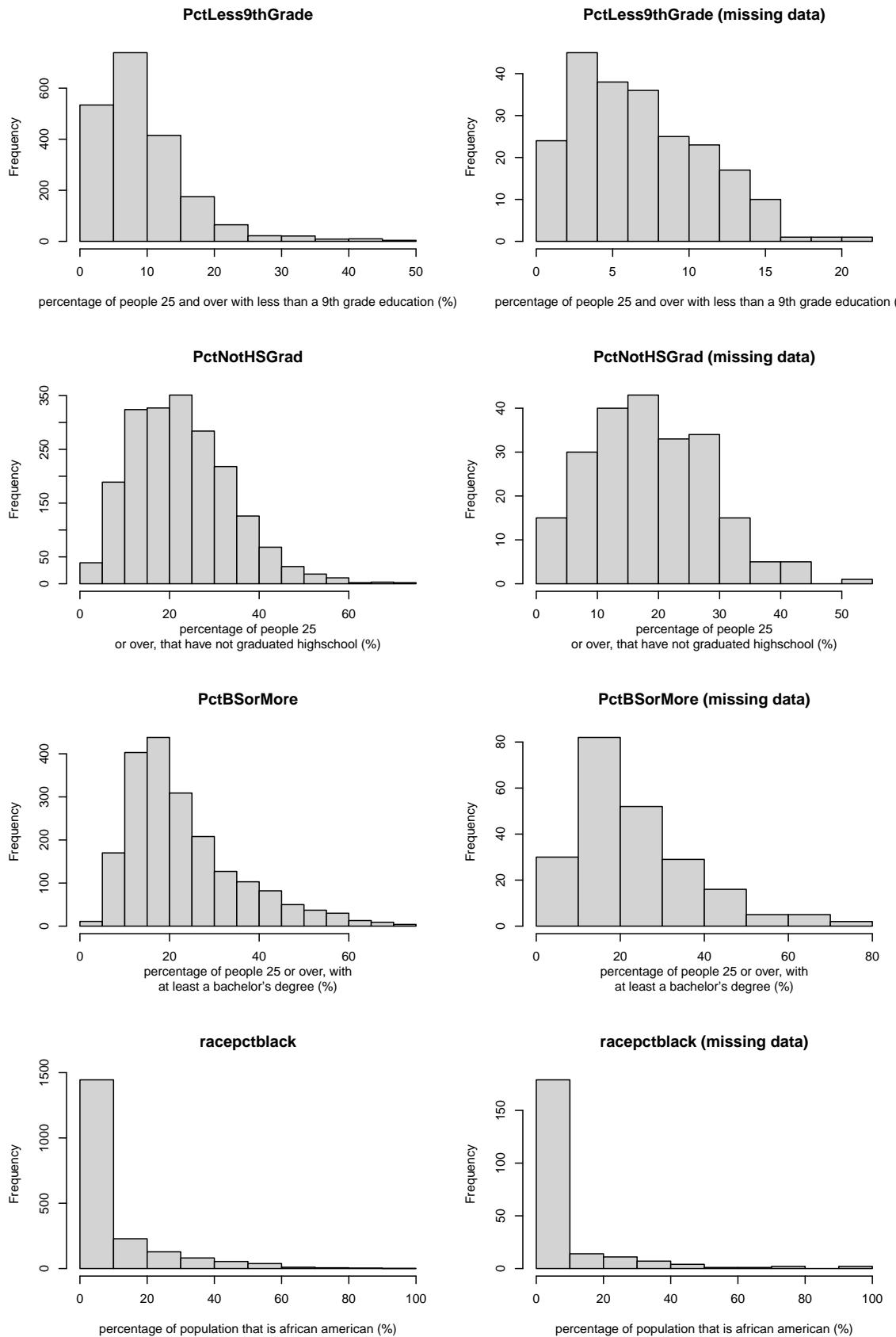


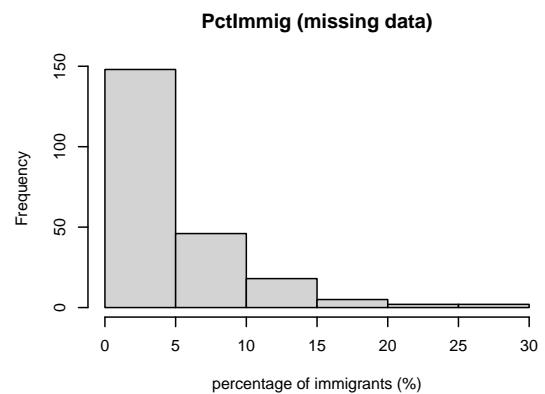
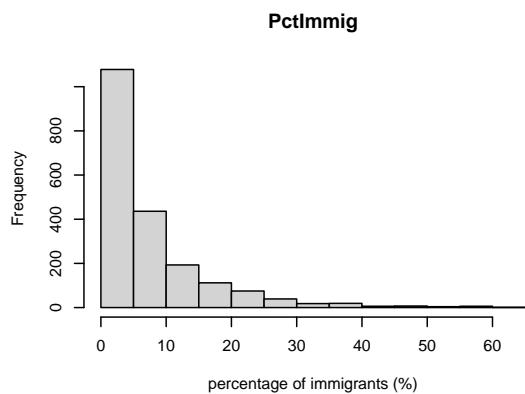
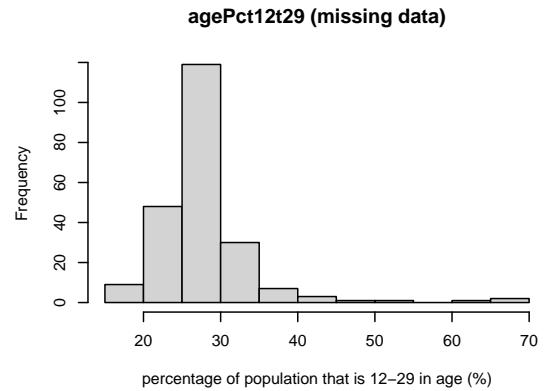
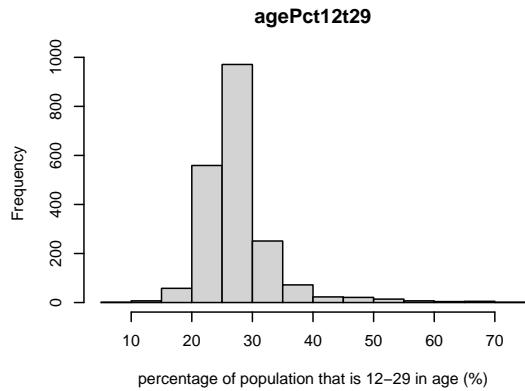




```
numeric_cols_na <- sapply(na_subset, is.numeric)
crimes_table_subset_num_na <- na_subset[, ..numeric_cols_na]
par(mfrow = c(4,2))
for(columnname in names(crimes_table_subset_num_na)){
  column <- crimes_table_subset_num_na[[columnname]]
  column_na <- crimes_table_subset_num_na[[columnname]]
  hist(column,
    main = columnname,
    xlab = get_label(column)
  )
  hist(column_na,
    main = paste(columnname, "(missing data)"),
    xlab = get_label(column)
  )
}
}
```







The correlation matrix shows the extent to which the variables in the dataset are correlated with each other. Below, all variables are listed, sorted from highest to lowest correlation with the outcome variable.

- *racepctblack* ($r = 0.63$)
- *PctPopUnderPov* ($r = 0.51$)
- *PctNotHSGrad* ($r = 0.47$)
- *PctLess9thGrade* ($r = 0.37$)
- *PctEmploy* ($r = -0.32$)
- *perCapInc* ($r = -0.32$)
- *PctBSorMore* ($r = 0.3$)
- *PctImmig* ($r = 0.19$)
- *agePct12t29* ($r = 0.11$)

It's important to mention that these correlations are indicators of an association, not of a causation.

The following predictors are highly correlated with each other. Therefore, it is best not to include them together in a model later.

- *PctNotHSGrad* and *PctLess9thGrade* ($r = 0.93$)
- *perCapInc* and *PctBSorMore* ($r = 0.77$)
- *PctNotHSGrad* and *PctBSorMore* ($r = -0.75$)

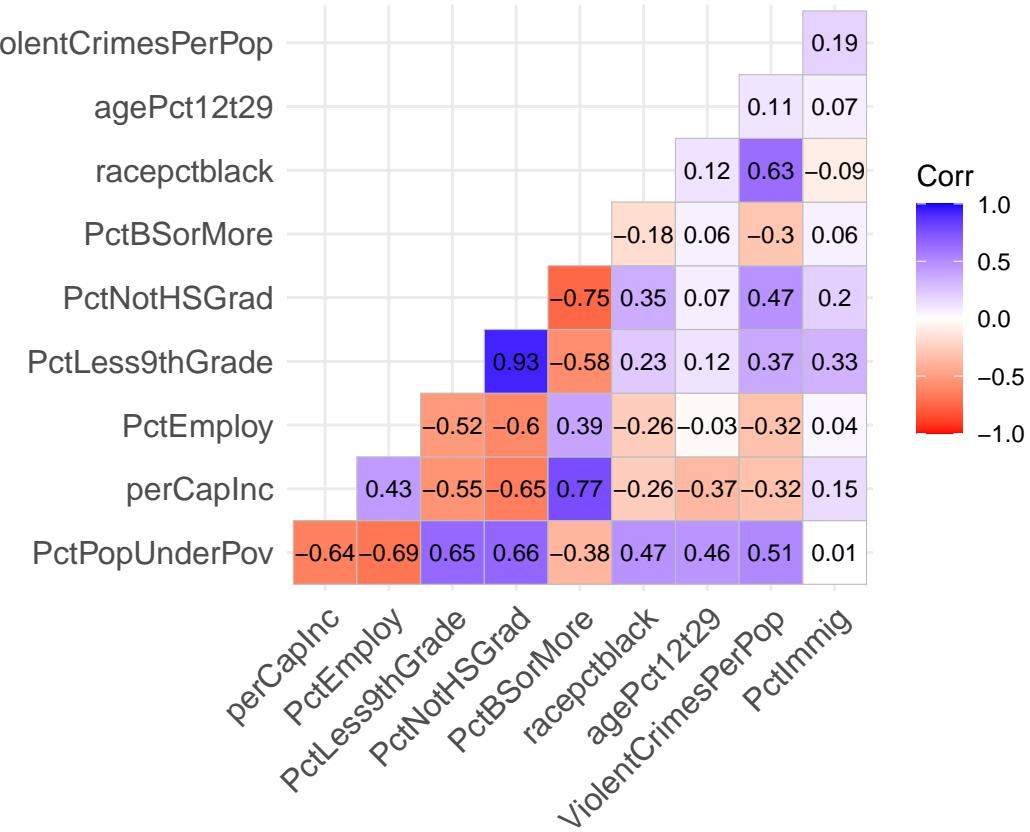
However the choice for predictors for the model will be dealt with thoroughly during the model building.

