

# Data Analysis

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## Load packages

```
library(tidyverse)
```

```
## -- Attaching packages -----
## v tibble  3.0.0          v purrr  0.3.3
## v tidyr   1.0.2.9000     v dplyr  0.8.5
## v readr   1.1.1          v forcats 0.3.0

## -- Conflicts ----- ti
## x lubridate::as.difftime() masks base::as.difftime()
## x lubridate::date()        masks base::date()
## x dplyr::filter()          masks stats::filter()
## x readr::guess_encoding()  masks rvest::guess_encoding()
## x lubridate::intersect()   masks base::intersect()
## x dplyr::lag()              masks stats::lag()
## x purrr::pluck()           masks rvest::pluck()
## x lubridate::setdiff()     masks base::setdiff()
## x lubridate::union()       masks base::union()
```

```
library(knitr)
library(broom)
library(ggplot2)
library(openintro)
library(nnet)
library(patchwork)
library(pROC)
```

```
## Type 'citation("pROC")' for a citation.
##
## Attaching package: 'pROC'
##
## The following objects are masked from 'package:stats':
##
##     cov, smooth, var
```

```
library(plotROC)
```

```
##
## Attaching package: 'plotROC'
##
## The following object is masked from 'package:pROC':
##
##     ggroc
```

```
library(psych)
```

```
##
## Attaching package: 'psych'
```

```

## The following objects are masked from 'package:ggplot2':
##
##      %+%, alpha
library(RColorBrewer) #custom color palettes
#wrap and model spatial data
library(sf)

## Linking to GEOS 3.5.1, GDAL 2.2.2, proj.4 4.9.2
library(sp)
library(spatialreg)

## Loading required package: spData

## To access larger datasets in this package, install the spDataLarge
## package with: `install.packages('spDataLarge',
## repos='https://nowosad.github.io/drat/', type='source')`

##
## Attaching package: 'spData'

## The following object is masked from 'package:openintro':
##
##      house

## Loading required package: Matrix

##
## Attaching package: 'Matrix'

## The following objects are masked from 'package:tidyr':
##
##      expand, pack, unpack
library(spdep)

##
## Attaching package: 'spdep'

## The following objects are masked from 'package:spatialreg':
##
##      GMarginImage, GMerrorsar, HPDinterval.lagImpact,
##      Hausman.test, Jacobian_W, LR.sarlm, LR1.sarlm, LR1.spautolm,
##      LU_prepermute_setup, LU_setup, MCMCsamp, ME, Matrix_J_setup,
##      Matrix_setup, SE_classic_setup, SE_interp_setup,
##      SE_whichMin_setup, SpatialFiltering, Wald1.sarlm, anova.sarlm,
##      as.spam.listw, as_dgRMatrix_listw, as_dsCMatrix_I,
##      as_dsCMatrix_IrW, as_dsTMatrix_listw, bptest.sarlm,
##      can.be.simmed, cheb_setup, coef.gmsar, coef.sarlm,
##      coef.spautolm, coef.stsls, create_WX, deviance.gmsar,
##      deviance.sarlm, deviance.spautolm, deviance.stsls, do_ldet,
##      eigen_pre_setup, eigen_setup, eigenw, errorsarlm,
##      fitted.ME_res, fitted.SFResult, fitted.gmsar, fitted.sarlm,
##      fitted.spautolm, get.ClusterOption, get.VerboseOption,
##      get.ZeroPolicyOption, get.coresOption, get.mcOption,
##      griffith_sone, gstsls, impacts, intImpacts, jacobianSetup,
##      l_max, lagmess, lagsarlm, lexB, leXS, leXW, lmSLX,
##      logLik.sarlm, logLik.spautolm, mcdet_setup, mom_calc,
##      mom_calc_int2, moments_setup, powerWeights, predict.SLX,

```

```
## predict.sarlm, print.ME_res, print.SFResult, print.gmsar,
## print.sarlm, print.sarlm.pred, print.spautolm, print.stsls,
## print.summary.gmsar, print.summary.sarlm,
## print.summary.spautolm, print.summary.stsls, residuals.gmsar,
## residuals.sarlm, residuals.spautolm, residuals.stsls,
## sacsarlm, set.ClusterOption, set.VerboseOption,
## set.ZeroPolicyOption, set.coresOption, set.mcOption,
## similar.listw, spBreg_lag, spam_setup, spam_update_setup,
## spautolm, stsls, subgraph_eigenw, summary.gmsar,
## summary.sarlm, summary.spautolm, summary.stsls, trW,
## vcov.sarlm

library(spData)
library(tibble)
library(dplyr)
```

## Loading and Manipulating the Data

```
gentdata <- read_csv("data/gentdata.csv", col_names = TRUE, col_types = cols())

manual <- read_csv("ImportR.csv", col_names = TRUE, col_types = cols())

manual <- manual %>%
  mutate(black = 100*(black17/total17 - black10/total10)) %>%
  mutate(collegewhite = 100*(collegewhite17/total17 - collegewhite10/total10)) %>%
  mutate(nodiploma = 100*(nodiploma17/total17 - nodiploma10/total10)) %>%
  mutate(highschoolgrad = 100*(highschoolgrad17/total17 - highschoolgrad10/total10)) %>%
  mutate(collegedegree = 100*(collegedegree17/total17 - collegedegree10/total10)) %>%
  mutate(collegedegree = 100*(collegedegree17/total17 - collegedegree10/total10)) %>%
  mutate(early_late = 100*(early_late17/employed17 - early_late10/employed10)) %>%
  mutate(privateschool = 100*(privateschool17/totalpop17 - privateschool10/totalpop10))
```

Mutating new variables to demonstrate change over time:

```
manual <- manual %>%
  mutate(moved17=as.numeric(moved17)) %>%
  mutate(moved10=as.numeric(moved10)) %>%
  mutate(moved = moved17-moved10) %>%
  mutate(homeprice17=as.numeric(homeprice17)) %>%
  mutate(homeprice10=as.numeric(homeprice10)) %>%
  mutate(homeprice_med = (homeprice17 - homeprice10)) %>%
  mutate(income2017=as.numeric(income2017)) %>%
  mutate(income2010=as.numeric(income2010)) %>%
  mutate(income_med = (income2017 - income2010))
```

```
## Warning: NAs introduced by coercion
```

```
## Warning: NAs introduced by coercion
```

```
## Warning: NAs introduced by coercion
```

```
## Warning: NAs introduced by coercion
```

```
## Warning: NAs introduced by coercion
```

```
## Warning: NAs introduced by coercion
```

```
names(manual)[1] <- "geoid"
```

