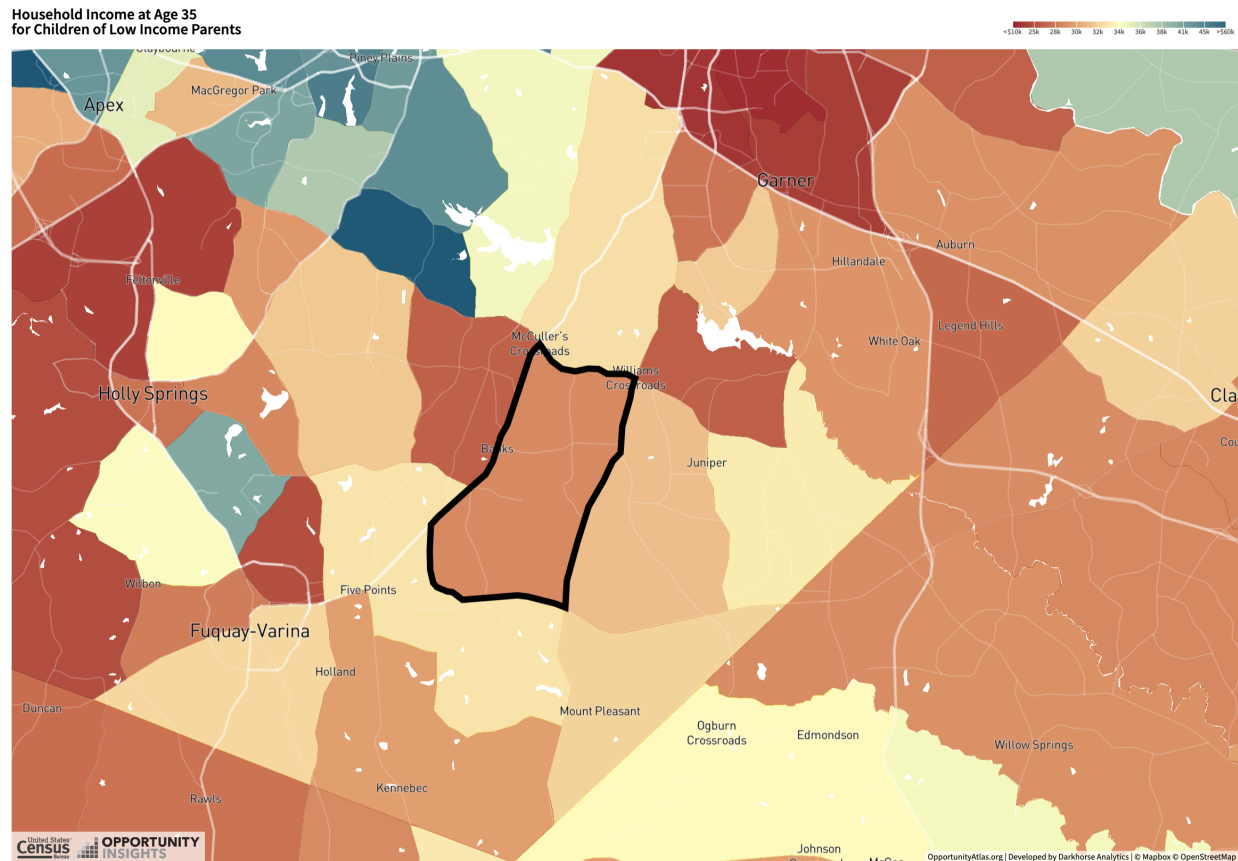


Project_Pt1

2023-02-25

Question 1 My name is Jose and I am from Willow Spring, North Carolina. My community is on the southern edge of Wake County. The map below shows adult household income for children born to low income parents. Compared to similar children nationwide, my neighborhood has below average outcomes.



Question 2: Summary statistics of all variables, including recording all missing values.
summary(atlas)

```
##      state      county      tract      tract_name
## Min.   : 1.00   Min.   : 1.00   Min.   : 100   Length:73199
## 1st Qu.:13.00   1st Qu.: 29.00   1st Qu.: 10300  Class :character
## Median :28.00   Median : 63.00   Median : 44702  Mode  :character
## Mean   :28.18   Mean   : 85.95   Mean   :255238
## 3rd Qu.:42.00   3rd Qu.:109.00   3rd Qu.:458650
## Max.   :72.00   Max.   :840.00   Max.   :989200
##
##      cz      czname      kfr_pooled_pooled_p25 kfr_natam_pooled_p25
## Min.   : 100   Length:73199   Min.   : -3.286   Min.   : -9.99
```

```

## 1st Qu.:11600   Class :character   1st Qu.: 38.070       1st Qu.: 27.80
## Median :19902   Mode  :character   Median : 42.520       Median : 33.41
## Mean    :21974                                     Mean    : 42.858       Mean    : 34.45
## 3rd Qu.:32000                                     3rd Qu.: 47.350       3rd Qu.: 40.08
## Max.    :90000                                     Max.    :103.349       Max.    :107.68
##                                     NA's    :1189          NA's    :71464
## kfr_asian_pooled_p25 kfr_black_pooled_p25 kfr_hisp_pooled_p25
## Min.    :-13.32   Min.    :-48.47       Min.    :-23.97
## 1st Qu.: 51.50    1st Qu.: 29.97       1st Qu.: 39.11
## Median : 57.96    Median : 33.03       Median : 43.27
## Mean    : 57.98    Mean    : 33.99       Mean    : 43.70
## 3rd Qu.: 64.32    3rd Qu.: 37.01       3rd Qu.: 47.65
## Max.    :113.61    Max.    : 99.52       Max.    :125.98
## NA's    :57759     NA's    :39111       NA's    :35581
## kfr_white_pooled_p25 kir_pooled_female_p25 kir_pooled_male_p25
## Min.    :-13.92   Min.    :-42.61       Min.    :-21.42
## 1st Qu.: 41.43    1st Qu.: 37.38       1st Qu.: 42.47
## Median : 45.74    Median : 40.98       Median : 46.81
## Mean    : 46.30    Mean    : 41.86       Mean    : 47.10
## 3rd Qu.: 50.72    3rd Qu.: 45.56       3rd Qu.: 51.44
## Max.    :103.18    Max.    : 87.90       Max.    :105.74
## NA's    :5221     NA's    :1554        NA's    :1512
## kir_natam_female_p25 kir_asian_female_p25 kir_black_female_p25
## Min.    :17.21    Min.    :-43.99       Min.    : 4.04
## 1st Qu.:30.80     1st Qu.: 51.04       1st Qu.:38.09
## Median :34.69     Median : 57.60       Median :41.35
## Mean    :35.12     Mean    : 57.49       Mean    :42.02
## 3rd Qu.:38.91     3rd Qu.: 63.87       3rd Qu.:45.32
## Max.    :76.50     Max.    :166.61       Max.    :89.37
## NA's    :72350     NA's    :65467       NA's    :47861
## kir_hisp_female_p25 kir_white_female_p25 kir_natam_male_p25 kir_asian_male_p25
## Min.    :-15.03   Min.    :-74.19       Min.    :15.66       Min.    :-124.87
## 1st Qu.: 38.64    1st Qu.: 36.45       1st Qu.:32.85       1st Qu.: 52.81
## Median : 42.74    Median : 40.83       Median :38.54       Median : 58.69
## Mean    : 43.12    Mean    : 41.83       Mean    :39.21       Mean    : 58.86
## 3rd Qu.: 47.20    3rd Qu.: 46.19       3rd Qu.:44.97       3rd Qu.: 65.02
## Max.    : 92.85    Max.    :109.12       Max.    :71.51       Max.    :114.93
## NA's    :47319     NA's    :7983        NA's    :72347       NA's    :65112
## kir_black_male_p25 kir_hisp_male_p25 kir_white_male_p25 jail_pooled_pooled_p25
## Min.    :-20.40   Min.    :-24.03       Min.    :-100.99     Min.    :-0.2145
## 1st Qu.: 35.17    1st Qu.: 45.23       1st Qu.: 44.82       1st Qu.: 0.0070
## Median : 39.05    Median : 49.62       Median : 49.38       Median : 0.0180
## Mean    : 39.57    Mean    : 49.80       Mean    : 49.84       Mean    : 0.0216
## 3rd Qu.: 43.42    3rd Qu.: 54.04       3rd Qu.: 54.48       3rd Qu.: 0.0325
## Max.    : 85.17    Max.    :119.73       Max.    :111.89       Max.    : 0.6909
## NA's    :48140     NA's    :47796       NA's    :7683        NA's    :1322
## jail_natam_pooled_p25 jail_asian_pooled_p25 jail_black_pooled_p25
## Min.    :-0.11    Min.    :-0.82       Min.    :-0.55
## 1st Qu.: 0.00     1st Qu.: -0.02       1st Qu.: 0.02
## Median : 0.03     Median : 0.00       Median : 0.05
## Mean    : 0.04     Mean    : 0.00       Mean    : 0.05
## 3rd Qu.: 0.06     3rd Qu.: 0.02       3rd Qu.: 0.08
## Max.    : 0.43     Max.    : 0.79       Max.    : 0.56
## NA's    :71790     NA's    :59738       NA's    :42014