It is noticeable that the variable *racepctblack* is the one most strongly correlated with the outcome variable ($r = 0.63$), even more than *PctPopUnderPov*, the head predictor that was chosen for this research.

It is noticeable the the variables *PctImmig* and *PctImmig* have correlation coefficients tthat are really low.

```
cor_matrix <- cor(crimes_table_subset_num[,-'population'])
cor_values <- as.data.frame(as.table(cor_matrix))

library(ggcormplot)
ggcorrplot(cor_matrix, lab = TRUE, type = "lower",
           lab_size = 3, colors = c("red", "white", "blue"))
```



The following scatter plots were generated:

- for each variable, a scatter plot showing the relationship with the outcome variable *ViolentCrimesPerPop*;
- for each variable, a scatter plot showing the relationship with the main predictor variable *PctPopUnderPov*.

The first series of scatter plots indicates that not all variables have a linear relationship with *ViolentCrimesPerPop*. In particular, the following variables do not appear to exhibit a clear linear trend:

- *perCapInc*
- *agePct12t29* (which also had a very low correlation coefficient)

Other variables show a somewhat linear pattern, although this trend is often distorted in the extreme regions of the x-axis.

The second series of scatter plots suggests that some variables exhibit a linear relationship with the main predictor *PctPopUnderPov*. In particular, the following variables appear to show a fairly linear trend:

- *PctEmploy*
- *PctLess9thGrade*
- *PctNotHSGrad*

This implies that these variables are probably not suitable as additional predictors when *PctPopUnderPov* is already included in the model.

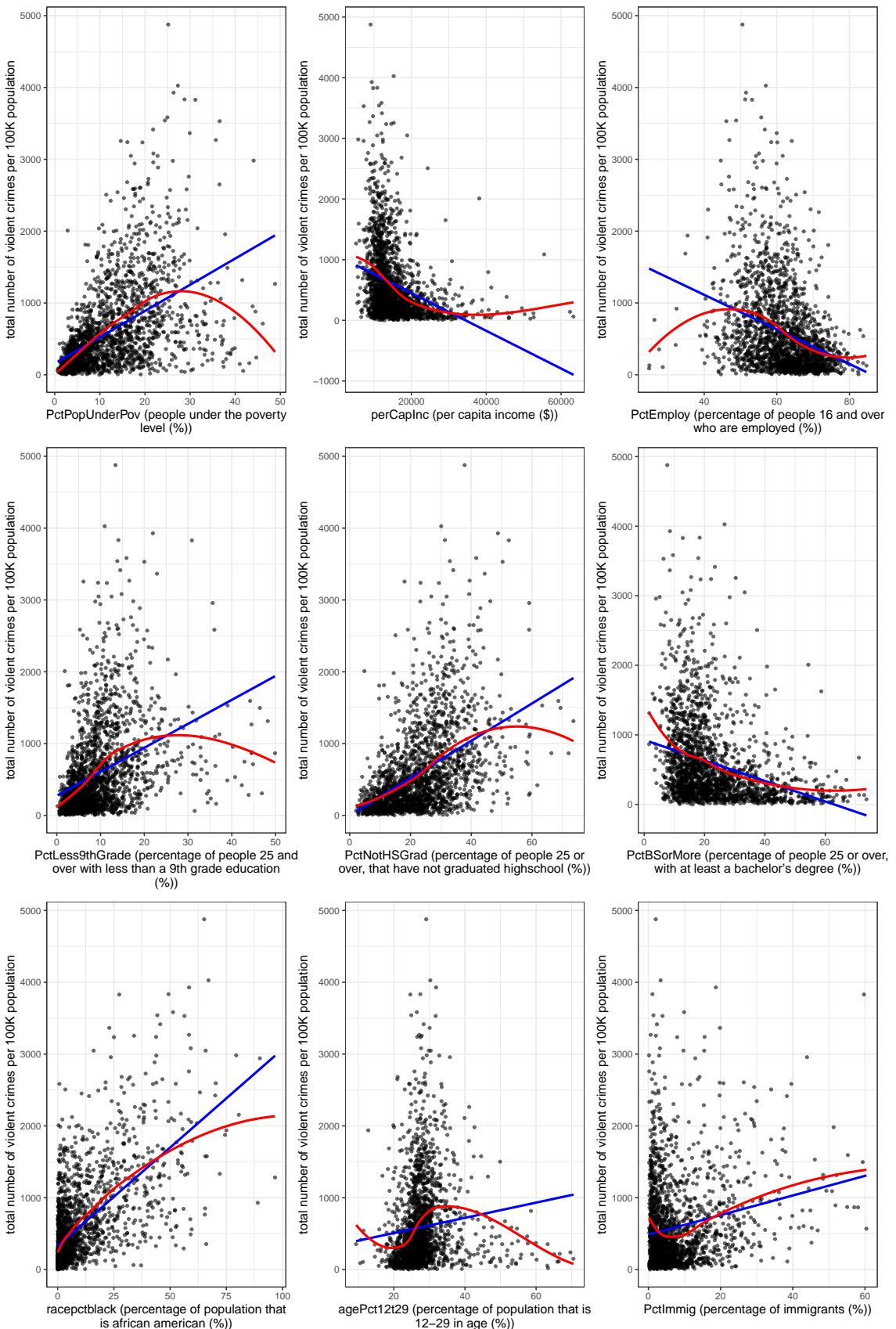
```

x_vars <- colnames(crimes_table_subset_num)
dict_labels <- setNames(sapply(x_vars, function(x_var) get_label(crimes_table_subset_num[[x_var]])), x_var)

library(ggplot2)
library(patchwork)
df <- crimes_table_subset_num[,-c("population", "fold")]
y_var <- "ViolentCrimesPerPop"
x_vars <- setdiff(colnames(df), y_var)
plots <- lapply(x_vars, function(x_var) {
  ggplot(df, aes_string(x_var,y_var)) +
  geom_point(alpha = 0.6, size = 0.7) +
  geom_smooth(method = "lm", color = "blue", se = FALSE) +
  geom_smooth(method = "loess", color = "red", se = FALSE) +
  theme_bw(base_size = 8) +
  labs(x = str_wrap(paste(x_var, " (", dict_labels[x_var], ")"), sep = ""), width = 45))
}
)

# Print 9 plots per pg
print(wrap_plots(plots, ncol = 3))