Recoding variables to be numeric:

```
manual <- manual %>%  
  mutate(income_med=as.numeric(income)) %>%  
  mutate(homeprice_med=as.numeric(homeprice)) %>%  
  mutate(collegewhite=as.numeric(collegewhite)) %>%  
  mutate(whitecollar=as.numeric(whitecollar)) %>%  
  mutate(early_late=as.numeric(early_late)) %>%  
  mutate(highschoolgrad=as.numeric(highschoolgrad)) %>%  
  mutate(collegedegree=as.numeric(collegedegree)) %>%  
  mutate(nodiploma=as.numeric(nodiploma)) %>%  
  mutate(black=as.numeric(black)) %>%  
  mutate(privateschool=as.numeric(privateschool))
```

```
## Warning: NAs introduced by coercion
```

```
## Warning: NAs introduced by coercion
```

```
## Warning: NAs introduced by coercion
```

Joining data sets:

```
gent_rural <- gentdata %>%  
  group_by(geoid) %>%  
  summarise(rural)
```

```
manual <- inner_join(manual, gent_rural, copy=T)
```

```
## Joining, by = "geoid"
```

manually imputing the mean value for homeprice

```
mean_homeprice <- manual %>%  
  summarise(mean = mean(homeprice_med, na.rm = T)) %>%  
  pull()
```

```
manual <- manual %>%  
  mutate(homeprice_med = if_else(is.na(homeprice_med), mean_homeprice, homeprice_med))
```

```
mean_income <- manual %>%  
  summarise(mean = mean(income_med, na.rm = T)) %>%  
  pull()
```

```
manual <- manual %>%  
  mutate(income_med = if_else(is.na(income_med), mean_income, income_med))
```

## EXPLORATORY DATA ANALYSIS

#Univariate analysis

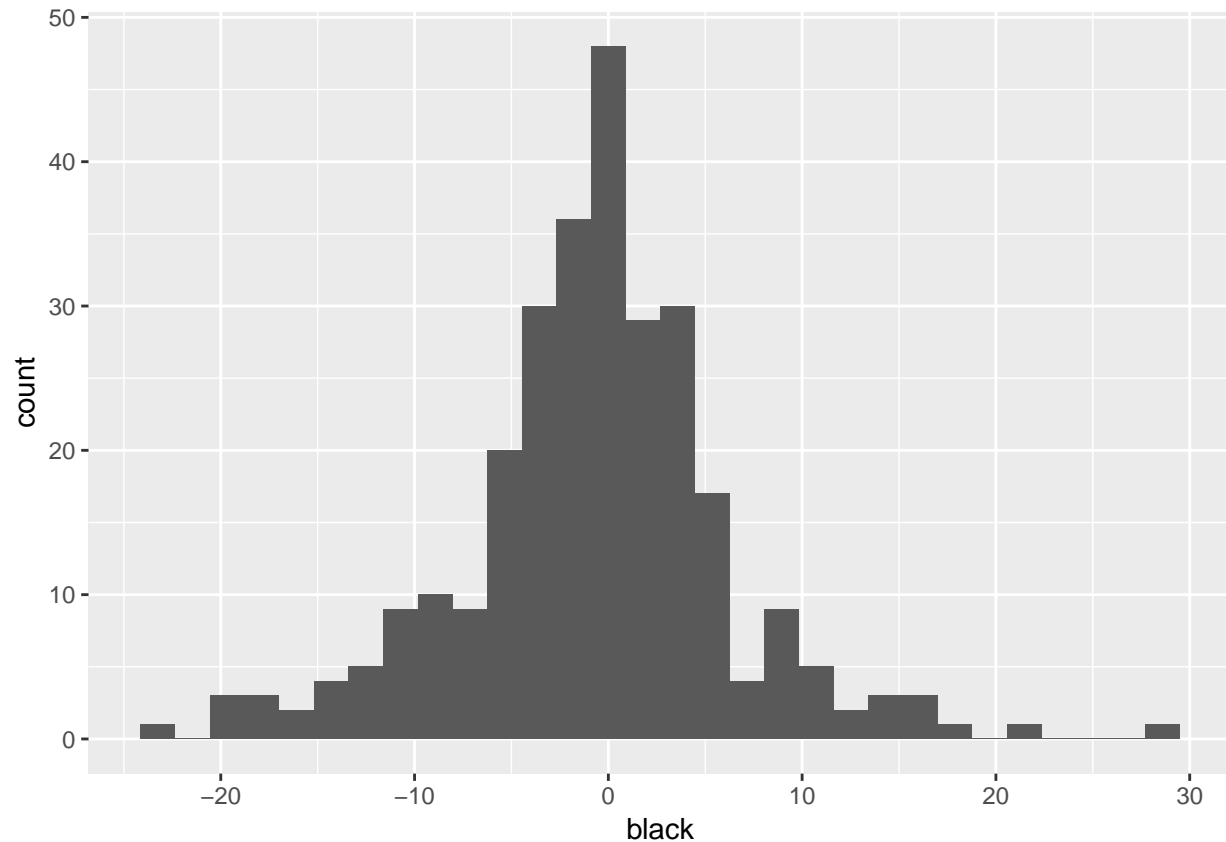
The variable “black” is the change in black population from 2010 to 2017. We will use this variable as our response to predict whether gentrification is occurring in a region in the research triangle.

The distribution of change in Black population:

```
ggplot(data = manual, mapping = aes(x = black)) +  
  geom_histogram()
```

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

```
## Warning: Removed 3 rows containing non-finite values (stat_bin).
```



```
typeof(manual$black)
```

```
## [1] "double"
```

```
(sd(manual$black))
```

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

std deviation is = 6.765474

More Univariate EDA:

```
p1 <- ggplot(data = manual, mapping = aes(x = privateschool)) +  
  geom_histogram()
```

```
p2 <-ggplot(data = manual, mapping = aes(x = collegewhite)) +  
  geom_histogram()
```

```
p3 <-ggplot(data = manual, mapping = aes(x = homeprice_med)) +  
  geom_histogram()
```