```

```

## jail_hisp_pooled_p25 jail_white_pooled_p25 jail_pooled_female_p25
## Min. :-0.40 Min. :-0.636 Min. :-0.5948
## 1st Qu.: 0.00 1st Qu.: 0.002 1st Qu.: -0.0036
## Median : 0.01 Median : 0.013 Median : 0.0029
## Mean : 0.02 Mean : 0.017 Mean : 0.0038
## 3rd Qu.: 0.03 3rd Qu.: 0.027 3rd Qu.: 0.0105
## Max. : 0.83 Max. : 0.820 Max. : 0.3893
## NA's :38468 NA's :5816 NA's :1723
## jail_pooled_male_p25 jail_natam_female_p25 jail_asian_female_p25
## Min. :-0.5659 Min. :-0.11 Min. :-0.46
## 1st Qu.: 0.0123 1st Qu.: -0.01 1st Qu.: -0.02
## Median : 0.0333 Median : 0.00 Median : 0.00
## Mean : 0.0409 Mean : 0.01 Mean : 0.00
## 3rd Qu.: 0.0618 3rd Qu.: 0.02 3rd Qu.: 0.02
## Max. : 1.1290 Max. : 0.16 Max. : 0.34
## NA's :1725 NA's :72505 NA's :66571
## jail_black_female_p25 jail_hisp_female_p25 jail_white_female_p25
## Min. :-0.16 Min. :-0.19 Min. :-1.300
## 1st Qu.: -0.01 1st Qu.: -0.01 1st Qu.: -0.008
## Median : 0.00 Median : 0.00 Median : 0.003
## Mean : 0.01 Mean : 0.00 Mean : 0.004
## 3rd Qu.: 0.02 3rd Qu.: 0.01 3rd Qu.: 0.015
## Max. : 0.32 Max. : 0.31 Max. : 1.355
## NA's :49929 NA's :49276 NA's :8717
## jail_natam_male_p25 jail_asian_male_p25 jail_black_male_p25 jail_hisp_male_p25
## Min. :-0.06 Min. :-2.04 Min. :-0.45 Min. :-1.27
## 1st Qu.: 0.01 1st Qu.: -0.02 1st Qu.: 0.06 1st Qu.: 0.00
## Median : 0.05 Median : 0.00 Median : 0.10 Median : 0.02
## Mean : 0.06 Mean : 0.01 Mean : 0.11 Mean : 0.03
## 3rd Qu.: 0.10 3rd Qu.: 0.03 3rd Qu.: 0.15 3rd Qu.: 0.05
## Max. : 0.42 Max. : 1.77 Max. : 0.71 Max. : 0.84
## NA's :72572 NA's :66368 NA's :51527 NA's :50725
## jail_white_male_p25 HOLC_A HOLC_B HOLC_C
## Min. :-1.044 Min. :0.00 Min. :0.00 Min. :0.00
## 1st Qu.: 0.004 1st Qu.:0.00 1st Qu.:0.00 1st Qu.:0.00
## Median : 0.023 Median :0.00 Median :0.00 Median :0.17
## Mean : 0.029 Mean :0.03 Mean :0.13 Mean :0.33
## 3rd Qu.: 0.048 3rd Qu.:0.00 3rd Qu.:0.11 3rd Qu.:0.62
## Max. : 1.212 Max. :1.00 Max. :1.00 Max. :1.00
## NA's :8623 NA's :63923 NA's :63923 NA's :63923
## HOLC_D pm25_1982 pm25_1990 pm25_2000
## Min. :0.00 Min. : 1.97 Min. : 1.501 Min. : 1.64
## 1st Qu.:0.00 1st Qu.:16.68 1st Qu.:13.494 1st Qu.:10.21
## Median :0.00 Median :20.85 Median :17.094 Median :12.71
## Mean :0.23 Mean :20.41 Mean :16.844 Mean :12.50
## 3rd Qu.:0.38 3rd Qu.:24.28 3rd Qu.:20.106 3rd Qu.:14.79
## Max. :1.00 Max. :35.92 Max. :32.744 Max. :25.37
## NA's :63923 NA's :1296 NA's :1296 NA's :1296
## pm25_2010 vegetation extreme_heat developed
## Min. : 1.531 Min. :-0.6337 Min. :-14.2736 Min. :0.0000
## 1st Qu.: 7.887 1st Qu.: -0.1392 1st Qu.: 0.2996 1st Qu.:0.1233
## Median : 9.506 Median :-0.0695 Median : 2.6096 Median :0.6734
## Mean : 9.286 Mean :-0.0838 Mean : 2.6758 Mean :0.5751
## 3rd Qu.:10.799 3rd Qu.: -0.0119 3rd Qu.: 5.2012 3rd Qu.:0.9983

```

## Max. :16.958	Max. : 0.3957	Max. : 14.6032	Max. :1.0000
## NA's :1296	NA's :2531	NA's :2531	NA's :2531
## hhinc_mean2000	mean_commutetime2000	frac_coll_plus2000	frac_coll_plus2010
## Min. : 7240	Min. : 2.50	Min. :0.0000	Min. :0.0000
## 1st Qu.: 57377	1st Qu.:22.04	1st Qu.:0.1108	1st Qu.:0.1310
## Median : 71721	Median :26.17	Median :0.1870	Median :0.2186
## Mean : 80335	Mean :26.95	Mean :0.2378	Mean :0.2694
## 3rd Qu.: 94675	3rd Qu.:31.05	3rd Qu.:0.3254	3rd Qu.:0.3709
## Max. :330042	Max. :80.03	Max. :1.0000	Max. :1.0000
## NA's :893	NA's :882	NA's :852	NA's :202
## foreign_share2010	med_hhinc1990	med_hhinc2016	popdensity2000
## Min. :0.0000	Min. : 4999	Min. : 3250	Min. : 0.00
## 1st Qu.:0.0230	1st Qu.: 22261	1st Qu.: 38809	1st Qu.: 97.78
## Median :0.0671	Median : 29810	Median : 52333	Median : 760.47
## Mean :0.1209	Mean : 32179	Mean : 58898	Mean : 1987.44
## 3rd Qu.:0.1689	3rd Qu.: 39409	3rd Qu.: 71935	3rd Qu.: 2005.30
## Max. :1.0000	Max. :150001	Max. :250001	Max. :205382.19
## NA's :916	NA's :882	NA's :436	NA's :726
## poor_share2010	poor_share2000	poor_share1990	share_white2010
## Min. :0.00000	Min. :0.0000	Min. :0.0000	Min. :0.0000
## 1st Qu.:0.05911	1st Qu.:0.0502	1st Qu.:0.0488	1st Qu.:0.4156
## Median :0.11581	Median :0.0956	Median :0.0955	Median :0.7355
## Mean :0.15092	Mean :0.1284	Mean :0.1322	Mean :0.6327
## 3rd Qu.:0.20493	3rd Qu.:0.1715	3rd Qu.:0.1752	3rd Qu.:0.8922
## Max. :1.00000	Max. :1.0000	Max. :1.0000	Max. :1.0000
## NA's :262	NA's :880	NA's :872	NA's :84
## share_black2010	share_hisp2010	share_asian2010	share_black2000
## Min. :0.00000	Min. :0.00000	Min. :0.0000	Min. :0.0000
## 1st Qu.:0.01336	1st Qu.:0.02443	1st Qu.:0.0039	1st Qu.:0.0090
## Median :0.04345	Median :0.06308	Median :0.0117	Median :0.0328
## Mean :0.14089	Mean :0.16099	Mean :0.0377	Mean :0.1316
## 3rd Qu.:0.15185	3rd Qu.:0.18790	3rd Qu.:0.0350	3rd Qu.:0.1297
## Max. :1.00000	Max. :1.00000	Max. :0.8918	Max. :1.0000
## NA's :84	NA's :84	NA's :1250	NA's :827
## share_white2000	share_hisp2000	share_asian2000	gsmn_math_g3_2013
## Min. :0.0000	Min. :0.0000	Min. :0.0000	Min. : -2.706
## 1st Qu.:0.5331	1st Qu.:0.0132	1st Qu.:0.0031	1st Qu.: 2.608
## Median :0.8132	Median :0.0361	Median :0.0092	Median : 3.220
## Mean :0.6943	Mean :0.1182	Mean :0.0306	Mean : 3.188
## 3rd Qu.:0.9302	3rd Qu.:0.1215	3rd Qu.:0.0280	3rd Qu.: 3.761
## Max. :1.0000	Max. :1.0000	Max. :0.9000	Max. : 6.878
## NA's :827	NA's :827	NA's :2146	NA's :1109
## rent_twobed2015	singleparent_share2010	singleparent_share2000	
## Min. : 99.0	Min. :0.0000	Min. :0.0000	
## 1st Qu.: 682.0	1st Qu.:0.1851	1st Qu.:0.1825	
## Median : 853.0	Median :0.2970	Median :0.2588	
## Mean : 951.2	Mean :0.3322	Mean :0.2935	
## 3rd Qu.:1125.0	3rd Qu.:0.4435	3rd Qu.:0.3663	
## Max. :3501.0	Max. :1.0000	Max. :1.0000	
## NA's :16592	NA's :631	NA's :910	
## singleparent_share1990	traveltime15_2010	emp2000	mail_return_rate2010
## Min. :0.0000	Min. :0.0000	Min. :0.0000	Min. : 0.00
## 1st Qu.:0.1242	1st Qu.:0.1842	1st Qu.:0.5331	1st Qu.: 74.30
## Median :0.1869	Median :0.2655	Median :0.6080	Median : 79.70

```
## Mean      :0.2288      Mean      :0.2950      Mean      :0.5945      Mean      : 78.73
## 3rd Qu.:0.2857      3rd Qu.:0.3762      3rd Qu.:0.6709      3rd Qu.: 84.30
## Max.      :1.0000      Max.      :1.0000      Max.      :1.0000      Max.      :100.00
## NA's      :999        NA's      :256        NA's      :851        NA's      :652
## ln_wage_growth_hs_grad jobs_total_5mi_2015 jobs_highpay_5mi_2015
## Min.      :-3.199      Min.      :      0      Min.      :      0
## 1st Qu.: -0.137      1st Qu.:  6522      1st Qu.:  2304
## Median :  0.046      Median :  42385      Median :  16981
## Mean      :  0.042      Mean      :111996      Mean      : 58566
## 3rd Qu.:  0.224      3rd Qu.:122916      3rd Qu.: 57828
## Max.      :  3.075      Max.      :2826437      Max.      :1794186
## NA's      :21563      NA's      :888        NA's      :888
## popdensity2010 ann_avg_job_growth_2004_2013 job_density_2013
## Min.      :      0.0    Min.      :-0.6067      Min.      :      0.0
## 1st Qu.:   319.4    1st Qu.: -0.0189      1st Qu.:    56.7
## Median :  2193.8    Median :  0.0085      Median :    412.6
## Mean      :  5236.2    Mean      : 0.0153      Mean      :   2157.0
## 3rd Qu.:  5277.5    3rd Qu.: 0.0410      3rd Qu.:  1371.1
## Max.      :543333.3    Max.      : 1.3365      Max.      :2905290.2
## NA's      :5        NA's      :2531      NA's      :736
```