```

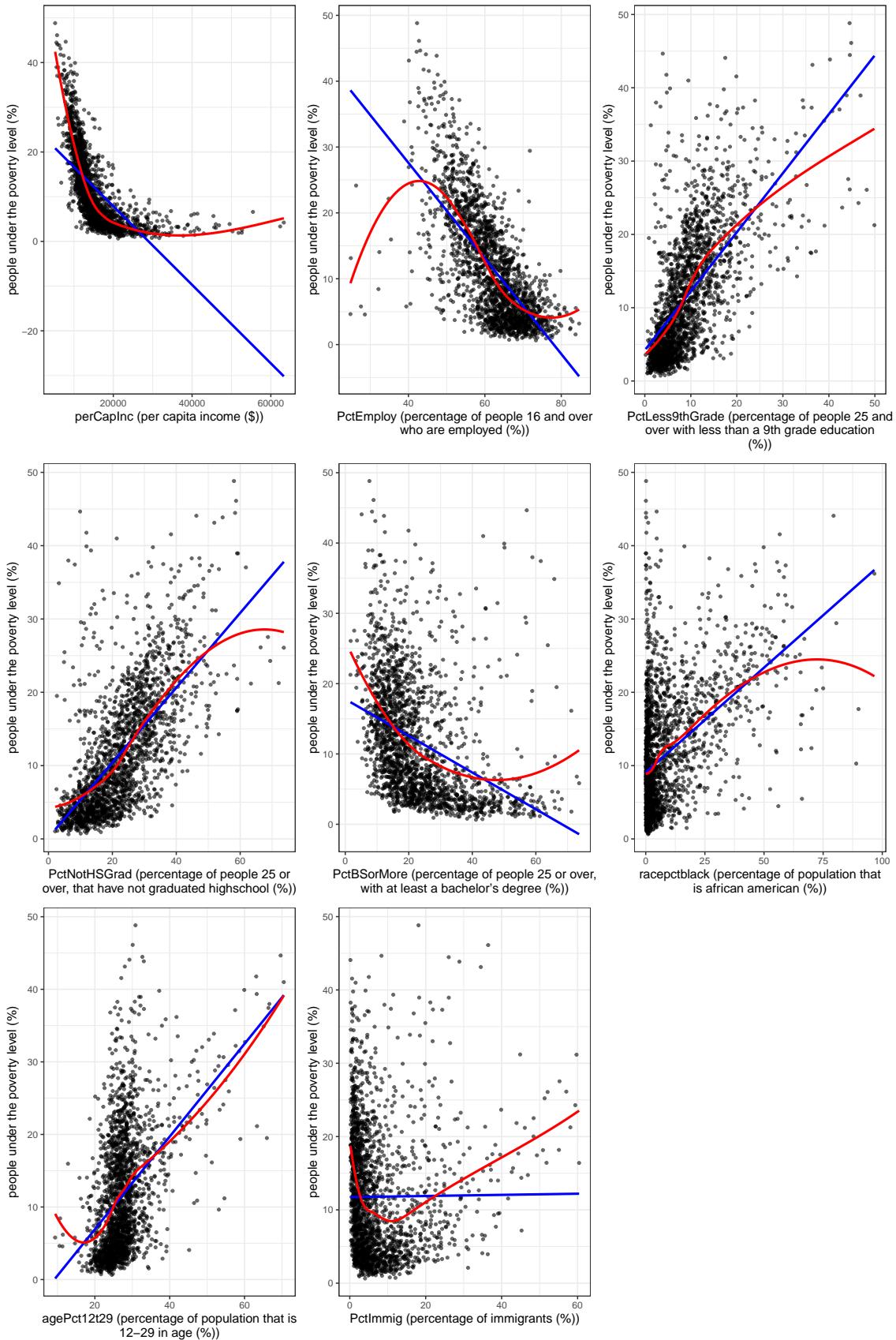


```

df <- crimes_table_subset_num[, -c("ViolentCrimesPerPop", "population", "fold")]
y_var <- "PctPopUnderPov"
x_vars <- setdiff(colnames(df), y_var)
plots <- lapply(x_vars, function(x_var) {
  ggplot(df, aes_string(x_var,y_var))+
  geom_point(alpha = 0.6, size = 0.7) +
  geom_smooth(method = "lm", color = "blue", se = FALSE)+
  geom_smooth(method = "loess", color = "red", se = FALSE) +
  theme_bw(base_size = 8) +
  labs(x = str_wrap(paste(x_var, " (", dict_labels[x_var], ")"), sep = ""), width = 45))
}
)

# Print 9 plots per pg
print(wrap_plots(plots, ncol = 3))

```



The scatter plots reveal one outlier for the ViolentCrimesPerPop variable. This outlier is the community Chestercity.

```
crimes_table_subset[order(crimes_table_subset$num$ViolentCrimesPerPop, decreasing=TRUE), , drop = FALSE]

##           communityname state countyCode communityCode population
##                <char> <char>     <char>        <char>      <int>
## 1:      Chestercity    PA         45       13208      41856
## 2:      Atlantacity    GA          ?          ?      394017
## 3:      Newarkcity     NJ         13       51000      275221
## 4:      Alexandriacity LA          ?          ?      49188
## 5:      Miamicity      FL          ?          ?      358548
## ---
## 1990:      Harvardtown MA         27       28950      12329
## 1991:      Ogdensburgcity NY         89       54485      13521
## 1992:      Cranberrytownship PA         19       16920      14816
## 1993:      Oswegocity   NY         75       55574      19195
## 1994:      Spencercity  IA         41       93955      11066
##           PctPopUnderPov perCapInc PctEmploy PctLess9thGrade PctNotHSGrad
##                <num>     <int>     <num>        <num>      <num>
## 1:      25.16      9115     50.54      13.42      37.84
## 2:      27.29     15279     56.97      10.96      30.13
## 3:      26.34      9424     51.51      21.97      48.79
## 4:      28.78     10887     51.28      14.08      31.37
## 5:      31.17      9799     53.19      30.89      52.37
## ---
## 1990:      3.88     17937     82.48       0.66      2.93
## 1991:     13.97     11213     44.61      10.81      32.03
## 1992:      2.54     16494     71.33       1.97      9.51
## 1993:     19.05     11758     51.56       8.71      26.92
## 1994:      9.87     12805     66.26       6.32      14.70
##           PctBSorMore racepctblack agePct12t29 ViolentCrimesPerPop PctImmig
##                <num>     <num>     <num>        <num>      <num>
## 1:      7.68      65.17     29.11      4877.06  2.0355505
## 2:     26.65      67.07     30.26      4026.59  3.3891939
## 3:      8.55      58.46     31.88      3928.03 18.6842574
## 4:     18.40      49.29     27.45      3834.10  1.1364560
## 5:     12.79      27.39     24.63      3829.21 59.7208742
## ---
## 1990:     42.39     12.22     39.12       7.79  4.5259145
## 1991:     11.85      8.27     29.81       7.60  5.0809851
## 1992:     29.48      0.47     25.19       6.64  1.4916307
## 1993:     19.63      0.73     32.15       5.35  2.4277156
## 1994:     16.09      0.05     24.23       0.00  0.4518344
```

Model Building

Before performing linear regression and building models, the dataset is randomly split into a training set (80% of the data) and a holdout set (20% of the data). This holdout set will be used to validate the final model.

```

n <- nrow(crimes_table_subset_num)
training <- sample(1:n, size = floor(0.8 * n))
train_data <- crimes_table_subset_num[training, ]
test_data <- crimes_table_subset_num[-training, ]

cat("Training set size:", nrow(train_data), "\n")

## Training set size: 1595

cat("Test set size:", nrow(test_data), "\n")

```

Test set size: 399

Univariate linear regression

The simple univariate regression equation we estimate with the training set is given as follows:

$$ViolentCrimesPerPop_i = \beta_0 + \beta_1 \cdot PctPopUnderPov_i + \epsilon_i$$

```

fit <- lm(ViolentCrimesPerPop ~ PctPopUnderPov, data = train_data)
summary(fit)

```

```

##
## Call:
## lm(formula = ViolentCrimesPerPop ~ PctPopUnderPov, data = train_data)
##
## Residuals:
##      Min       1Q   Median       3Q      Max 
## -1517.9  -244.9   -97.7   153.7  3831.6 
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)    
## (Intercept) 157.594    21.853   7.212 8.52e-13 ***
## PctPopUnderPov 35.289     1.512  23.332 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 515.2 on 1593 degrees of freedom
## Multiple R-squared:  0.2547, Adjusted R-squared:  0.2542 
## F-statistic: 544.4 on 1 and 1593 DF,  p-value: < 2.2e-16

```

We show the relevant statistics to be discussed in this section:

```

cat("Regression equation: ViolentCrimesPerPop =",
    round(coef(fit)[1], 2), "+",
    round(coef(fit)[2], 2), "* PctPopUnderPov\n\n")

## Regression equation: ViolentCrimesPerPop = 157.59 + 35.29 * PctPopUnderPov

```

```

# R-squared
cat("R-squared:", round(summary(fit)$r.squared, 4), "\n")

## R-squared: 0.2547

cat("Adjusted R-squared:", round(summary(fit)$adj.r.squared, 4), "\n")

## Adjusted R-squared: 0.2542

# MSE
mse_simple <- mean(fit$residuals^2)
cat("MSE:", round(mse_simple, 2), "\n")

## MSE: 265053

# Confidence intervals for coefficients
cat("\n95% Confidence Intervals:\n")

## 
## 95% Confidence Intervals:

print(confint(fit))

##                   2.5 %    97.5 %
## (Intercept) 114.73033 200.45775
## PctPopUnderPov 32.32253 38.25589

```

PctPopUnderPov = increase in violent crimes per 100K population if poverty rate increases by one percentage point