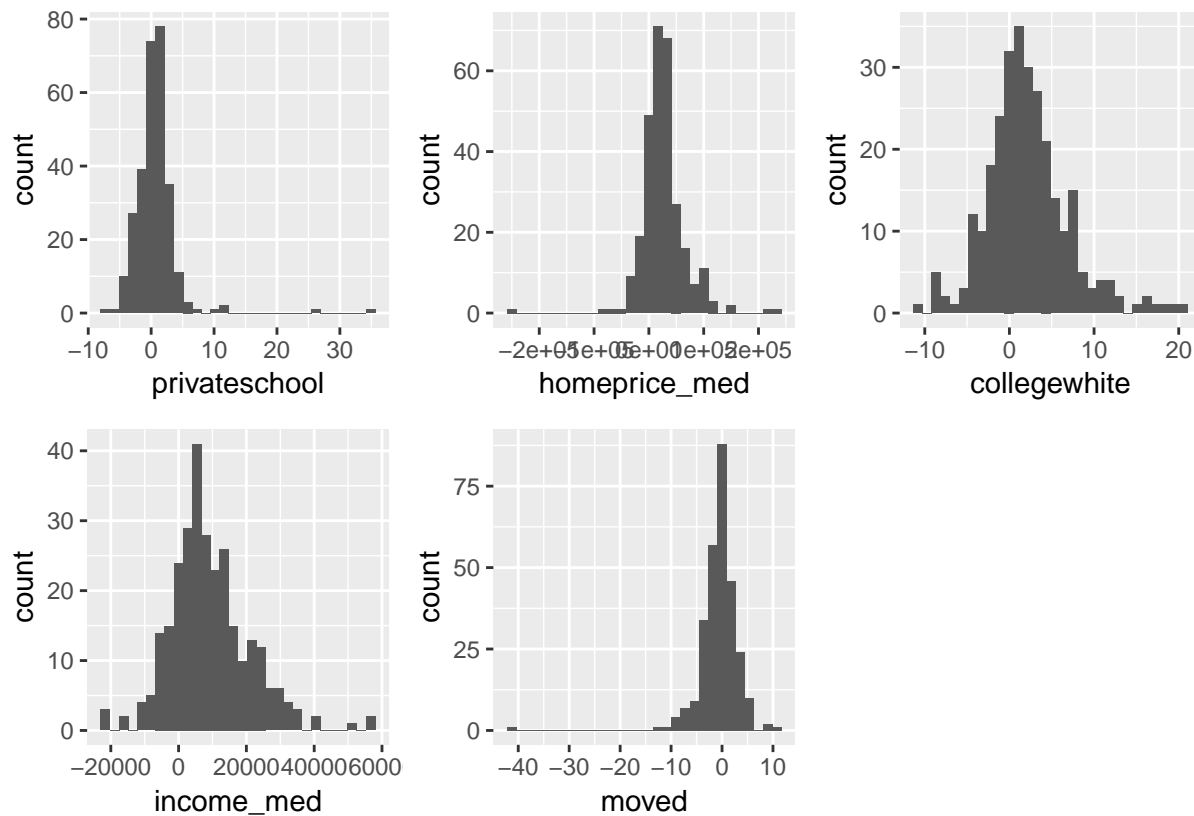
```
p4 <-ggplot(data = manual, mapping = aes(x = income_med)) +
```

```
geom_histogram()

p5 <- ggplot(data = manual, mapping = aes(x = moved)) +
  geom_histogram()

p1 + p3 + p2 + p4 + p5

## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
## Warning: Removed 3 rows containing non-finite values (stat_bin).
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
## Warning: Removed 3 rows containing non-finite values (stat_bin).
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
## Warning: Removed 3 rows containing non-finite values (stat_bin).
```



Bivariate EDA:

```
p6 <- ggplot(data = manual, mapping = aes(x = black, y = privateschool)) +
  geom_boxplot()

p7 <- ggplot(data = manual, mapping = aes(x = black, y = collegewhite)) +
  geom_boxplot()

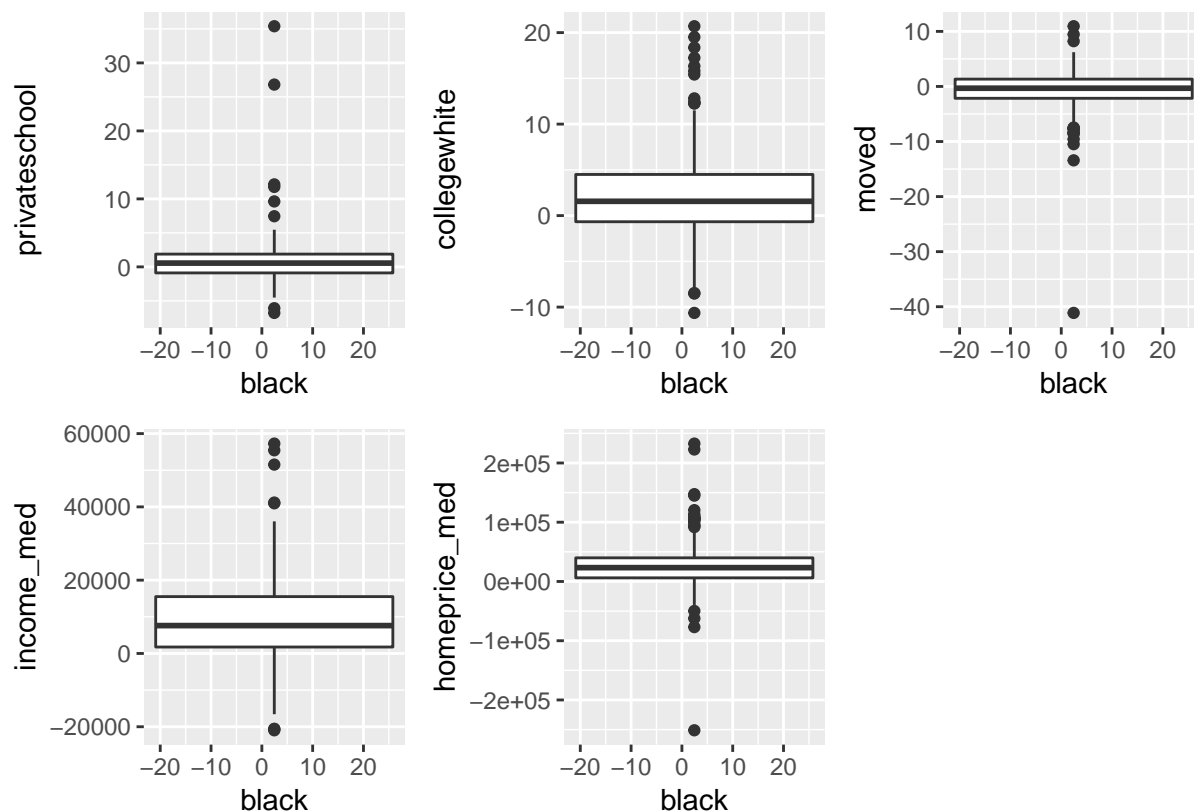
p8 <- ggplot(data = manual, mapping = aes(x = black, y = moved)) +
  geom_boxplot()
```

```
p9 <- ggplot(data = manual, mapping = aes(x = black, y = income_med)) +
  geom_boxplot()

p10 <- ggplot(data = manual, mapping = aes(x = black, y = homeprice_med)) +
  geom_boxplot()

p6 + p7 + p8 + p9 + p10
```

```
## Warning: Continuous x aesthetic -- did you forget aes(group=...)?
## Warning: Removed 3 rows containing missing values (stat_boxplot).
## Warning: Continuous x aesthetic -- did you forget aes(group=...)?
## Warning: Removed 3 rows containing missing values (stat_boxplot).
## Warning: Continuous x aesthetic -- did you forget aes(group=...)?
## Warning: Removed 3 rows containing missing values (stat_boxplot).
## Warning: Continuous x aesthetic -- did you forget aes(group=...)?
## Warning: Removed 3 rows containing missing values (stat_boxplot).
## Warning: Continuous x aesthetic -- did you forget aes(group=...)?
## Warning: Removed 3 rows containing missing values (stat_boxplot).
```



###Part I: Location of Gentrification

In part I, the following research question will be examined:

Where in the Research Triangle (counties including Durham, Wake, Orange and Chatham) is gentrification occurring the most?