Most variables have at least some number of missing values. Many of them lack values for almost all tracts, especially the variables that pool data at the household level for different racial groups. This may be because many racial tracts do not have enough people belonging to that racial group to collect enough aggregate data for that given variable.

Question 3 The `kfr_pooled_pooled_p25` variable calculates the mean percentile rank for household income for people born to parents at the 25th percentile of the national distribution of household income. In other words, the higher the value for this variable, the higher absolute mobility. This is calculated with using a linear model to capture the effect that being born to parents at the 25th percentile of household income will have on income outcomes as adults, pooled at the census tract level.

#Question 4: Histogram of Absolute Mobility at the 25th percentile

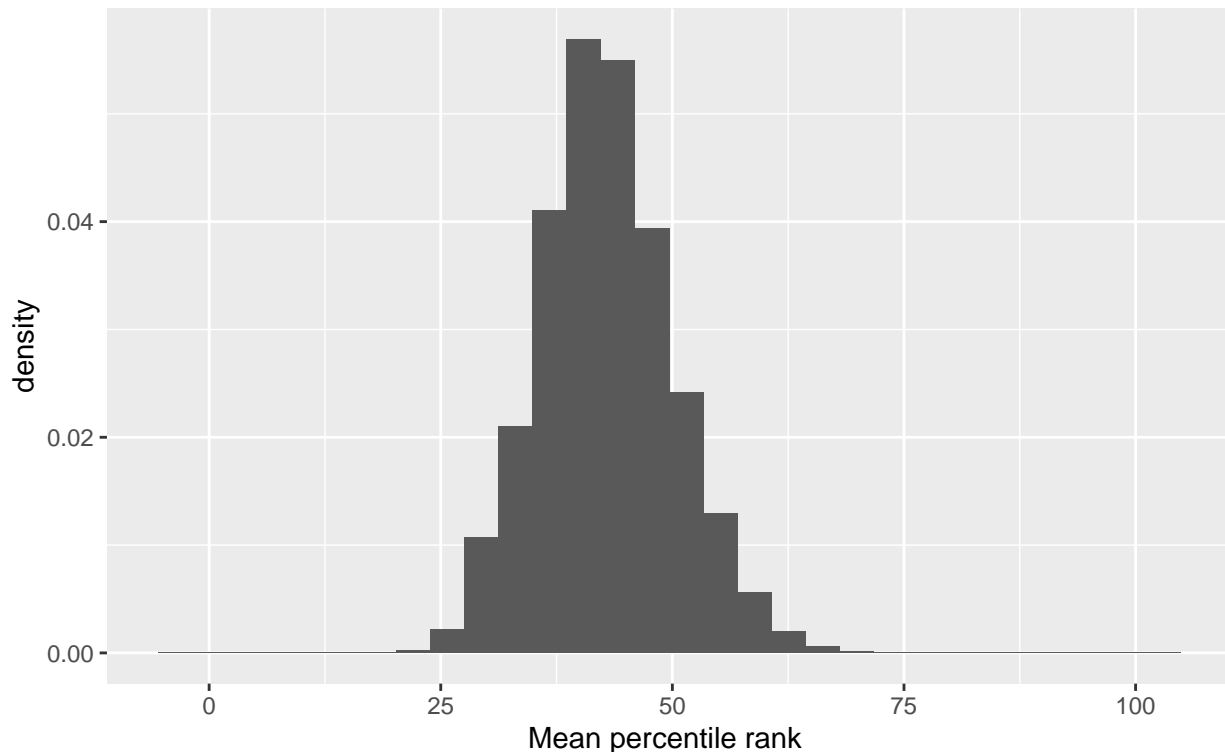
```
absmob_histo <- atlas |>
  ggplot() +
  geom_histogram(aes(x = kfr_pooled_pooled_p25, y = after_stat(density))) +
  labs(title = "Ranked Absolute Mobility at the 25th Percentile",
       subtitle = "For household income 2014-15",
       x = "Mean percentile rank ")
absmob_histo
```

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

```
## Warning: Removed 1189 rows containing non-finite values (`stat_bin()`).
```

Ranked Absolute Mobility at the 25th Percentile

For household income 2014–15



```
ggsave("absmob_histo.png")
```

```
## Saving 6.5 x 4.5 in image
```

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

```
## Warning: Removed 1189 rows containing non-finite values (`stat_bin()`).
```

The histogram shows an approximately normal distribution of mean percentile rank of absolute mobility at the 25th percentile, slightly left skewed. The majority of pooled census tract absolute mobility ranks are between 25 and 50.

#Question 5: Summary statistics for kfr_pooled_pooled_p25

```
sumstats <- summary(atlas$kfr_pooled_pooled_p25, na.rm = TRUE)
```

```
sdstats <- sd(atlas$kfr_pooled_pooled_p25, na.rm = TRUE)
```

#creating a table of those values

```
sumstat <- data.frame(c("Min", "1st Qu.", "Median", "Mean", "3rd Qu.", "Max", "SD", "NAs"),
                      c(-3.286, 38.070, 42.520, 42.858, 47.350, 103.349, 7.126422, 1189))
```

```
names(sumstat)[1] <- "Summary Stats"
```

```
names(sumstat)[2] <- "Values"
```

```
sumstat <- sumstat |>
  mutate(Values = round(Values, 2))
```

```
sumstat
```

```
##   Summary Stats  Values
## 1           Min   -3.29
## 2        1st Qu.  38.07
## 3          Median  42.52
## 4           Mean  42.86
```

```
## 5      3rd Qu.    47.35
## 6          Max   103.35
## 7          SD     7.13
## 8          NAs 1189.00
```

Question 6 `kfr_pooled_pooled_p25` can be negative or above 100 in these data because of the limitations of a simple linear model. The model does not know that we are trying to create a standardized percentile rank variable that (logically) starts at 0 and ends at 100. It simply receives a variable (in this case, parents' percentile rank of national household income pooled at the census tract) and uses it to predict the values of a dependent variable (kid percentile rank of household income). Because we are working with a large dataset, it is natural that the model, even when using standardized percentile values, will report values that are below 0 and above 100.