Assumption checks

We check assumptions linearity, independence of errors, homoscedasticity, and normality of errors.

```

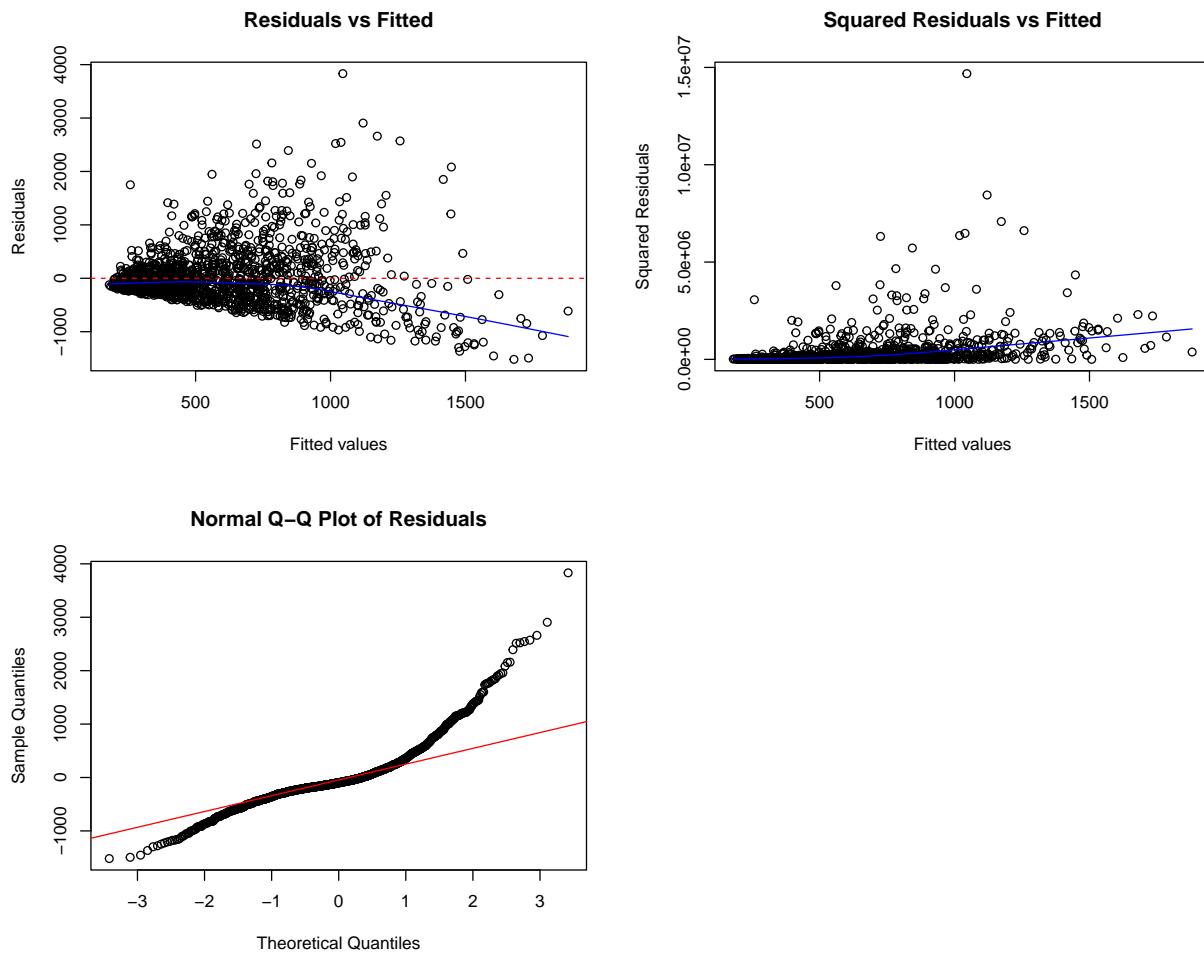
par(mfrow = c(2, 2))

#Residuals vs Fitted
plot(fit$fitted.values, fit$residuals,
      xlab = "Fitted values", ylab = "Residuals",
      main = "Residuals vs Fitted")
abline(h = 0, col = "red", lty = 2)
lines(lowess(fit$fitted.values, fit$residuals), col = "blue")

# Squared residuals vs Fitted
plot(fit$fitted.values, fit$residuals^2,
      xlab = "Fitted values", ylab = "Squared Residuals",
      main = "Squared Residuals vs Fitted")
lines(lowess(fit$fitted.values, fit$residuals^2), col = "blue")

```

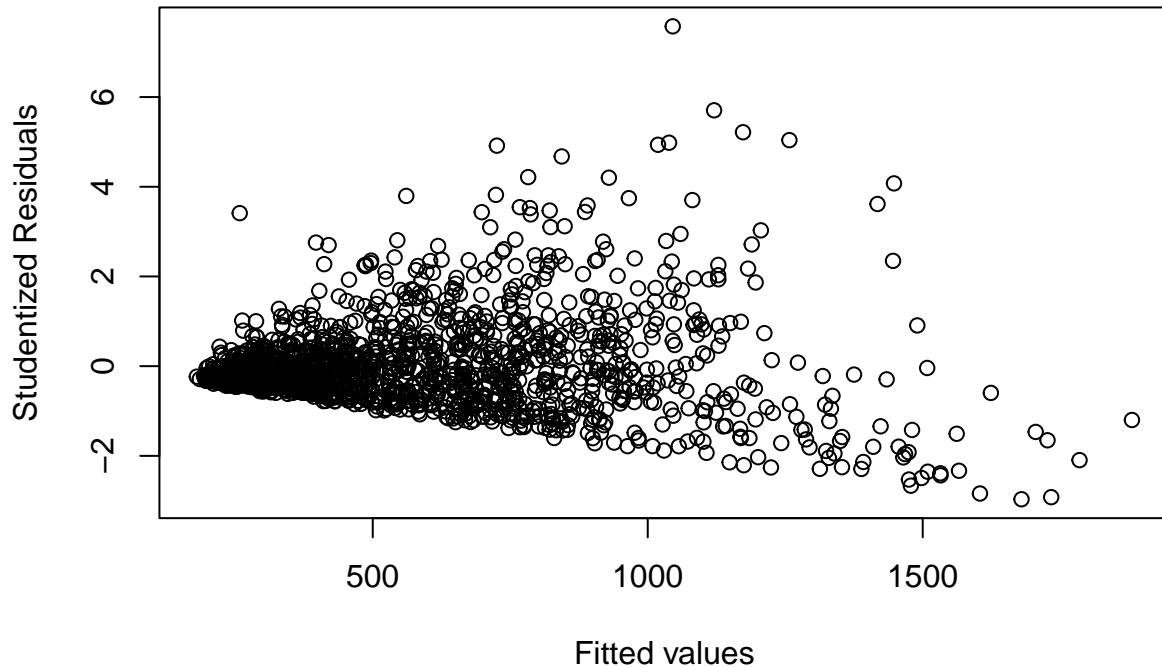
```
# QQ-plot of residuals (normality)
qqnorm(fit$residuals, main = "Normal Q-Q Plot of Residuals")
qqline(fit$residuals, col = "red")
par(mfrow = c(1, 1))
```



Studentized residuals

```
# Studentized residuals plot
stud_res <- rstudent(fit)
plot(fit$fitted.values, stud_res,
      xlab = "Fitted values", ylab = "Studentized Residuals",
      main = "Studentized Residuals vs Fitted")
```

Studentized Residuals vs Fitted



```
outliers_simple <- which(abs(stud_res) > 2)
```

Table to get a visual illustration of whether outliers are more common in small pop

```
# outliers are more present in small pop?
if(length(outliers_simple) > 0) {
  outlier_data <- train_data[outliers_simple, .(population, ViolentCrimesPerPop, PctPopUnderPov)]
  print(outlier_data)
}
```

	population	ViolentCrimesPerPop	PctPopUnderPov
## 1:	30996	1938.43	15.96
## 2:	98052	2109.96	21.17
## 3:	29925	2594.03	17.80
## 4:	10170	137.89	32.74
## 5:	12822	161.69	43.13
## 6:	27334	1726.58	14.93
## 7:	10005	2119.93	24.78
## 8:	21265	239.75	44.66
## 9:	26326	1607.89	10.36
## 10:	109602	2078.85	16.47
## 11:	13547	39.87	28.82
## 12:	33892	2572.91	25.56
## 13:	13024	2649.80	20.64

## 14:	20651	712.74	46.12
## 15:	280015	3235.45	19.44
## 16:	130474	2127.02	21.29
## 17:	723959	1810.07	12.66
## 18:	86905	1651.73	9.35
## 19:	358548	3829.21	31.17
## 20:	28653	178.75	37.33
## 21:	222103	1807.31	7.43
## 22:	34590	112.41	37.43
## 23:	10690	295.76	34.98
## 24:	13051	63.81	30.22
## 25:	20592	2209.01	17.05
## 26:	15464	1640.60	9.28
## 27:	164693	2304.49	15.76
## 28:	88675	2065.29	18.05
## 29:	73552	2941.62	17.71
## 30:	45549	2344.68	21.57
## 31:	141686	1908.96	17.07
## 32:	33497	2758.90	29.70
## 33:	11874	1634.34	9.33
## 34:	87425	3239.20	16.10
## 35:	152466	1960.67	18.18
## 36:	72411	2682.28	16.05
## 37:	10815	307.42	38.95
## 38:	574283	1968.89	18.71
## 39:	12001	2451.06	19.59
## 40:	75695	2089.28	18.75
## 41:	18666	1678.99	11.93
## 42:	12652	284.75	38.96
## 43:	736014	3081.26	21.87
## 44:	87492	3530.78	36.56
## 45:	37986	3583.48	24.97
## 46:	52456	219.36	37.98
## 47:	61815	1707.41	9.62
## 48:	95706	2299.87	29.04
## 49:	17363	2210.88	23.20
## 50:	741952	1682.47	12.47
## 51:	265968	2466.68	24.82
## 52:	395934	1787.20	10.83
## 53:	14903	2018.97	19.62
## 54:	10398	278.92	33.20
## 55:	30705	302.47	38.30
## 56:	435146	2460.11	15.32
## 57:	10201	2650.26	36.51
## 58:	10588	1889.44	14.65
## 59:	21080	2586.29	17.28
## 60:	36118	2507.51	11.43
## 61:	81245	1764.17	15.91
## 62:	23755	3268.26	35.71
## 63:	30326	1884.24	18.65
## 64:	7322564	2097.71	19.29
## 65:	139739	2288.32	27.51
## 66:	26623	3540.57	24.40
## 67:	394017	4026.59	27.29

```

##   68:      36830      2581.96      29.23
##   69:      12200      1938.17      20.54
##   70:      70218      2264.09      21.73
##   71:     219531      2978.69      26.17
##   72:     228537      2017.65      18.88
##   73:      33830      370.67      39.91
##   74:     90454      2169.92      27.50
##   75:     40949      1812.62       6.78
##   76:     16027      1730.67      12.03
##   77:     15023      1987.15      10.97
##   78:     49188      3834.10      28.78
##   79:     41856      4877.06      25.16
##   80:    672971      1681.08      12.81
##   81:     12915      150.31      40.99
##   82:     22754      2523.46      17.84
##   83:     10014      47.11      28.09
##   84:     19378      2008.66       2.85
##   85:     27331      1996.15      13.07
##   86:    3485398      2414.77      18.86
##   87:    124773      1981.45      22.32
##   88:     35701      1678.08       9.60
##   89:     49998      1818.82      15.48
##   90:     22122      2728.14      20.77
##   91:     50961      1846.45      13.25
##   92:     41643      2047.92      16.38
##   93:     12849      420.91      37.04
##   94:     18942      2241.85      25.11
##   95:     86835      157.99      29.56
##   96:     12361      196.79      33.88
##   97:     20807      2885.57      22.90
##   98:    372242      2601.60      18.82
##   99:     11751      1581.25       7.20
## 100:     23478      212.45      34.88
## population ViolentCrimesPerPop PctPopUnderPov