Recoding our response variable to “1” if change in black population is  $\leq$  “-6.765” or one standard deviation below the mean and equal to “0” if  $>$  “-6.765” in order visualize and eventually create a logistic model:

```
manual <- manual %>%
  mutate(gent = case_when(black>(-6.765474) ~ 0, black<=(-6.765474) ~ 1))

manual <- manual %>%
  mutate(gent = if_else(is.na(gent), 0, gent))

manual %>%
  count(gent)

## # A tibble: 2 x 2
##   gent     n
##   <dbl> <int>
## 1     0   244
## 2     1    44
typeof(manual$gent)

## [1] "double"

manual

## # A tibble: 288 x 52
##   geoid County total17 total10 black17 black10 `Change in blac~ income2017
##   <chr> <chr>   <int>   <int>   <int>   <int>   <int>   <dbl>
## 1 1400~ Chath~   4557   3784   301    276     25   83750
## 2 1400~ Chath~   4841   4546   123    139    -16   84638
## 3 1400~ Chath~   4334   2342   112    182    -70   87773
## 4 1400~ Chath~   4483   3548   429    514    -85   78897
## 5 1400~ Chath~   7792   7864   803    562    241   64987
## 6 1400~ Chath~   2708   2814   470    509    -39   43594
## 7 1400~ Chath~   5769   5365  1279   1026   253   35952
## 8 1400~ Chath~   5016   5457  1380   1184   196   23848
## 9 1400~ Chath~   3635   4001   254    349   -95   44070
## 10 1400~ Chath~   5720   5811  1105   1135   -30   54914
## # ... with 278 more rows, and 44 more variables: income2010 <dbl>,
## #   income <chr>, collegewhite17 <int>, collegewhite10 <int>,
## #   nodiploma17 <int>, highschoolgrad17 <int>, collegedegree17 <int>,
## #   nodiploma10 <int>, highschoolgrad10 <int>, collegedegree10 <int>,
## #   homeprice17 <dbl>, homeprice10 <dbl>, homeprice <chr>,
## #   employed17 <int>, white1_17 <chr>, white2_17 <chr>, white3_17 <chr>,
## #   whitecollar17 <chr>, employed10 <int>, white1_10 <chr>,
## #   white2_10 <chr>, white3_10 <chr>, whitecollar10 <chr>,
## #   whitecollar <dbl>, early_late17 <int>, early_late10 <int>,
## #   privateschool17 <int>, totalpop17 <int>, privateschool10 <int>,
## #   totalpop10 <int>, moved17 <dbl>, moved10 <dbl>, black <dbl>,
## #   collegewhite <dbl>, nodiploma <dbl>, highschoolgrad <dbl>,
## #   collegedegree <dbl>, early_late <dbl>, privateschool <dbl>,
## #   moved <dbl>, homeprice_med <dbl>, income_med <dbl>, rural <chr>,
## #   gent <dbl>
```

Reading in spatial data



```
# read the shapefile
shape <- read_sf(dsn = "data", layer = "triangletracts")

## Error: All columns in a tibble must be vectors.
## x Column `geometry` is a `sfc_POLYGON/sfc` object.
# convert RegionID to numeric before we join and drop some columns that we don't need

shape <- shape %>%
  mutate(geoid = as.character(AFFGEOID))
```

```
## Error in eval(lhs, parent, parent): object 'shape' not found
# merge keeping only those in both data sets
```

```
merged <- inner_join(shape, manual, by = "geoid")
```

```
## Error in inner_join(shape, manual, by = "geoid"): object 'shape' not found
Plotting research triangle area:
```

```
ggplot(data = merged) +
  geom_sf()
```

```
## Error in ggplot(data = merged): object 'merged' not found
```

Plotting research triangle area by which regions have experienced gentrification:

```
ggplot(data = merged, aes(fill = gent)) +
  geom_sf() +
  labs(title = "Research Triangle",
       subtitle = "Gentrification by census tract") +
  theme_void() +
  scale_fill_distiller(palette = 'RdBu', guide = "legend")
```

```
## Error in ggplot(data = merged, aes(fill = gent)): object 'merged' not found
```

### Part 2: Factors Associated with Gentrification

In part 2, the following research question will be examined:

What factors are associated with and the strongest predictors of the gentrification of these areas.

We already determined a model using aic and drop in deviance tests ....

### Using Logistic Regression

Creating the logistic model using mutated variable “gent” as our response variable. :

```
model <- glm(gent ~ collegewhite + whitecollar + privateschool + nodiploma + highschoolgrad + collegedept,
             data = manual, family="binomial")

tidy(model, conf.int = TRUE) %>%
  kable(format = "markdown", digits = 5)
```

term	estimate	std.error	statistic	p.value	conf.low	conf.high
(Intercept)	-3.00216	1.76126	-1.70455	0.08828	-6.50546	0.42759
collegewhite	0.14487	0.06063	2.38927	0.01688	0.02889	0.26801
whitecollar	-0.05600	0.03098	-1.80772	0.07065	-0.11837	0.00337
privateschool	0.01855	0.05029	0.36882	0.71226	-0.09732	0.10785
nodiploma	0.12286	0.06408	1.91736	0.05519	0.00077	0.25299

term	estimate	std.error	statistic	p.value	conf.low	conf.high
highschoolgrad	0.05705	0.04424	1.28955	0.19721	-0.02836	0.14590
collegedegree	-0.03647	0.06220	-0.58638	0.55762	-0.16062	0.08438
income_med	-0.00002	0.00002	-0.86680	0.38605	-0.00005	0.00002
homeprice_med	0.00001	0.00000	2.44891	0.01433	0.00000	0.00002
early_late	-0.00940	0.01852	-0.50727	0.61197	-0.04578	0.02715
moved	0.00344	0.04475	0.07697	0.93865	-0.07898	0.10019