#Question 7: comparing absolute mobility at the 25th percentile in my census tract, state, and across the nation

```
usabsmob <- mean(atlas$kfr_pooled_pooled_p25, na.rm = TRUE) #mean of absolute mobility nationwide
ncabsmob <- mean(atlas$kfr_pooled_pooled_p25[atlas$state == "37"], na.rm = TRUE) #mean of absolute mobility in NC
myhood <- atlas %>% subset(state == "37" & county == "183" & tract == "53110") #creating data frame of my neighborhood
myhoodabsmob <- mean(myhood$kfr_pooled_pooled_p25, na.rm = TRUE) #mean of absolute mobility in my neighborhood

abscomp <- data.frame(c("My Neighborhood", "North Carolina", "United States"),
                      c(myhoodabsmob, ncabsmob, usabsmob))
names(abscomp)[1] <- "Level"
names(abscomp)[2] <- "Absolute Mobility at the 25th Percentile"
abscomp
```

```
##           Level Absolute Mobility at the 25th Percentile
## 1 My Neighborhood                38.18818
## 2 North Carolina                 37.89704
## 3 United States                  42.85813
```

My neighborhood/census tract has a slightly higher level of absolute mobility (at the 25th percentile) compared to my home state of North Carolina. Both my census tract and North Carolina have a lower level of absolute mobility than the national average.

#Question 8: calculating and comparing standard deviations of absolute mobility in my home county, state, and across the nation

```
usasd <- sd(atlas$kfr_pooled_pooled_p25, na.rm = TRUE) #sd of absolute mobility nationwide
ncsd <- sd(atlas$kfr_pooled_pooled_p25[atlas$state == "37"], na.rm = TRUE) #sd of absolute mobility in NC
wake_co <- atlas %>% subset(state == "37" & county == "183") #creating data frame of just my tract
countysd <- sd(wake_co$kfr_pooled_pooled_p25, na.rm = TRUE) #sd of absolute mobility in my county

sdcomp <- data.frame(c("Wake County", "North Carolina", "United States"),
                     c(countysd, ncsd, usasd))
names(sdcomp)[1] <- "Level"
names(sdcomp)[2] <- "Std Dev of Absolute Mobility at the 25th Percentile"
sdcomp
```

```
##           Level Std Dev of Absolute Mobility at the 25th Percentile
## 1 Wake County                7.692496
## 2 North Carolina             5.756876
## 3 United States              7.126422
```

The standard deviation of mobility outcomes for my Wake County, NC (7.69) is slightly higher than the standard deviation at the national level. North Carolina's mobility outcome standard deviation of 5.75 is lower than the amount of spread at the county or national level. This gives us a sense of how results may vary, which makes sense because of the increasing sample size for each successive level of analysis.

#Question 9: plotting the relationship between upward mobility and rent in my neighborhood

#9a: Scatter plot of my home county

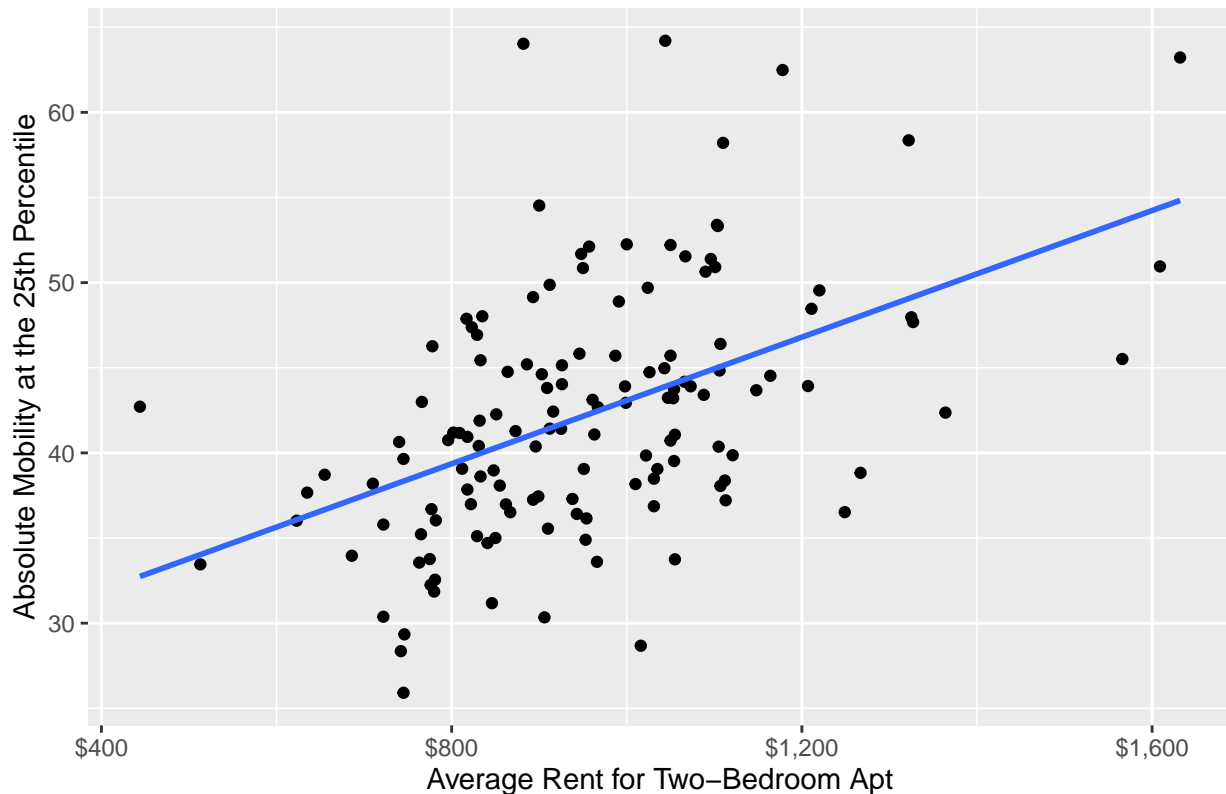
```
scatterrent <- atlas |>
  filter(state == 37,
         county == 183) |>
  ggplot() +
  geom_point(aes(x = rent_twobed2015, y = kfr_pooled_pooled_p25)) +
  geom_smooth(aes(x = rent_twobed2015, y = kfr_pooled_pooled_p25), method = "lm", se = F) +
  labs(x = "Average Rent for Two-Bedroom Apt",
       y = "Absolute Mobility at the 25th Percentile",
       title = "Relationship between rent and absolute mobility in Wake County, NC") +
  scale_x_continuous(labels = scales::dollar_format())
scatterrent
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 50 rows containing non-finite values (`stat_smooth()`).
```

```
## Warning: Removed 50 rows containing missing values (`geom_point()`).
```

Relationship between rent and absolute mobility in Wake County, NC



```
ggsave("scatterrent.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 50 rows containing non-finite values (`stat_smooth()`).
```

```
## Removed 50 rows containing missing values (`geom_point()`).
```


Question 9B There is an apparent positive correlation between rent and absolute mobility at the 25th percentile. The line of best fit suggests that as rent increases in my home county, the percentile rank of absolute mobility increases as well. This suggests that low income children living in higher-priced tracts experience higher mobility than low income children living in cheaper (and, presumably, more economically and socially disadvantaged) tracts.

Question 9C Opportunity bargains are neighborhoods with cheaper than average rents and above average absolute mobility outcomes. For this question, I am defining a given neighborhood an opportunity bargain if it has under \$1500 in median two-bedroom rent and the highest percentile rank for absolute mobility at the 25th percentile for my given county.

#Question 9c: Defining Opportunity Bargains

```
myhood <- atlas %>% subset(state == "37" & county == "183" & tract == "53110") #creating data frame of my neighborhood
myhoodabsmob
```

```
## [1] 38.18818
```

```
myhoodrent <- mean(myhood$rent_twobed2015, na.rm = TRUE) #myneighborhood rent
myhoodrent
```

```
## [1] 710
```

With an absolute mobility rank (at the 25th percentile) of 38 and a median two-bedroom apartment rent of \$710, my neighborhood can perhaps be considered an opportunity tract, but we would need to compare it to other tracts in my county to make a more practical judgment. What about other census tracts in my home county of Wake County?