```

Model selection

We use an all-possible regressions procedure to select predictor variables. We include models with a maximum of 5 predictor variables and only models with 0, 1, or 2 education predictor variables. The best model is chosen based on the Bayesian Information Criterion.

```

library(leaps)

# Define predictor variables for model selection
predictors <- c("PctPopUnderPov", "perCapInc", "PctEmploy",
                 "PctLess9thGrade", "PctNotHSGrad", "PctBSorMore",
                 "racepctblack", "agePct12t29", "PctImmig")

# Educ variables
educ <- c("PctLess9thGrade", "PctNotHSGrad", "PctBSorMore")

# data for model selection
data <- train_data[, c("ViolentCrimesPerPop", predictors), with = FALSE]

```

```

# Test reg with max 5 predictors
all_combos <- regsubsets(ViolentCrimesPerPop ~ ., data = data,
                           nvmax = 5, nbest = 10, method = "exhaustive")
all_combos_sum <- summary(all_combos)

# Create results dataframe
results <- data.frame(
  n_predictors = apply(all_combos_sum$which[, -1], 1, sum),
  predictors = apply(all_combos_sum$which[, -1], 1, function(x)
    paste(names(x)[x], collapse = ", ")),
  rsq = all_combos_sum$rsq,
  adjrsq = all_combos_sum$adjr2,
  cp = all_combos_sum$cp,
  bic = all_combos_sum$bic
)

# Count education variables in each model
count_edu_vars <- function(pred_string) {
  sum(sapply(educ, function(v) grep1(v, pred_string)))
}
results$n_edu <- sapply(results$predictors, count_edu_vars)

# Filter: only models with 0, 1, or 2 education variables
BIC_ranking <- results[results$n_edu <= 2, ]

# Sort by BIC (lower is better)
BIC_ranking<- BIC_ranking[order(BIC_ranking$bic), ]

print(head(BIC_ranking, 10))

##      n_predictors
## 40          5
## 41          5
## 42          5
## 43          5
## 30          4
## 44          5
## 45          5
## 46          5
## 31          4
## 47          5
##
##                                predictors
## 40 PctPopUnderPov, PctLess9thGrade, PctNotHSGrad, racepctblack, PctImmig
## 41  PctPopUnderPov, PctLess9thGrade, PctBSorMore, racepctblack, PctImmig
## 42      PctPopUnderPov, PctBSorMore, racepctblack, agePct12t29, PctImmig
## 43          PctPopUnderPov, perCapInc, PctBSorMore, racepctblack, PctImmig
## 30              PctPopUnderPov, PctBSorMore, racepctblack, PctImmig
## 44      PctPopUnderPov, PctNotHSGrad, PctBSorMore, racepctblack, PctImmig
## 45          PctPopUnderPov, PctEmploy, PctBSorMore, racepctblack, PctImmig
## 46      PctPopUnderPov, perCapInc, racepctblack, agePct12t29, PctImmig
## 31          PctPopUnderPov, racepctblack, agePct12t29, PctImmig
## 47  PctPopUnderPov, PctLess9thGrade, racepctblack, agePct12t29, PctImmig
##      rsq      adjrsq        cp        bic n_edu

```

```

## 40 0.5405806 0.5391350 55.53845 -1196.330      2
## 41 0.5333709 0.5319026 81.25195 -1171.494      2
## 42 0.5299952 0.5285163 93.29165 -1159.997      1
## 43 0.5281873 0.5267027 99.73965 -1153.873      1
## 30 0.5248257 0.5236303 109.72874 -1149.924      1
## 44 0.5259742 0.5244826 107.63283 -1146.409      2
## 45 0.5248994 0.5234044 111.46607 -1142.797      1
## 46 0.5244528 0.5229564 113.05905 -1141.298      0
## 31 0.5196014 0.5183928 128.36173 -1132.484      0
## 47 0.5214140 0.5199080 123.89694 -1131.139      1

```

We then run the multivariate regression equation

```

# Print best model
best <- which.min(BIC_ranking$bic)
best_pred <- BIC_ranking$predictors[best]
cat("\nBest model:\n")

##
## Best model:

cat("Predictors:", best_pred, "\n")

## Predictors: PctPopUnderPov, PctLess9thGrade, PctNotHSGrad, racepctblack, PctImmig

cat("BIC:", round(BIC_ranking$bic[best], 2), "\n")

## BIC: -1196.33

cat("Adjusted R2:", round(BIC_ranking$adjrsq[best], 4), "\n")

## Adjusted R2: 0.5391

# store pred
best_pred_sel <- strsplit(best_pred, " ", "")[[1]]

# multivariate regression with these predictors
formula_multi <- as.formula(paste("ViolentCrimesPerPop ~",
                                    paste(best_pred_sel, collapse = " + ")))
fit_multi <- lm(formula_multi, data = train_data)
summary(fit_multi)

##
## Call:
## lm(formula = formula_multi, data = train_data)
##
## Residuals:
##       Min        1Q    Median        3Q       Max
## -1430.10   -205.07   -46.96   136.02  2752.99
##
```

```

## Coefficients:
##                               Estimate Std. Error t value Pr(>|t|)
## (Intercept)           -210.9603   30.9090 -6.825  1.25e-11 ***
## PctPopUnderPov      16.7847    1.8095  9.276 < 2e-16 ***
## PctLess9thGrade     -45.0986   4.7495 -9.495 < 2e-16 ***
## PctNotHSGrad        29.5885   2.8013 10.563 < 2e-16 ***
## racepctblack       20.8437   0.8947 23.297 < 2e-16 ***
## PctImmig            19.6195   1.3468 14.568 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 405 on 1589 degrees of freedom
## Multiple R-squared:  0.5406, Adjusted R-squared:  0.5391
## F-statistic: 373.9 on 5 and 1589 DF,  p-value: < 2.2e-16

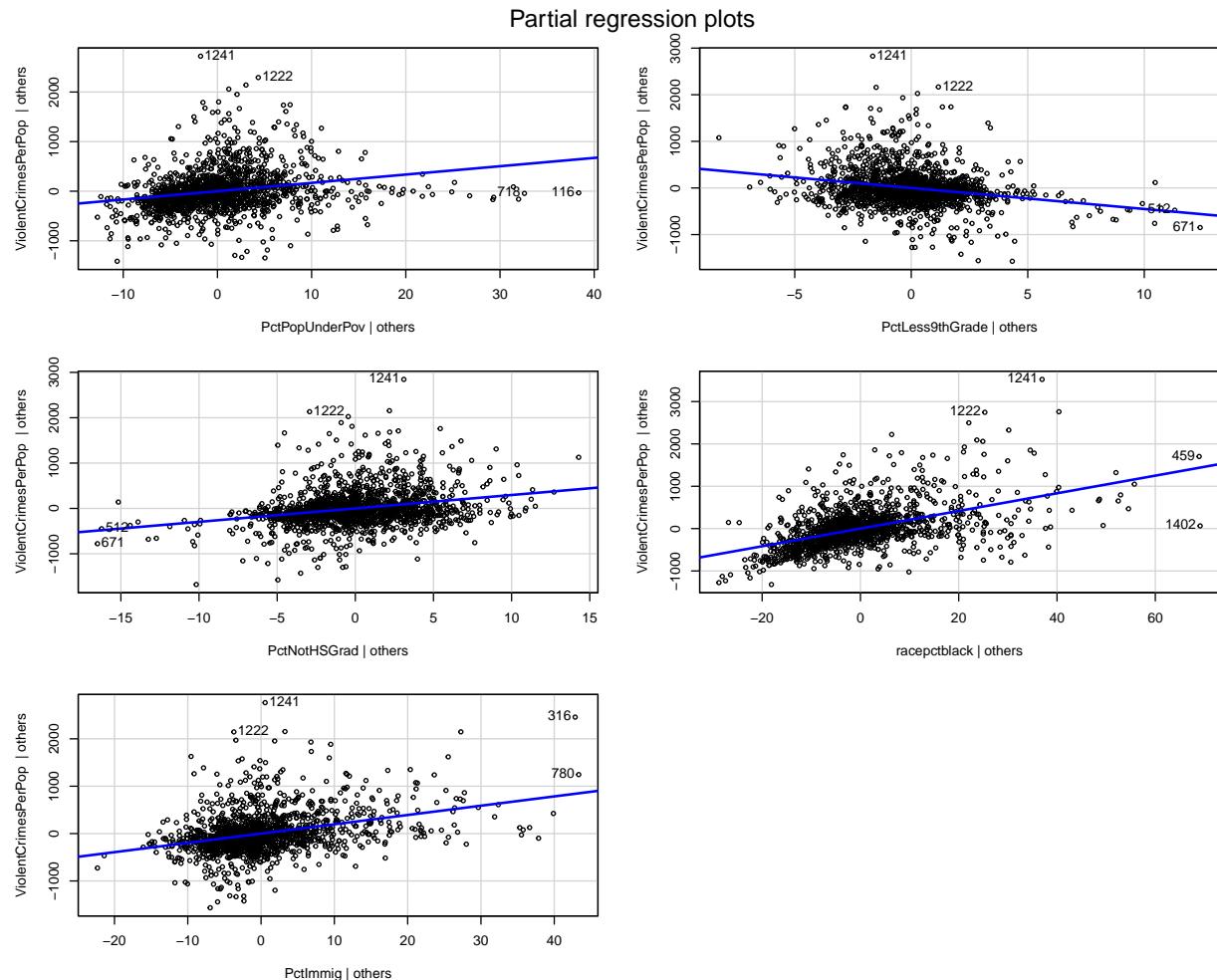
```