Using backward selection to find the optimal model:

```
model_aic <- step(model, direction = "backward", conf.int=T)
```

```
## Start:  AIC=241.3
## gent ~ collegewhite + whitecollar + privateschool + nodiploma +
##       highschoolgrad + collegedegree + income_med + homeprice_med +
##       early_late + moved
##
##               Df Deviance    AIC
## - moved          1    219.31 239.31
## - privateschool   1    219.43 239.43
## - early_late      1    219.56 239.56
## - collegedegree   1    219.65 239.65
## - income_med      1    220.07 240.07
## - highschoolgrad  1    221.00 241.00
## <none>              219.30 241.30
## - whitecollar     1    222.71 242.71
## - nodiploma        1    223.19 243.19
## - collegewhite     1    225.34 245.34
## - homeprice_med    1    225.64 245.64
##
## Step:  AIC=239.31
## gent ~ collegewhite + whitecollar + privateschool + nodiploma +
##       highschoolgrad + collegedegree + income_med + homeprice_med +
##       early_late
##
##               Df Deviance    AIC
## - privateschool   1    219.44 237.44
## - early_late      1    219.56 237.56
## - collegedegree   1    219.66 237.66
## - income_med      1    220.08 238.08
## - highschoolgrad  1    221.00 239.00
## <none>              219.31 239.31
## - whitecollar     1    222.73 240.73
## - nodiploma        1    223.19 241.19
## - collegewhite     1    225.38 243.38
## - homeprice_med    1    225.65 243.65
##
## Step:  AIC=237.44
## gent ~ collegewhite + whitecollar + nodiploma + highschoolgrad +
##       collegedegree + income_med + homeprice_med + early_late
##
##               Df Deviance    AIC
## - early_late      1    219.66 235.66
```

```

## - collegedegree 1 219.76 235.76
## - income_med 1 220.21 236.21
## - highschoolgrad 1 221.11 237.11
## <none> 219.44 237.44
## - whitecollar 1 222.86 238.86
## - nodiploma 1 223.25 239.25
## - collegewhite 1 225.44 241.44
## - homeprice_med 1 225.82 241.82
##
## Step: AIC=235.66
## gent ~ collegewhite + whitecollar + nodiploma + highschoolgrad +
## collegedegree + income_med + homeprice_med
##
## Df Deviance AIC
## - collegedegree 1 219.98 233.98
## - income_med 1 220.52 234.52
## - highschoolgrad 1 221.25 235.25
## <none> 219.66 235.66
## - whitecollar 1 223.04 237.04
## - nodiploma 1 223.43 237.43
## - collegewhite 1 225.57 239.57
## - homeprice_med 1 226.71 240.71
##
## Step: AIC=233.98
## gent ~ collegewhite + whitecollar + nodiploma + highschoolgrad +
## income_med + homeprice_med
##
## Df Deviance AIC
## - income_med 1 220.93 232.93
## <none> 219.98 233.98
## - highschoolgrad 1 222.65 234.65
## - whitecollar 1 223.39 235.39
## - nodiploma 1 224.51 236.51
## - homeprice_med 1 226.87 238.87
## - collegewhite 1 228.82 240.82
##
## Step: AIC=232.93
## gent ~ collegewhite + whitecollar + nodiploma + highschoolgrad +
## homeprice_med
##
## Df Deviance AIC
## <none> 220.93 232.93
## - highschoolgrad 1 224.01 234.01
## - whitecollar 1 224.36 234.36
## - nodiploma 1 225.70 235.70
## - homeprice_med 1 226.98 236.98
## - collegewhite 1 228.92 238.92

```

```

tidy(model_aic, conf.int = TRUE) %>%
  kable(format = "markdown", digits = 5)

```

term	estimate	std.error	statistic	p.value	conf.low	conf.high
(Intercept)	-2.23080	0.24515	-9.09985	0.00000	-2.74176	-1.77646
collegewhite	0.10841	0.03943	2.74954	0.00597	0.03290	0.18858

term	estimate	std.error	statistic	p.value	conf.low	conf.high
whitecollar	-0.05692	0.03137	-1.81461	0.06958	-0.11996	0.00321
nodiploma	0.13470	0.06339	2.12476	0.03361	0.01355	0.26289
highschoolgrad	0.06938	0.04021	1.72554	0.08443	-0.00792	0.15022
homeprice_med	0.00001	0.00000	2.37718	0.01745	0.00000	0.00002

Creating a full model to determine if we should add “rural” to the model:

```
model_aic_full <- glm(gent ~ collegewhite + whitecollar + nodiploma + highschoolgrad + homeprice_med + rural, data = gent_data)
tidy(model_aic_full)
```

```
## # A tibble: 7 x 5
##   term          estimate std.error statistic  p.value
##   <chr>          <dbl>    <dbl>    <dbl>    <dbl>
## 1 (Intercept)   -2.88      0.450     -6.39 1.64e-10
## 2 collegewhite  0.0939    0.0400      2.35 1.88e- 2
## 3 whitecollar  -0.0546    0.0318     -1.72 8.61e- 2
## 4 nodiploma     0.118     0.0631      1.87 6.16e- 2
## 5 highschoolgrad 0.0606    0.0406      1.49 1.35e- 1
## 6 homeprice_med 0.0000108 0.00000468  2.31 2.10e- 2
## 7 ruralUrban    0.891     0.476      1.87 6.12e- 2
```

Drop in deviance test:

```
(dev_m <- glance(model_aic)$deviance)

## [1] 220.9284

(dev_full <- glance(model_aic_full)$deviance)

## [1] 216.9145

(test_stat <- dev_m - dev_full)
```

```
## [1] 4.013908
```

p-value:

```
1- pchisq(test_stat, 1)
```

```
## [1] 0.04512643
```

Since the chisq p-value for adding “Rural” to the model is less than .05, we reject the null hypothesis that “Rural” is not a significant predictor of whether or not a region has experienced gentrification.

Therefore we will continue with this full model for the remained of our analysis.

### Assumptions

In order to use the full model with the predictor variables collegewhite, whitecollar, nodiploma, highschoolgrad, homeprice\_med, and rural, we must first test how well this model satisfies assumptions.

For testing linearity, we will augment the model with predicted probabilities and residuals in order to examine binned residual plots for predicted probability and numeric variables.

```
model_aug <- augment(model_aic_full, type.predict = "response", type.residuals = "response")
```