#Finding other opportunity bargains in my home county

```
wake_co <- atlas %>% subset(state == "37" & county == "183") #creating data frame of just my tract
wake_co$bargain_index <- wake_co$kfr_pooled_pooled_p25/wake_co$rent_twobed2015
wake_bargains <- wake_co |>
  select(tract_name, tract, kfr_pooled_pooled_p25, rent_twobed2015, bargain_index) |>
  arrange(desc(bargain_index))
print(wake_bargains)
```

```
## # A tibble: 186 x 5
##   tract_name          tract kfr_pooled_pooled_p25 rent_two~1 barga~2
##   <chr>              <dbl>          <dbl>          <dbl>    <dbl>
## 1 Holly Springs, NC    53206             42.7            444  0.0962
## 2 Brookgreen Forest, Cary, NC 53423             64.0            882  0.0726
## 3 Wendell, NC          54403             33.5            513  0.0652
## 4 Umstead, Raleigh, NC 53717             64.2           1044  0.0615
## 5 North Raleigh, Raleigh, NC 53715             54.5            900  0.0606
## 6 Wake Forest, NC      54204             46.3            778  0.0595
## 7 Wendell, NC          54402             37.7            635  0.0593
## 8 Wendell, NC          54111             38.7            655  0.0591
## 9 Southwest Raleigh, Raleigh, NC 52404             47.9            817  0.0586
## 10 Zebulon, NC         54301             36.0            623  0.0578
## # ... with 176 more rows, and abbreviated variable names 1: rent_twobed2015,
## # 2: bargain_index
```

Using the opportunity bargain index method derived by dividing the absolute mobility rank by the median two-bedroom rent, we see that my census tract is #21 out of the 186 tracts in my home county. Furthermore, we see that my tract's bargain index variable is influenced by the very cheap median rent in my neighborhood - but cheap rent alone does not make an opportunity bargain. Therefore, I would not classify my neighborhood/census tract as an opportunity bargain. I will go ahead and determine opportunity bargains those census tracts with rents under \$1500 and an absolute mobility rank of over 50. Let's see where they are, if any.

```
#Finding opportunity bargains in Wake County
top_bargains <- wake_bargains |>
  filter(rent_twobed2015 <= 1500,
         kfr_pooled_pooled_p25 >= 50)
print(top_bargains)
```

```
## # A tibble: 17 x 5
##   tract_name                tract kfr_pooled_pooled_p25 rent_two~1 barga-2
##   <chr>                    <dbl>          <dbl>          <dbl>    <dbl>
## 1 Brookgreen Forest, Cary, NC 53423            64.0            882  0.0726
## 2 Umstead, Raleigh, NC      53717            64.2           1044  0.0615
## 3 North Raleigh, Raleigh, NC 53715            54.5            900  0.0606
## 4 Northwest Raleigh, Raleigh, NC 53712            51.7            948  0.0545
## 5 Dutchess Village, Cary, NC 53524            52.1            957  0.0545
## 6 North Hills, Raleigh, NC   51501            50.9            950  0.0535
## 7 Cary, NC                  53603            62.5           1178  0.0530
## 8 Breckenridge, Raleigh, NC 53725            58.2           1110  0.0524
## 9 North Raleigh, Raleigh, NC 54011            52.2           1000  0.0522
## 10 North Raleigh, Raleigh, NC 54016            52.2           1050  0.0497
## 11 Northwest Raleigh, Raleigh, NC 53724            53.4           1103  0.0484
## 12 Southwest Raleigh, Raleigh, NC 52301            51.5           1067  0.0483
## 13 Cary, NC                 53425            53.3           1104  0.0483
## 14 Lochmere, Cary, NC       53004            51.4           1096  0.0469
## 15 Hillisdale Forest, Cary, NC 53505            50.6           1090  0.0465
## 16 Stonebridge, Raleigh, NC 53807            50.9           1101  0.0462
## 17 Medfield Estates, Raleigh, NC 53521            58.4           1322  0.0441
## # ... with abbreviated variable names 1: rent_twobed2015, 2: bargain_index
```

So now we see a list of 17 census tracts - from my experience, most of them clustered in the neighborhoods of upper middle-class North Raleigh and rapidly growing Cary - that we can reasonably say are opportunity bargains. Let's highlight them on the scatterplot of rent and absolute mobility to place them in the context of other census tracts. Do they stand out?

```
#Highlighting opportunity bargains in Wake County, NC
```

```
scatterbargain <- atlas |>
  filter(state == 37,
         county == 183) |>
  ggplot() +
  geom_point(aes(x = rent_twobed2015, y = kfr_pooled_pooled_p25)) +
  geom_point(data = top_bargains, aes(x = rent_twobed2015, y = kfr_pooled_pooled_p25),
            color = "red", size = 3) +
  geom_text(aes(x = rent_twobed2015, y = kfr_pooled_pooled_p25, label = tract), check_overlap = TRUE,
            size = 3, nudge_x = 2, nudge_y = 1) +
  geom_smooth(aes(x = rent_twobed2015, y = kfr_pooled_pooled_p25), method = "lm", se = F) +
  labs(x = "Average Rent for Two-Bedroom Apt",
       y = "Absolute Mobility at the 25th Percentile",
       title = "Relationship between rent and absolute mobility in Wake County, NC",
       subtitle = "Opportunity bargains highlighted in red") +
  scale_x_continuous(labels = scales::dollar_format())
scatterbargain
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 50 rows containing non-finite values (`stat_smooth()`).
```

```
## Warning: Removed 50 rows containing missing values (`geom_text()`).
```

Opportunity bargains highlighted in red



```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 50 rows containing missing values (`geom_point()`).
```

```
## Warning: Removed 50 rows containing missing values (`geom_text()`).
```

And there they are! These are the bargain tracts with cheap rent and above average mobility outcomes in Wake County, NC.

#10A: scatter plot of poverty rate in 1990 and 2010 for Wake County

11

```

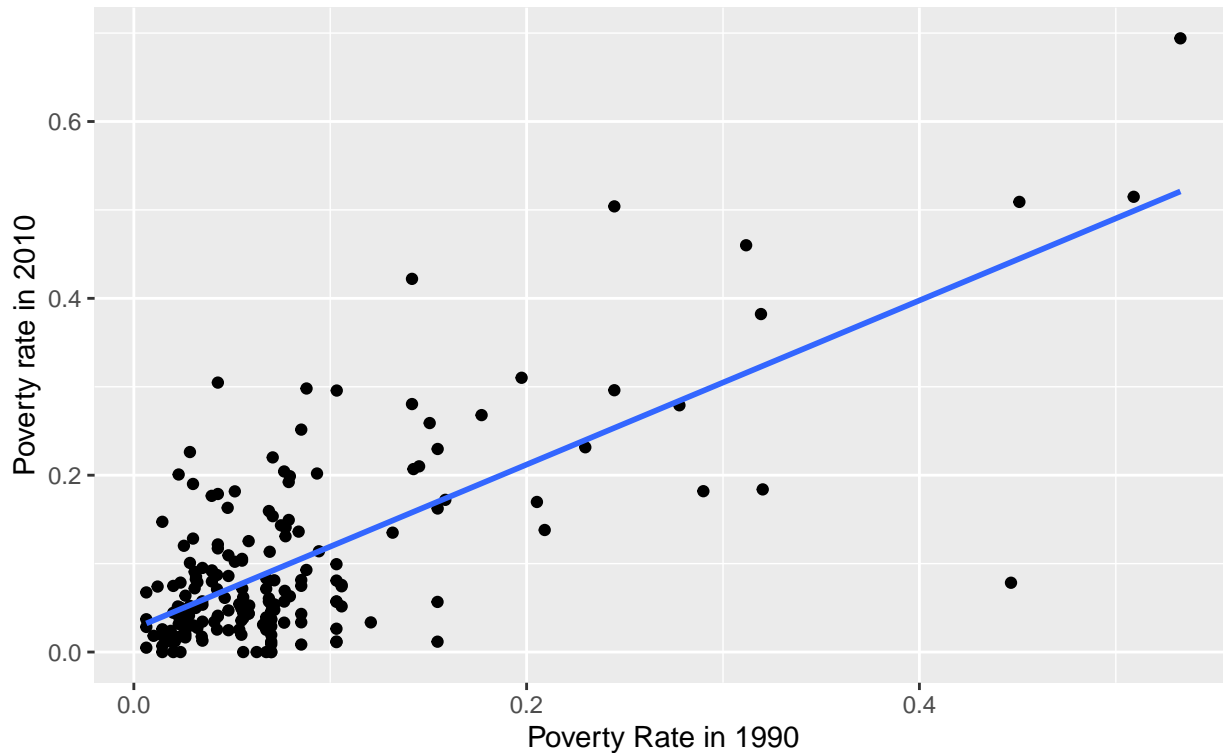
      subtitle = "Most tracts saw few changes in their poverty rate over time")
wakepoverty

```

```
## `geom_smooth()` using formula = 'y ~ x'
```

Poverty Rate, 1990–2010 in Wake County, NC

Most tracts saw few changes in their poverty rate over time



```
ggsave("wakepoverty.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

#10A: scatter plot of Black population in 1990 and 2010 for Wake County