Partial Regression Plots

```

library(car)
avPlots(fit_multi, main = "Partial regression plots")

```



Interaction Terms Selection

Following the protocol, we add all interaction terms of the predictor variables with our main predictor variable *PctPopUnderPov* and evaluate their significance. We choose the best one.

```
# store all other pred except for main pred
other <- setdiff(best_pred_sel, "PctPopUnderPov")

# Create interaction terms
interaction_terms <- paste("PctPopUnderPov", other, sep = ":")

# Evaluate each interaction term individually
interaction_results <- data.frame(
  interaction = interaction_terms,
  t_value = NA,
  p_value = NA,
  delta_adjrsq = NA
)

for(i in seq_along(interaction_terms)) {
  formula_single_int <- as.formula(paste("ViolentCrimesPerPop ~",
                                           paste(best_pred_sel, collapse = " + "), "+",
                                           interaction_terms[i]))
  fit_single_int <- lm(formula_single_int, data = train_data)
  coef_summary <- summary(fit_single_int)$coefficients
  int_row <- nrow(coef_summary)
  interaction_results$t_value[i] <- coef_summary[int_row, "t value"]
  interaction_results$p_value[i] <- coef_summary[int_row, "Pr(>|t|)"]
  interaction_results$delta_adjrsq[i] <- summary(fit_single_int)$adj.r.squared -
    summary(fit_multi)$adj.r.squared
}

interaction_results <- interaction_results[order(interaction_results$p_value), ]
cat("Interaction terms ranked by p-value:\n")

## Interaction terms ranked by p-value:

print(interaction_results)

##          interaction      t_value      p_value  delta_adjrsq
## 4 PctPopUnderPov:PctImmig  4.3495099 1.451276e-05  5.138960e-03
## 2 PctPopUnderPov:PctNotHSGrad 1.1106339 2.668941e-01  6.771535e-05
## 1 PctPopUnderPov:PctLess9thGrade -0.6994640 4.843646e-01 -1.481829e-04
## 3 PctPopUnderPov:racepctblack -0.4819476 6.299096e-01 -2.227749e-04

# Select best interaction
best_interaction <- interaction_results$interaction[1]
cat("\nSelected interaction term:", best_interaction, "\n")

##
## Selected interaction term: PctPopUnderPov:PctImmig
```

Multivariate Model

Based on the model selection procedure, we fit the multivariate model including the selected interaction term.

```
# Estimate model with interaction
formula_final <- as.formula(paste("ViolentCrimesPerPop ~",
                                paste(best_pred_sel, collapse = " + "), "+",
                                best_interaction))

fit_final <- lm(formula_final, data = train_data)
summary(fit_final)

##
## Call:
## lm(formula = formula_final, data = train_data)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1362.64  -197.20   -47.71  131.99  2736.70
##
## Coefficients:
##                               Estimate Std. Error t value Pr(>|t|)
## (Intercept)                -146.2336   34.1492 -4.282 1.96e-05 ***
## PctPopUnderPov               12.3935    2.0632  6.007 2.34e-09 ***
## PctLess9thGrade              -51.0327   4.9161 -10.381 < 2e-16 ***
## PctNotHSGrad                  31.5151   2.8206  11.173 < 2e-16 ***
## racepctblack                  21.6761   0.9100  23.819 < 2e-16 ***
## PctImmig                      11.1444   2.3644   4.713 2.65e-06 ***
## PctPopUnderPov:PctImmig        0.6206   0.1427   4.350 1.45e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 402.7 on 1588 degrees of freedom
## Multiple R-squared:  0.546, Adjusted R-squared:  0.5443
## F-statistic: 318.3 on 6 and 1588 DF,  p-value: < 2.2e-16

# confint
print(confint(fit_final))

##
##                               2.5 %      97.5 %
## (Intercept)                -213.2159357 -79.2512316
## PctPopUnderPov                 8.3465416  16.4404847
## PctLess9thGrade              -60.6753498 -41.3900261
## PctNotHSGrad                  25.9826622  37.0476213
## racepctblack                  19.8911021  23.4611324
## PctImmig                      6.5067351  15.7820906
## PctPopUnderPov:PctImmig        0.3407421  0.9004886
```

Multicollinearity Check

Before checking model assumptions, we first assess multicollinearity using the Variance Inflation Factor (VIF) and remove a variable if necessary.

```

library(car)
vif <- vif(fit_final)
print(vif)

##          PctPopUnderPov      PctLess9thGrade      PctNotHSGrad
##            3.045230           10.914754           9.379288
##      racepctblack      PctImmig PctPopUnderPov:PctImmig
##            1.549970           4.036734           5.791617

```

There is multicollinearity, as PctNotHSGrad and PctLess9thGrade are highly correlated (also already derived before from Ilja's plots). Thus, we remove the variable with the highest VIF (PctLess9thGrade). We then do our model evaluation again, and find that the most relevant interaction term to include is now PctPopUnderPov*PctLess9thGrade

```

# Highest vif variable
main_effects_vif <- vif[!grepl(":", names(vif))]
highest_vif_var <- names(which.max(main_effects_vif))

# remove
best_predictors_reduced <- setdiff(best_pred_sel, highest_vif_var)

# Re-evaluate interaction terms without the removed variable
other_predictors_reduced <- setdiff(best_predictors_reduced, "PctPopUnderPov")
interaction_terms_reduced <- paste("PctPopUnderPov", other_predictors_reduced, sep = ":")

interaction_results_reduced <- data.frame(
  interaction = interaction_terms_reduced,
  t_value = NA,
  p_value = NA
)

for(i in seq_along(interaction_terms_reduced)) {
  formula_int <- as.formula(paste("ViolentCrimesPerPop ~",
                                   paste(best_predictors_reduced, collapse = " + "), "+",
                                   interaction_terms_reduced[i]))
  fit_int <- lm(formula_int, data = train_data)
  coef_summary <- summary(fit_int)$coefficients
  int_row <- nrow(coef_summary)
  interaction_results_reduced$t_value[i] <- coef_summary[int_row, "t value"]
  interaction_results_reduced$p_value[i] <- coef_summary[int_row, "Pr(>|t|)"]
}

interaction_results_reduced <- interaction_results_reduced[order(interaction_results_reduced$p_value), ]
cat("Interaction terms ranked by p-value:\n")

## Interaction terms ranked by p-value:

print(interaction_results_reduced)

##          interaction      t_value      p_value
## 1 PctPopUnderPov:PctNotHSGrad -3.7295862 0.0001985316
## 3      PctPopUnderPov:PctImmig  1.4797650 0.1391341948
## 2 PctPopUnderPov:racepctblack -0.5451314 0.5857396230

```

```

best_interaction_reduced <- interaction_results_reduced$interaction[1]
cat("\nSelected interaction term:", best_interaction_reduced, "\n")

##
## Selected interaction term: PctPopUnderPov:PctNotHSGrad

# fit reduced model
formula_final_reduced <- as.formula(paste("ViolentCrimesPerPop ~",
                                             paste(best_predictors_reduced, collapse = " + "), "+",
                                             best_interaction_reduced))

fit_final_reduced <- lm(formula_final_reduced, data = train_data)
summary(fit_final_reduced)

##
## Call:
## lm(formula = formula_final_reduced, data = train_data)
##
## Residuals:
##      Min       1Q   Median       3Q      Max 
## -1575.81  -204.42   -54.01   136.82  2854.96 
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)    
## (Intercept) -132.32009  37.91231 -3.490 0.000496 ***
## PctPopUnderPov 20.63786   3.00596  6.866 9.46e-12 ***
## PctNotHSGrad  10.78928   1.84738  5.840 6.31e-09 ***
## racepctblack  23.44057   0.87140 26.900 < 2e-16 ***
## PctImmig     15.27991   1.28034 11.934 < 2e-16 ***
## PctPopUnderPov:PctNotHSGrad -0.34853   0.09345 -3.730 0.000199 *** 
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1 

## Residual standard error: 414.5 on 1589 degrees of freedom
## Multiple R-squared:  0.5187, Adjusted R-squared:  0.5172 
## F-statistic: 342.5 on 5 and 1589 DF,  p-value: < 2.2e-16

cat("\n95% Confidence Intervals:\n")

##
## 95% Confidence Intervals:

print(confint(fit_final_reduced))

##                               2.5 %      97.5 %
## (Intercept) -206.6834945 -57.9566894
## PctPopUnderPov 14.7417902  26.5339267
## PctNotHSGrad  7.1657326  14.4128322
## racepctblack 21.7313481  25.1497957
## PctImmig     12.7685823  17.7912380
## PctPopUnderPov:PctNotHSGrad -0.5318214 -0.1652295

```

```
# check vif again-->Correct
vif_adapted <- vif(fit_final_reduced)
print(vif_adapted)
```

##	PctPopUnderPov	PctNotHSGrad
	6.101430	3.797917
##	racepctblack	PctImmig
##	1.341479	1.117333
## PctPopUnderPov:PctNotHSGrad		
##	10.659845	