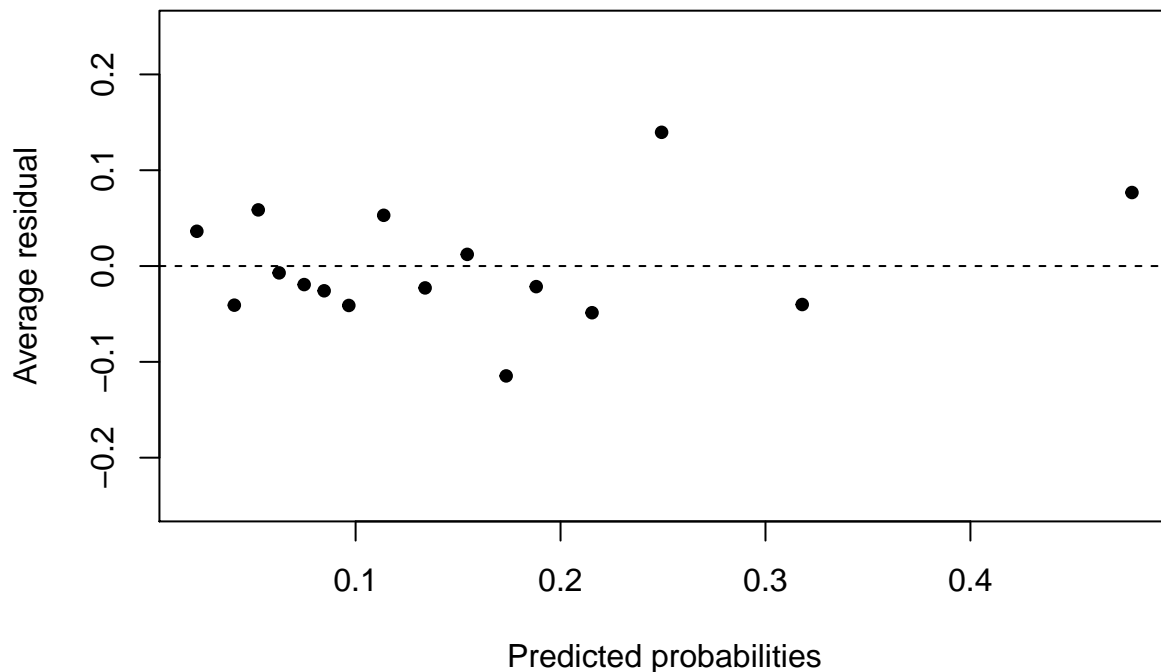
```
model_aug
```

```
## # A tibble: 285 x 15
```

```
##      .rownames  gent collegewhite whitecollar nodiploma highschoolgrad
##      <chr>      <dbl>      <dbl>      <dbl>      <dbl>      <dbl>
##  1 1          0      -8.46      -16.4      7.20      -4.89
##  2 2          0       2.96       3.2     -0.640     -0.658
##  3 3          0     -0.735       4.3     0.807     -1.44
##  4 4          0     -4.08       6.6     0.0800    -1.86
##  5 5          0       2.31      -2.8     1.16     -4.08
##  6 6          0     -1.25       0.5    -0.388     -4.55
##  7 7          0     -0.224       4.3    -7.19      5.87
##  8 8          0       2.17      18.4    -5.19     11.3
##  9 9          0     -1.92      -5.6     2.05     0.321
## 10 10         0       5.44       6.9    -0.837     0.687
## # ... with 275 more rows, and 9 more variables: homeprice_med <dbl>,
## #   rural <chr>, .fitted <dbl>, .se.fit <dbl>, .resid <dbl>, .hat <dbl>,
## #   .sigma <dbl>, .cooksad <dbl>, .std.resid <dbl>
```

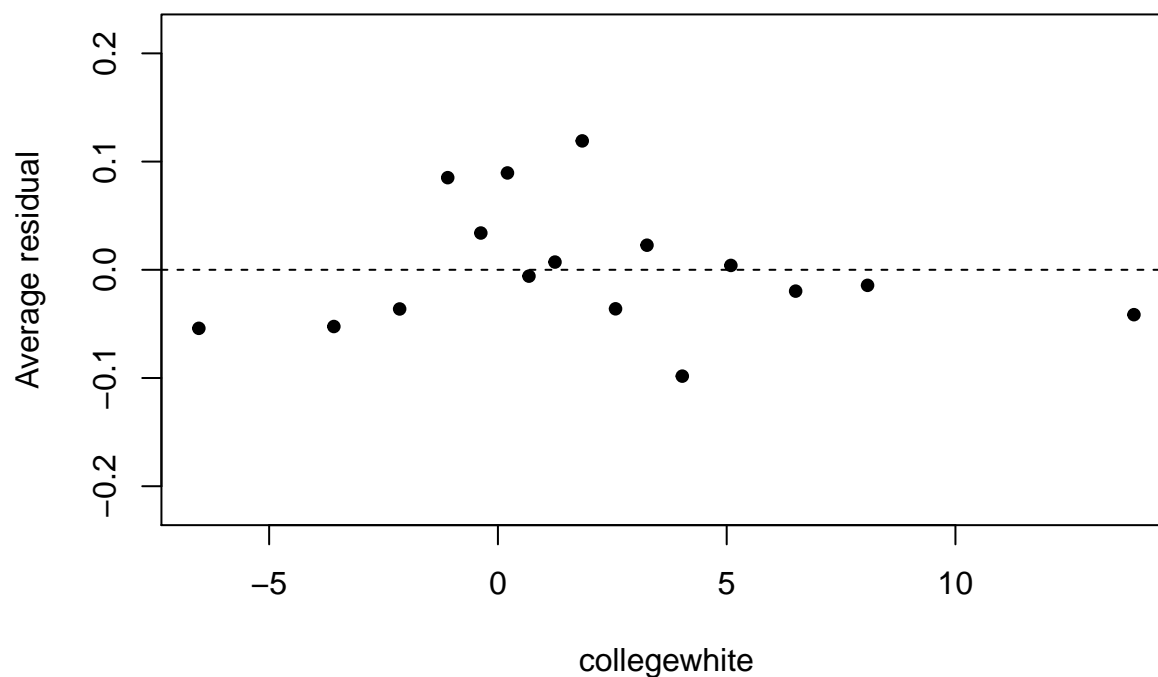
```
arm::binnedplot(x = model_aug$.fitted,
  y = model_aug$.resid,
  col.int = FALSE,
  xlab = "Predicted probabilities",
  main = "Binned Residual vs. Predicted Probability")
```