```

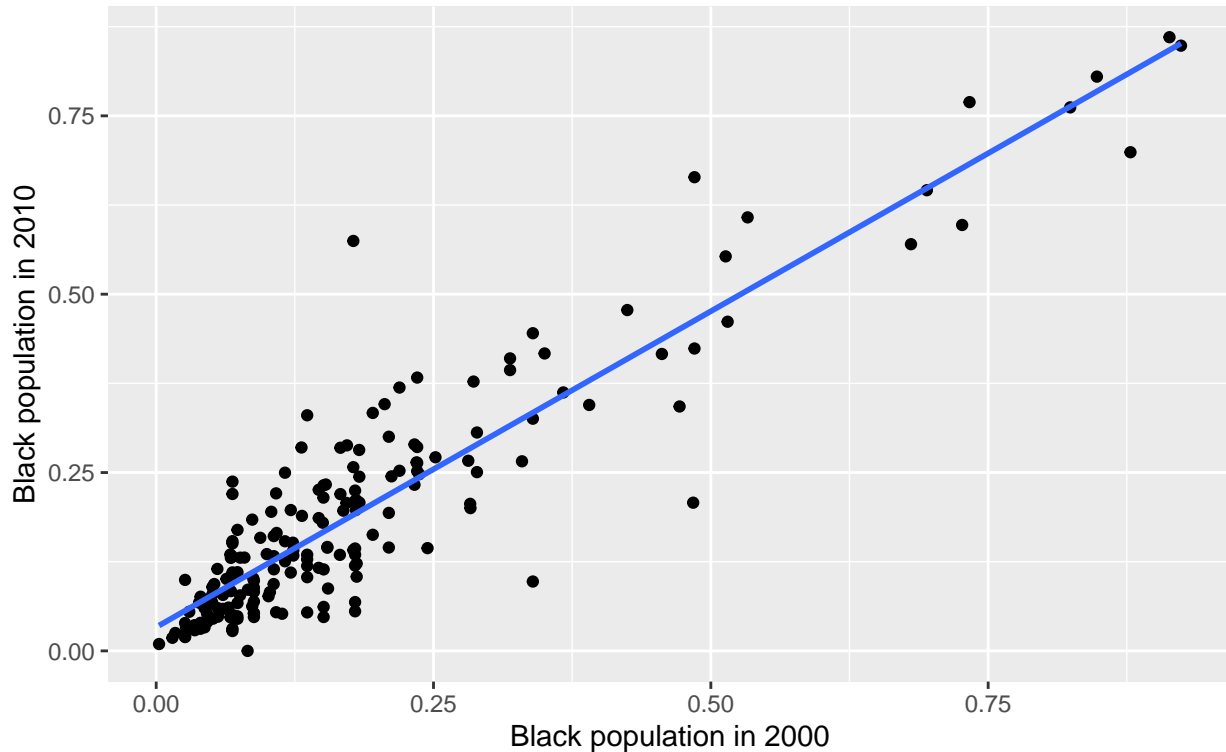
wakeblack <- atlas |>
  filter(state == 37,
         county == 183) |>
  ggplot(aes(x = share_black2000, y = share_black2010)) +
  geom_point() +
  geom_smooth(method = "lm", se = F) +
  labs(x = "Black population in 2000",
       y = "Black population in 2010",
       title = "Black population share 2000–2010 in Wake County, NC",
       subtitle = "Roughly equal number of census tracts saw their Black population share rise
wakeblack

```

```
## `geom_smooth()` using formula = 'y ~ x'
```

Black population share 2000–2010 in Wake County, NC

Roughly equal number of census tracts saw their Black population share rise or fall over



```
ggsave("wakeblack.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

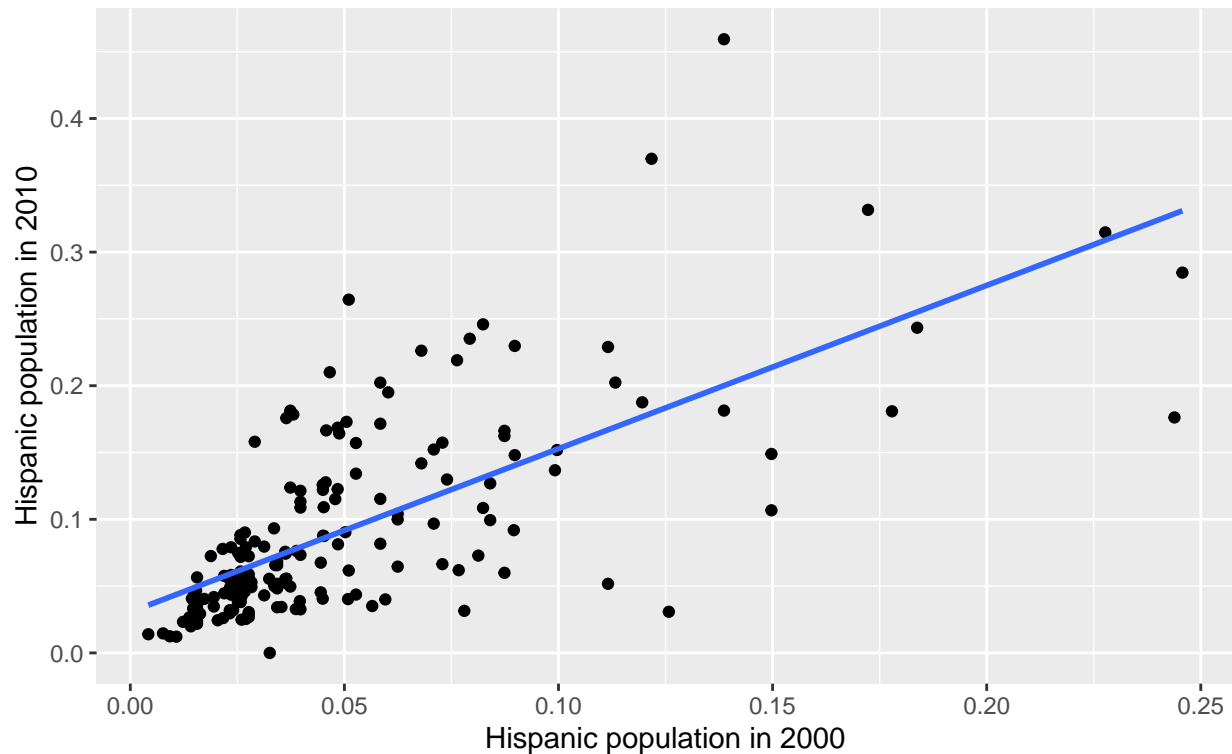
```
#10A: scatter plot of Hispanic population in 1990 and 2010 for Wake County
```

```
wakehisp <- atlas |>
  filter(state == 37,
         county == 183) |>
  ggplot(aes(x = share_hisp2000, y = share_hisp2010)) +
  geom_point() +
  geom_smooth(method = "lm", se = F) +
  labs(x = "Hispanic population in 2000",
       y = "Hispanic population in 2010",
       title = "Hispanic population share 2000-2010 in Wake County, NC",
       subtitle = "Several tracts saw their Hispanic population grow significantly over time")
wakehisp
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

Hispanic population share 2000–2010 in Wake County, NC

Several tracts saw their Hispanic population grow significantly over time



```
ggsave("wakehisp.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
#10A: scatter plot of Asian population in 1990 and 2010 for Wake County
```

```
wakeasian <- atlas |>
  filter(state == 37,
         county == 183) |>
  ggplot(aes(x = share_asian2000, y = share_asian2010)) +
  geom_point() +
  geom_smooth(method = "lm", se = F) +
  labs(x = "Asian population in 2000",
       y = "Asian population in 2010",
       title = "Asian population share 2000–2010 in Wake County, NC",
       subtitle = "Select number of tracts saw their Asian population more than double over time")
wakeasian
```

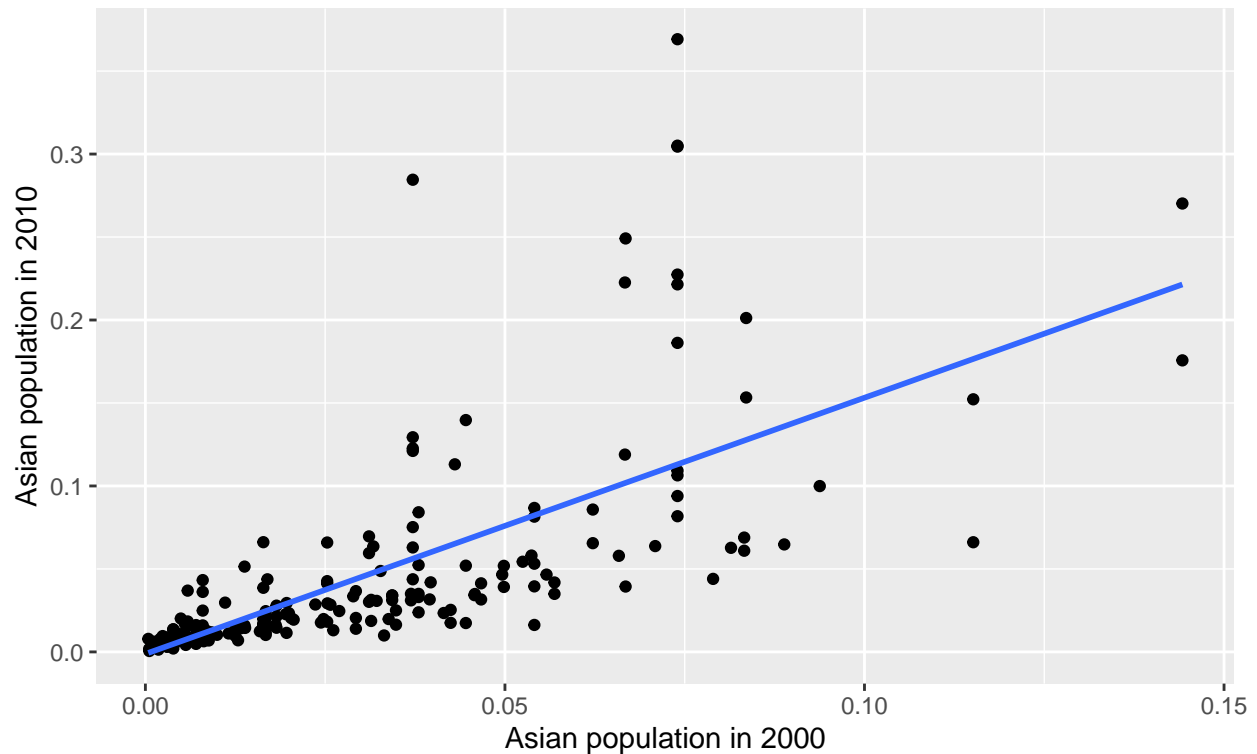
```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 2 rows containing non-finite values (`stat_smooth()`).
```