We see that our model performs only a little less well, but this way we did account for multicollinearity and our estimates are correct.

```
# Compare models
cat("Comparison of models:\n")

## Comparison of models:

cat("Original model adjusted R2: ", round(summary(fit_multi)$adj.r.squared, 4), "\n")

## Original model adjusted R2: 0.5391

cat("Reduced model adjusted R2: ", round(summary(fit_final_reduced)$adj.r.squared, 4), "\n")

## Reduced model adjusted R2: 0.5172

# Update fit_final
fit_final <- fit_final_reduced
formula_final <- formula_final_reduced
best_predictors <- best_predictors_reduced
```

Assumption checks final model

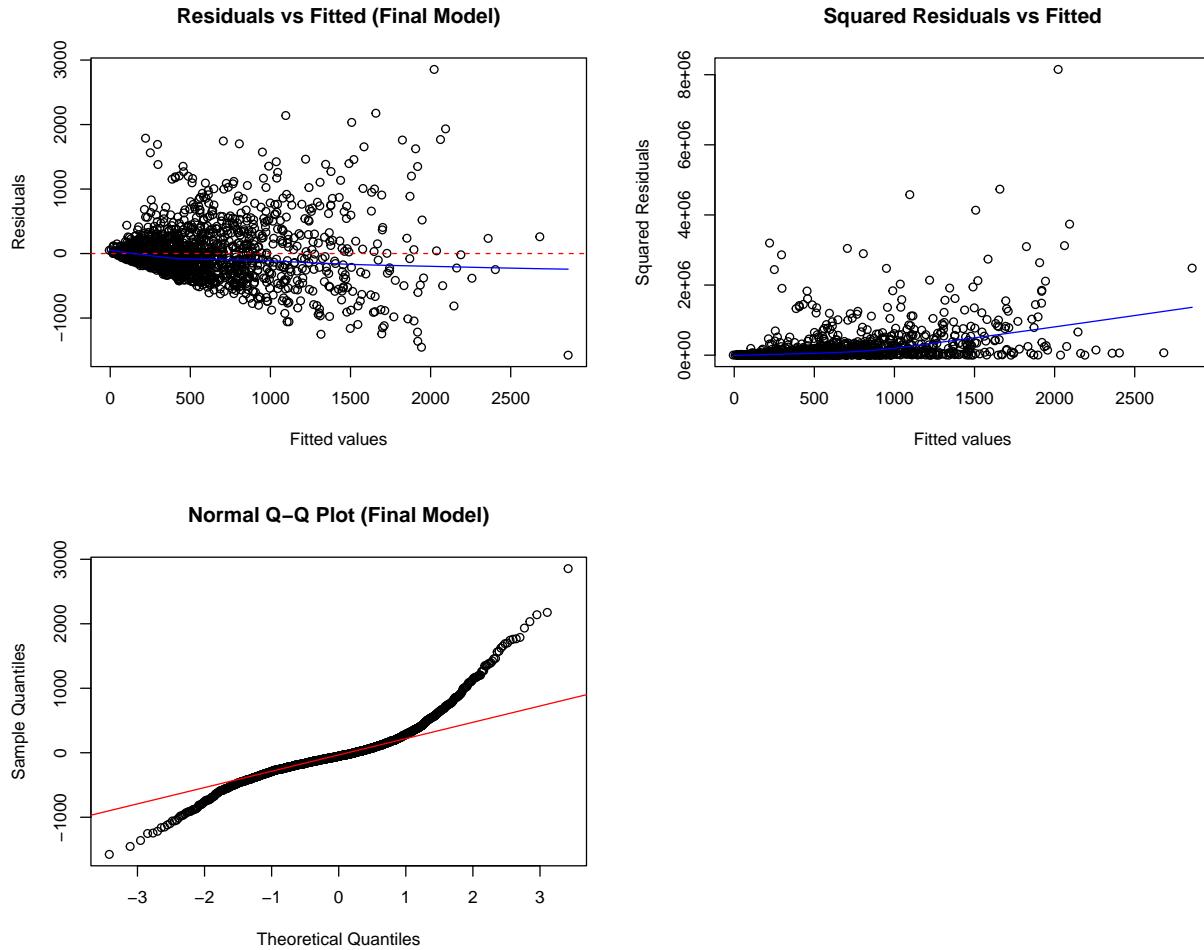
```
par(mfrow = c(2, 2))

#Residuals vs Fitted
plot(fit_final$fitted.values, fit_final$residuals,
      xlab = "Fitted values", ylab = "Residuals",
      main = "Residuals vs Fitted (Final Model)")
abline(h = 0, col = "red", lty = 2)
lines(lowess(fit_final$fitted.values, fit_final$residuals), col = "blue")

# Squared residuals vs Fitted
plot(fit_final$fitted.values, fit_final$residuals^2,
      xlab = "Fitted values", ylab = "Squared Residuals",
      main = "Squared Residuals vs Fitted")
lines(lowess(fit_final$fitted.values, fit_final$residuals^2), col = "blue")
```

```
# QQ-plot
qqnorm(fit_final$residuals, main = "Normal Q-Q Plot (Final Model)")
qqline(fit_final$residuals, col = "red")

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



Robustness checks

For the final regression function, we included robust regression for outliers and heterosced-robust standard errors.

```
library(sandwich)
library(lmtest)
library(MASS)

robust <- rlm(formula_final, data = train_data)
robust_se <- coeftest(robust, vcov = vcovHC(robust, type = "HC3"))
print(robust_se)
```

```
##
```

```

## z test of coefficients:
##
##                               Estimate Std. Error z value Pr(>|z|)
## (Intercept)           -102.40098   30.14713 -3.3967 0.000682 ***
## PctPopUnderPov        17.02242    4.00072  4.2548 2.092e-05 ***
## PctNotHSGrad          8.95953    1.56000  5.7433 9.285e-09 ***
## racepctblack         21.96445    1.40385 15.6459 < 2.2e-16 ***
## PctImmig              13.60872    1.57840  8.6218 < 2.2e-16 ***
## PctPopUnderPov:PctNotHSGrad -0.25088   0.11940 -2.1012 0.035626 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

#idk of die robustness nodig is of er een andere methode geprefereerd is, dit is comparison maar blijft er niet in

```

# Compare OLS vs Robust coefficients
comparison <- data.frame(
  OLS = coef(fit_final),
  Robust = coef(robust),
  Difference = coef(fit_final) - coef(robust)
)
print(round(comparison, 4))

```

	OLS	Robust	Difference
## (Intercept)	-132.3201	-102.4010	-29.9191
## PctPopUnderPov	20.6379	17.0224	3.6154
## PctNotHSGrad	10.7893	8.9595	1.8298
## racepctblack	23.4406	21.9644	1.4761
## PctImmig	15.2799	13.6087	1.6712
## PctPopUnderPov:PctNotHSGrad	-0.3485	-0.2509	-0.0976

Outlier and Influence Diagnostics

We use several diagnostic measures to identify influential observations

```

# Calculate diagnostics
stud_res_final <- rstudent(fit_final)
leverage <- hatvalues(fit_final)
p <- length(coef(fit_final))
n_train <- nrow(train_data)
leverage_threshold <- 2 * p / n_train
cooks_d <- cooks.distance(fit_final)
dffits_val <- dffits(fit_final)
dffits_threshold <- 2 * sqrt(p / n_train)
dfbetas_val <- dfbetas(fit_final)
dfbetas_threshold <- 2 / sqrt(n_train)

# data frame
diagnostics <- data.frame(
  obs = 1:n_train,
  population = train_data$population,
  stud_residual = stud_res_final,