## Binned Residual vs. Predicted Probability



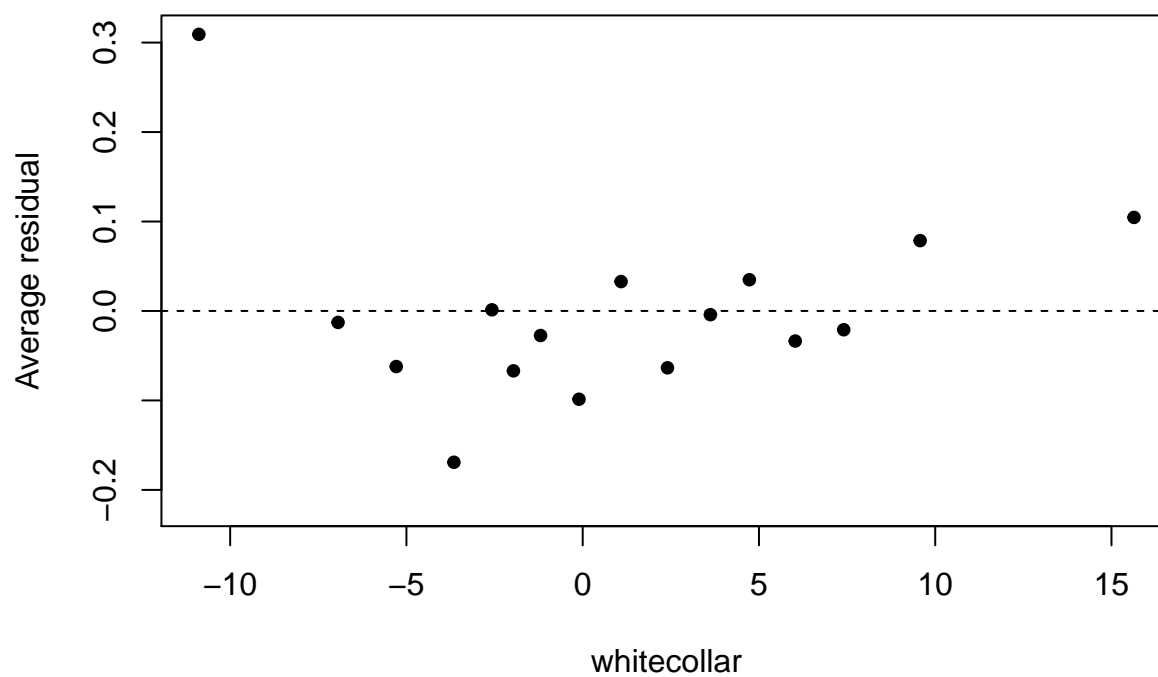
```
arm::binnedplot(x = model_aug$collegewhite,
  y = model_aug$.resid,
  col.int = FALSE,
  xlab = "collegewhite",
  main = "Binned Residual vs. collegewhite")
```

**Binned Residual vs. collegewhite**



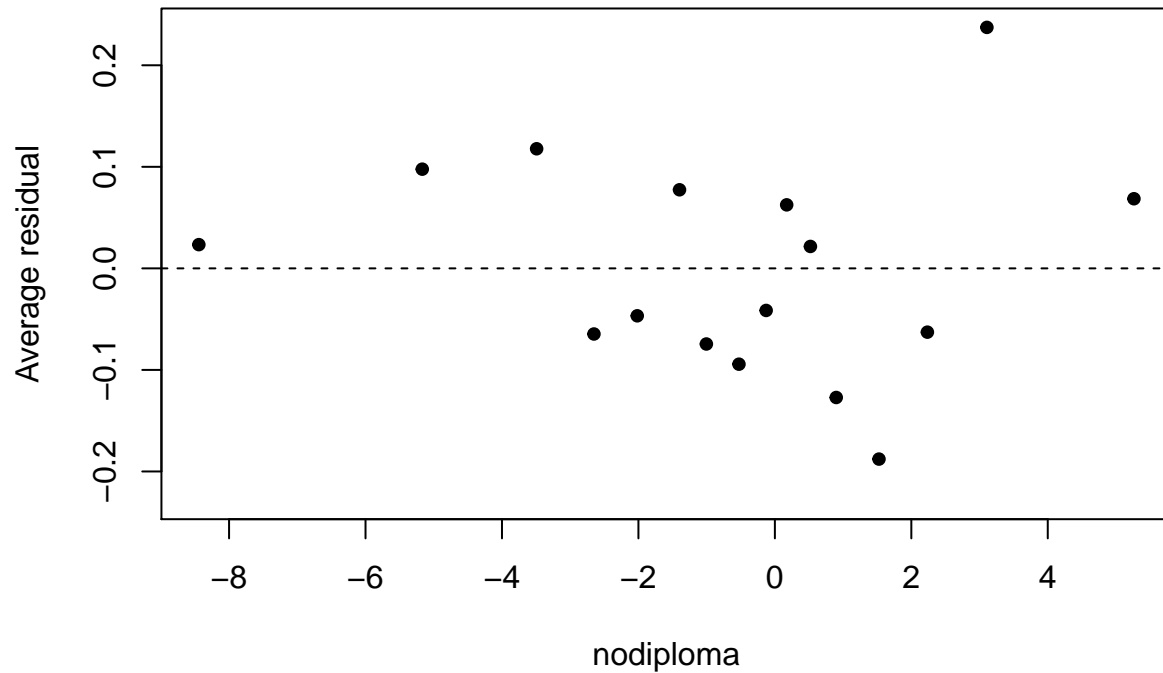
```
arm::binnedplot(x = model_aug$whitecollar,  
  y = model_aug$.resid,  
  col.int = FALSE,  
  xlab = "whitecollar",  
  main = "Binned Residual vs. whitecollar")
```

**Binned Residual vs. whitecollar**



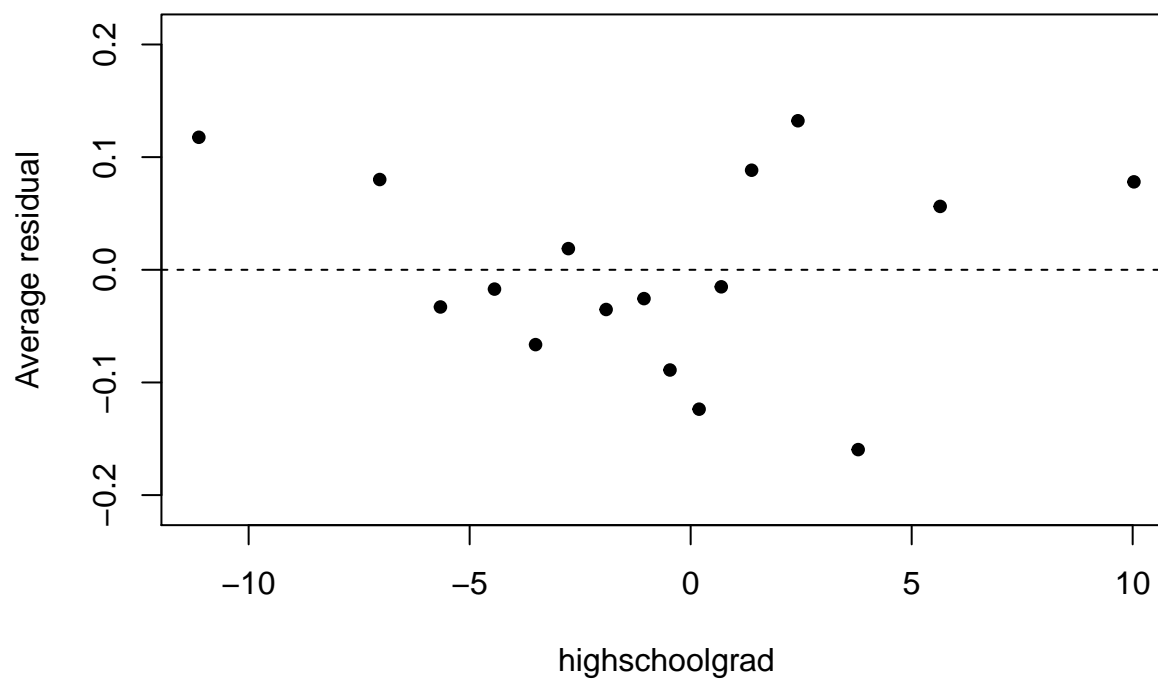
```
arm::binnedplot(x = model_aug$nodiploma,
  y = model_aug$.resid,
  col.int = FALSE,
  xlab = "nodiploma",
  main = "Binned Residual vs. nodiploma")
```