```
## Warning: Removed 2 rows containing missing values (`geom_point()`).
```

Asian population share 2000–2010 in Wake County, NC

Select number of tracts saw their Asian population more than double over time



```
ggsave("wakeasian.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 2 rows containing non-finite values (`stat_smooth()`).
```

```
## Removed 2 rows containing missing values (`geom_point()`).
```

```
#10A: scatter plot of White population in 1990 and 2010 for Wake County
```

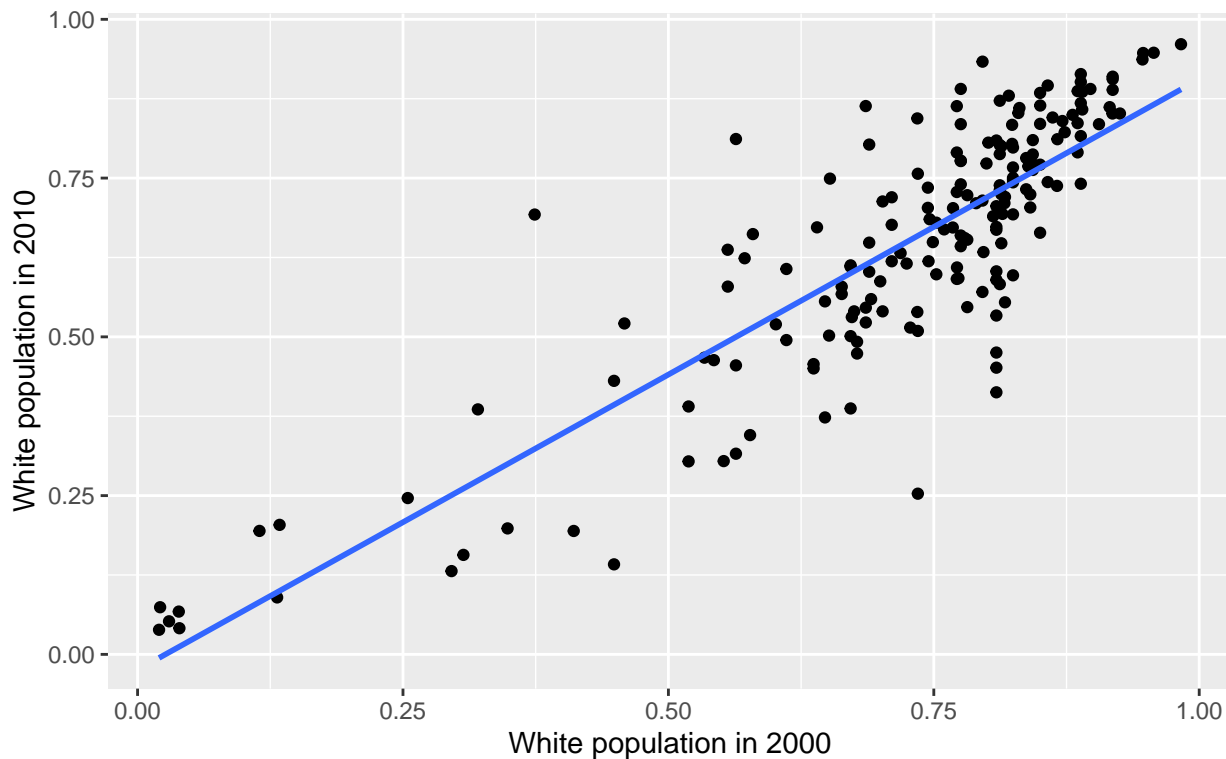
```
wakewhite <- atlas |>
  filter(state == 37,
         county == 183) |>
  ggplot(aes(x = share_white2000, y = share_white2010)) +
  geom_point() +
  geom_smooth(method = "lm", se = F) +
  labs(x = "White population in 2000",
       y = "White population in 2010",
       title = "White population share 2000–2010 in Wake County, NC",
       subtitle = "Several tracts are no longer majority white")
```

```
wakewhite
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

White population share 2000–2010 in Wake County, NC

Several tracts are no longer majority white



```
ggsave("wakewhite.png")
```

```
## Saving 6.5 x 4.5 in image
## `geom_smooth()` using formula = 'y ~ x'
```

Question 10B Between 1990, 2000, and 2010, Wake County saw relatively minor changes to its poverty rate, but significant changes to its demographic composition. Most tracts saw fairly little change in their poverty rate between 1990 and 2010. However, several tracts saw significant increases in their share of Asian and Hispanic populations between 2000 and 2010. At the same time, many tracts saw their share of the White population fall under 50%. This highlights the growing share of nonwhite populations in Wake County over the last few decades.

#Question 11a: Average absolute mobility rate for neighborhoods grouped by HOLC grade (A, B, C, and D)

```
holcmobil <- atlas |>
  mutate(grade = case_when(HOLC_A > 0.5 & !is.na(HOLC_A) ~ 'A',
                           HOLC_B > 0.5 & !is.na(HOLC_B) ~ 'B',
                           HOLC_C > 0.5 & !is.na(HOLC_C) ~ 'C',
                           HOLC_D > 0.5 & !is.na(HOLC_D) ~ 'D')) |>
  group_by(grade) |>
  summarize(mean_absolute_mobility = mean(kfr_pooled_pooled_p25, na.rm = TRUE))
```

```
holcmobil
```

```
## # A tibble: 5 x 2
##   grade mean_absolute_mobility
##   <chr>          <dbl>
## 1 A             44.0
## 2 B             42.5
```



```
## 3 C 39.9
## 4 D 36.2
## 5 <NA> 43.2
```

There is a clear pattern of increasing upward mobility for children born in the 1980s in Census tracts with higher 1930s HOLC grades. For tracts where a majority of its area was graded A, the average mobility rank at the 25th percentile is roughly 44. As we start to look at tracts with majority of their areas graded B through D, we see a consistent pattern of a decrease of 2-3 mobility percentile ranks. This suggests that practices such as redlining may have had an adverse effect on mobility outcomes in various neighborhoods that persists to this day.

#Q 11b: Black population share in 2000 for neighborhoods grouped by HOLC grade

```
holcblack <- atlas |>
  mutate(grade = case_when(HOLC_A > 0.5 & !is.na(HOLC_A) ~ 'A',
                           HOLC_B > 0.5 & !is.na(HOLC_B) ~ 'B',
                           HOLC_C > 0.5 & !is.na(HOLC_C) ~ 'C',
                           HOLC_D > 0.5 & !is.na(HOLC_D) ~ 'D')) |>
  group_by(grade) |>
  summarize(mean_blackpop = mean(share_black2000, na.rm = TRUE))
```

```
holcblack
```

```
## # A tibble: 5 x 2
##   grade mean_blackpop
##   <chr>      <dbl>
## 1 A         0.201
## 2 B         0.289
## 3 C         0.333
## 4 D         0.466
## 5 <NA>      0.111
```

It is clear that when looking at the racial composition of HOLC graded neighborhoods, we see that neighborhoods graded A have less than half of Black population share as we see in D rated neighborhoods. There is a clear pattern of increasing Black population share as we move across neighborhoods by their HOLC grade. We cannot rule out race as a confounding variable in driving mobility outcomes.