```

```

        leverage = leverage,
        cooks_d = cooks_d,
        dffits = dffits_val
    )

    # Flag observations
    diagnostics$outlier_residual <- abs(diagnostics$stud_residual) > 2
    diagnostics$high_leverage <- diagnostics$leverage > leverage_threshold
    diagnostics$high_cooks <- diagnostics$cooks_d > 4 / n_train
    diagnostics$high_dffits <- abs(diagnostics$dffits) > dffits_threshold

    # summ
    cat("Outliers by studentized residuals (|r*| > 2):", sum(diagnostics$outlier_residual), "\n")

## Outliers by studentized residuals (|r*| > 2): 98

cat("High leverage observations (h >", round(leverage_threshold, 4), "):",
    sum(diagnostics$high_leverage), "\n")

## High leverage observations (h > 0.0075 ): 150

cat("High Cook's distance (D >", round(4/n_train, 4), "):",
    sum(diagnostics$high_cooks), "\n")

## High Cook's distance (D > 0.0025 ): 134

cat("High DFFITS (|DFFITS| >", round(dffits_threshold, 4), "):",
    sum(diagnostics$high_dffits), "\n")

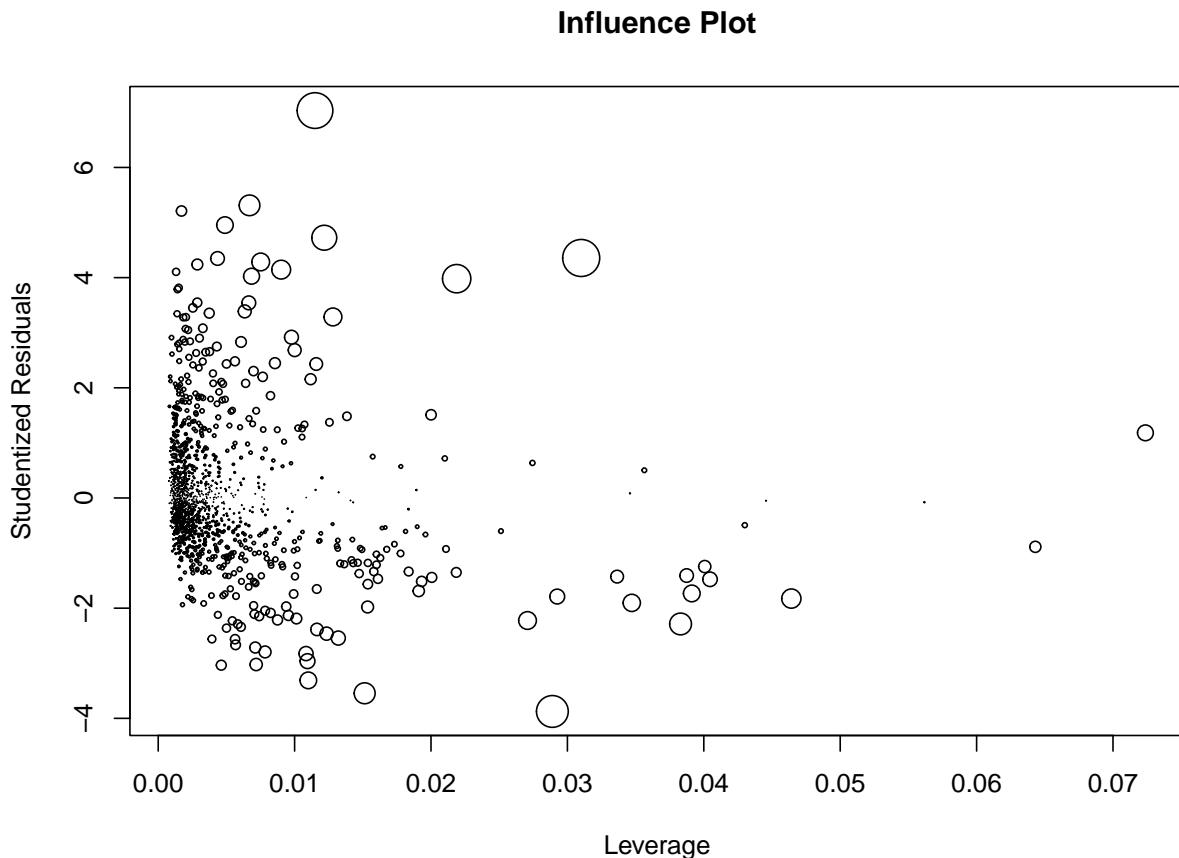
## High DFFITS (|DFFITS| > 0.1227 ): 134

# find influent obs
influential <- diagnostics[diagnostics$high_cooks | diagnostics$high_dffits, ]
influential <- influential[order(-influential$cooks_d), ]
print(head(influential[, c("obs", "population", "stud_residual", "leverage", "cooks_d", "dffits")], 10))

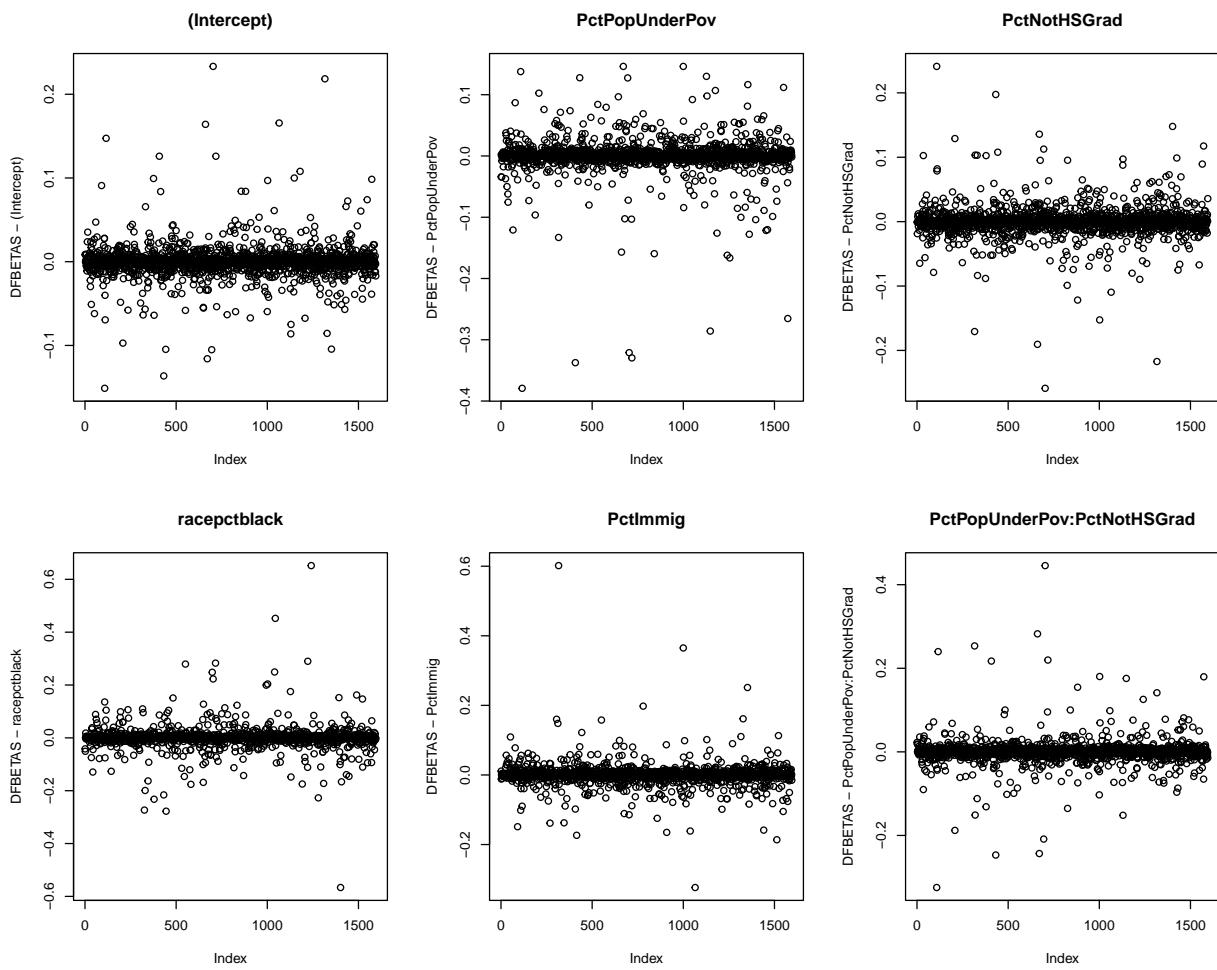
##      obs population stud_residual      leverage      cooks_d      dffits
## 316   316      358548     4.356588 0.031010169 0.10010140  0.7793616
## 1241  1241      41856      7.032657 0.011495577 0.09302365  0.7583950
## 1402  1402      12257     -3.874930 0.028896884 0.07381564 -0.6684311
## 703   703      87492      3.980953 0.021880943 0.05854069  0.5954211
## 1044  1044      394017      4.724200 0.012175272 0.04523935  0.5244785
## 108   108      12822     -2.286852 0.038295628 0.03461609 -0.4563435
## 432   432      18906     -3.545055 0.015136325 0.03195868 -0.4394860
## 1222  1222      49188      5.312197 0.006694863 0.03116590  0.4361177
## 116   116      21265     -1.826202 0.046417616 0.02701677 -0.4029130
## 1000  1000      36118      4.145069 0.009015939 0.02579029  0.3953702

```

```
# Influence plot
plot(leverage, stud_res_final,
      xlab = "Leverage", ylab = "Studentized Residuals",
      main = "Influence Plot",
      cex = sqrt(cooks_d) * 10)
```



```
# DFBETAS plots
par(mfrow = c(2, ceiling(ncol(dfbetas_val)/2)))
for(j in 1:ncol(dfbetas_val)) {
  plot(dfbetas_val[, j],
        ylab = paste("DFBETAS - ", colnames(dfbetas_val)[j]),
        main = colnames(dfbetas_val)[j])
}
```



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

Summary

Dusja multivariate model stuk beter dan univariate model als je kijkt naar de tabel

```
# summary

mse_final <- mean(fit_final$residuals^2)
summary_results <- data.frame(
  Model = c("Simple (PctPopUnderPov only)", "Final Multivariate"),
  R_squared = c(round(summary(fit)$r.squared, 4),
                round(summary(fit_final)$r.squared, 4)),
  Adj_R_squared = c(round(summary(fit)$adj.r.squared, 4),
                    round(summary(fit_final)$adj.r.squared, 4)),
  MSE = c(round(mse_simple, 2), round(mse_final, 2))
)

kable(summary_results, caption = "Comparison of Simple and Final Multivariate Models")
```

Table 2: Comparison of Simple and Final Multivariate Models

Model	R_squared	Adj_R_squared	MSE
Simple (PctPopUnderPov only)	0.2547	0.2542	265053.0
Final Multivariate	0.5187	0.5172	171155.5

References

Becker GS (1968) Crime and Punishment: An Economic Approach. *J Polit Econ* 76: 169–217

References dataset

U. S. Department of Commerce, Bureau of the Census, Census Of Population And Housing 1990 United States: Summary Tape File 1a & 3a (Computer Files),

U.S. Department Of Commerce, Bureau Of The Census Producer, Washington, DC and Inter-university Consortium for Political and Social Research Ann Arbor, Michigan. (1992)

U.S. Department of Justice, Bureau of Justice Statistics, Law Enforcement Management And Administrative Statistics (Computer File) U.S. Department Of Commerce, Bureau Of The Census Producer, Washington, DC and Inter-university Consortium for Political and Social Research Ann Arbor, Michigan. (1992)

U.S. Department of Justice, Federal Bureau of Investigation, Crime in the United States (Computer File) (1995)

Redmond, M. A. and A. Baveja: A Data-Driven Software Tool for Enabling Cooperative Information Sharing Among Police Departments. *European Journal of Operational Research* 141 (2002) 660-678.