### Binned Residual vs. nodiploma



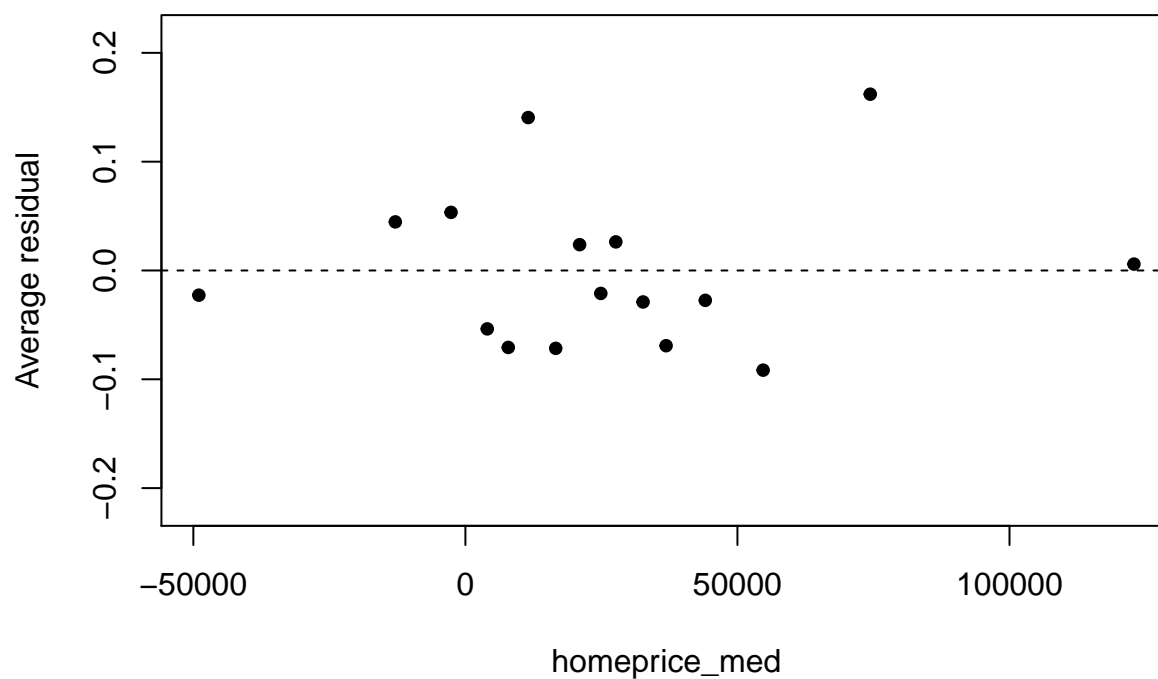
```
arm::binnedplot(x = model_aug$highschoolgrad,
  y = model_aug$.resid,
  col.int = FALSE,
  xlab = "highschoolgrad",
  main = "Binned Residual vs. highschoolgrad")
```

**Binned Residual vs. highschoolgrad**



```
arm::binnedplot(x = model_aug$homeprice_med,  
  y = model_aug$.resid,  
  col.int = FALSE,  
  xlab = "homeprice_med",  
  main = "Binned Residual vs. homeprice_med")
```

**Binned Residual vs. homeprice\_med**





```
model_aug %>%
  group_by(rural) %>%
  summarise(mean_resid = mean(.resid))
```

```
## # A tibble: 2 x 2
##   rural mean_resid
##   <chr>      <dbl>
## 1 Rural  -5.38e-11
## 2 Urban  -4.71e-12
```

The linearity assumption is satisfied. The binned residuals vs. predicted probability plot shows irregularity with a very slight clustering of residual values below 0.0. The binned residuals vs. collegewhite plot shows irregularity. The binned residuals vs. whitecollar plot shows irregularity, with a slight clustering of residual values below 0.0 and a slight increase in residual values as you move right. The binned residuals vs. nodiploma, binned residuals vs. highschoolgrad, and binned residuals vs. homeprice\_med show complete irregularity. For the predictor variable rural, which has two categories rural and urban, both mean residuals are very close to zero. There is no strong indication of nonlinearity; therefore, we can assume that there is a linear relationship between log(gent) and the predictor variables.

To discuss randomness and independence, we must go back to the source of our data. All of the data we are using is sourced from the Census Bureau's annual American Community Survey and official North Carolina demographic data. According to the census sampling techniques and methodology, we can reasonably assume that randomness and independence are satisfied. Read more here: <https://www.census.gov/programs-surveys/sipp/methodology.html>

## Interpreting Model Coefficients

Now that we've confirmed that it satisfies assumptions, let's take a look at our chosen logistic model again:

```
tidy(model_aic_full, conf.int = TRUE, exponentiate = FALSE) %>%
  kable(digits = 3, format = "markdown")
```

term	estimate	std.error	statistic	p.value	conf.low	conf.high
(Intercept)	-2.877	0.450	-6.392	0.000	-3.867	-2.076
collegewhite	0.094	0.040	2.349	0.019	0.017	0.175
whitecollar	-0.055	0.032	-1.716	0.086	-0.119	0.006
nodiploma	0.118	0.063	1.869	0.062	-0.002	0.246
highschoolgrad	0.061	0.041	1.494	0.135	-0.018	0.142
homeprice_med	0.000	0.000	2.309	0.021	0.000	0.000
ruralUrban	0.891	0.476	1.872	0.061	0.018	1.915

We would like to discuss the variables that have the most impact on the response variable gent. Therefore, we will discuss variables with p-values of  $<0.05$ . The variable collegewhite seems to have a reliably strong impact on gent: holding all other variables constant, a unit change in collewhite causes the odds of gentrification are expected to multiply by a factor of  $\exp(0.089) = 1.093$ . However, this impact is not as strong as that of the rural variable. According to the model coefficient for the term ruralUrban, holding all other variables constant, the odds of gentrification for an urban area is expected to be 2.55 that of a rural locale. We would like to suggest that the change in college-educated whites in a county and urban character likely greatly impact “gentrification” as we have classified it (a significant decrease in black population).