#Q11C: Average of Black and White pooled absolute mobility for neighborhoods grouped by HOLC grade

```
holcracefixed <- atlas |>
  mutate(grade = case_when(HOLC_A > 0.5 & !is.na(HOLC_A) ~ 'A',
                           HOLC_B > 0.5 & !is.na(HOLC_B) ~ 'B',
                           HOLC_C > 0.5 & !is.na(HOLC_C) ~ 'C',
                           HOLC_D > 0.5 & !is.na(HOLC_D) ~ 'D')) |>
  group_by(grade) |>
  summarize(mean_blackmobil = mean(kfr_black_pooled_p25, na.rm = TRUE),
           mean_whitemobil = mean(kfr_white_pooled_p25, na.rm = TRUE))
```

```
holcracefixed
```

```
## # A tibble: 5 x 3
##   grade mean_blackmobil mean_whitemobil
##   <chr>      <dbl>      <dbl>
## 1 A         34.4         50.4
## 2 B         34.5         48.8
## 3 C         33.2         46.3
## 4 D         31.6         44.1
```

```
## 5 <NA>          34.1          46.3
```

When we look at absolute mobility rates by HOLC neighborhood grade and race, racial composition cannot be a confounder in this analysis because we are isolating racial variables. When we look solely at Black mobility rates or white mobility rates, we are looking solely at Black child outcomes, so the potential effect that former HOLC neighborhood grade may have on mobility outcomes should be more clear.

Looking at Black and white pooled mobility rates, it's evident that mobility percentile ranks decrease as neighborhood HOLC grades decrease. There is a decrease of roughly 3 percentile ranks for Black children's mobility outcomes between grade A and D neighborhoods. Similarly, there is a decrease of roughly 6 percentile ranks for White children's outcomes between A and D neighborhoods. What's notable is that the average mobility rank for white children living in D neighborhoods remains higher (44 vs. 34) than the average mobility rank for Black children who lived in A neighborhoods. Clearly, HOLC grade and race have separate but similar impacts on mobility outcomes.

#Q11D: Analyzing potential causal relationship between HOLC neighborhood grade and vegetation, extreme

```
holccausal <- atlas |>
  mutate(grade = case_when(HOLC_A > 0.5 & !is.na(HOLC_A) ~ 'A',
                           HOLC_B > 0.5 & !is.na(HOLC_B) ~ 'B',
                           HOLC_C > 0.5 & !is.na(HOLC_C) ~ 'C',
                           HOLC_D > 0.5 & !is.na(HOLC_D) ~ 'D')) |>
  group_by(grade) |>
  summarize(mean_veg = mean(vegetation, na.rm = TRUE),
            mean_heat = mean(extreme_heat, na.rm = TRUE),
            mean_developed = mean(developed, na.rm = TRUE))
```

```
holccausal
```

```
## # A tibble: 5 x 4
##   grade mean_veg mean_heat mean_developed
##   <chr>   <dbl>   <dbl>         <dbl>
## 1 A      -0.0802    4.22          0.892
## 2 B      -0.205     5.73          0.940
## 3 C      -0.211     5.76          0.943
## 4 D      -0.263     6.39          0.945
## 5 <NA>  -0.0721     2.40          0.544
```

The analysis shows that neighborhoods with higher HOLC grades have more vegetation, lower extreme heat temperatures, and slightly less developed land (all three variables relative to a baseline). While all graded neighborhood levels have lower levels of vegetation than the baseline (which makes sense, since graded neighborhoods were more likely to be in urban areas), the level of vegetation gets lower with each decreasing HOLC grade. As HOLC grades decrease, there is a rise in extreme land surface heat during summer. Finally, while all graded tracts are fairly highly developed, lower graded tracts have higher rates of land development. This suggests that factors like less green space and investment in public parks are a legacy of redlining, and potentially contributing to lower mobility outcomes today.

#Q11E: Visualizing variables means affected by HOLC neighborhood grades

#Absolute mobility pooled for all races and genders, grouped by HOLC neighborhood grade

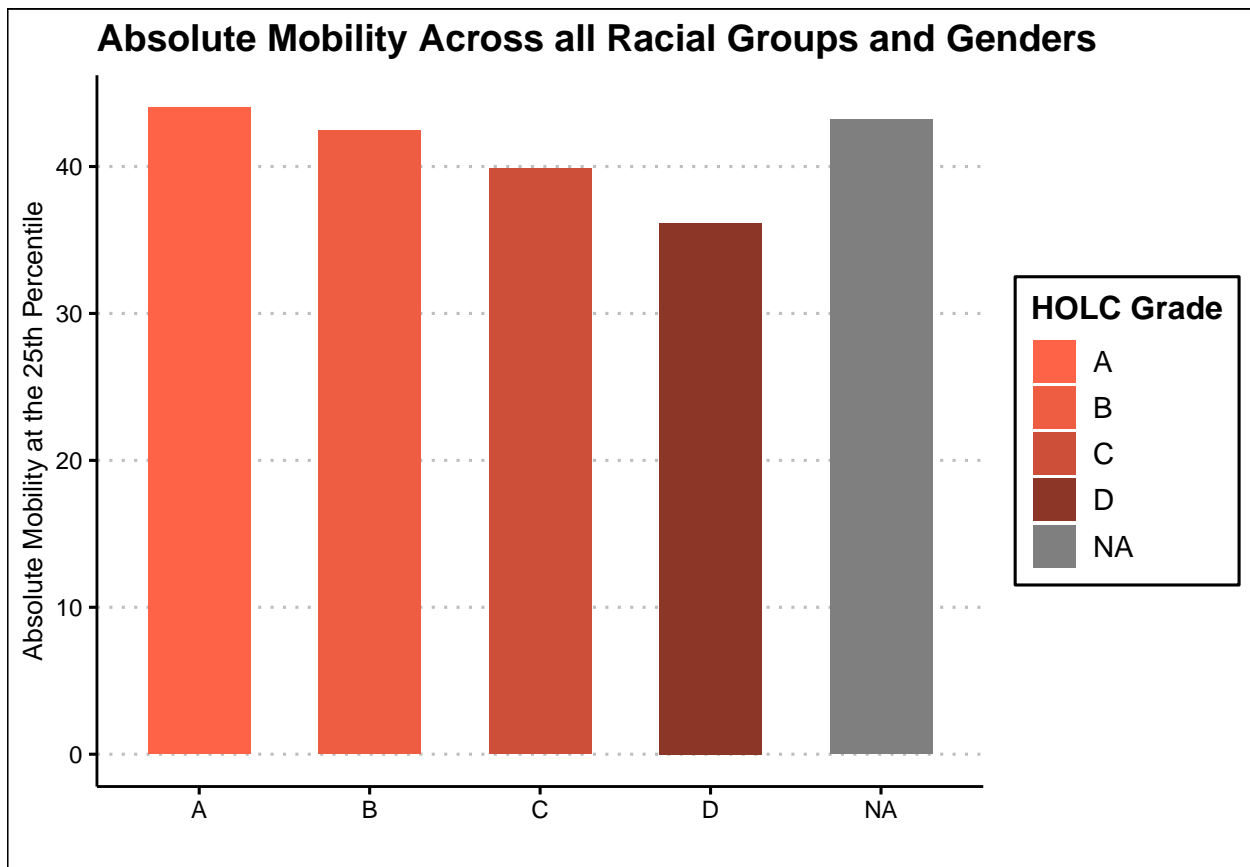
```
mobilgraph <- atlas |>
  mutate(grade = case_when(HOLC_A > 0.5 & !is.na(HOLC_A) ~ 'A',
                           HOLC_B > 0.5 & !is.na(HOLC_B) ~ 'B',
                           HOLC_C > 0.5 & !is.na(HOLC_C) ~ 'C',
                           HOLC_D > 0.5 & !is.na(HOLC_D) ~ 'D')) |>
  group_by(grade) |>
```

```

summarize(mean_absolute_mobility = mean(kfr_pooled_pooled_p25, na.rm = TRUE)) |>
ggplot(aes(x = grade, y = mean_absolute_mobility, fill = grade)) +
geom_bar(stat = "identity", show.legend = TRUE, width = 0.6) +
scale_fill_manual(values = c("tomato1", "tomato2", "tomato3", "tomato4")) +
labs(x = "",
     y = "Absolute Mobility at the 25th Percentile",
     title = "Absolute Mobility Across all Racial Groups and Genders",
     fill = "HOLC Grade") +
theme_clean()

```

mobilgraph



```
ggsave("mobilgraph.png")
```

Saving 6.5 x 4.5 in image

#Visualizing Black population share across census tracts grouped by HOLC neighborhood grade

```

holcblackgraph <- atlas |>
  mutate(grade = case_when(HOLC_A > 0.5 & !is.na(HOLC_A) ~ 'A',
                           HOLC_B > 0.5 & !is.na(HOLC_B) ~ 'B',
                           HOLC_C > 0.5 & !is.na(HOLC_C) ~ 'C',
                           HOLC_D > 0.5 & !is.na(HOLC_D) ~ 'D')) |>
  group_by(grade) |>
  summarize(mean_blackpop = mean(share_black2000, na.rm = TRUE))|>
  ggplot(aes(x = grade, y = mean_blackpop, fill = grade)) +
  geom_bar(stat = "identity", show.legend = TRUE, width = 0.6) +

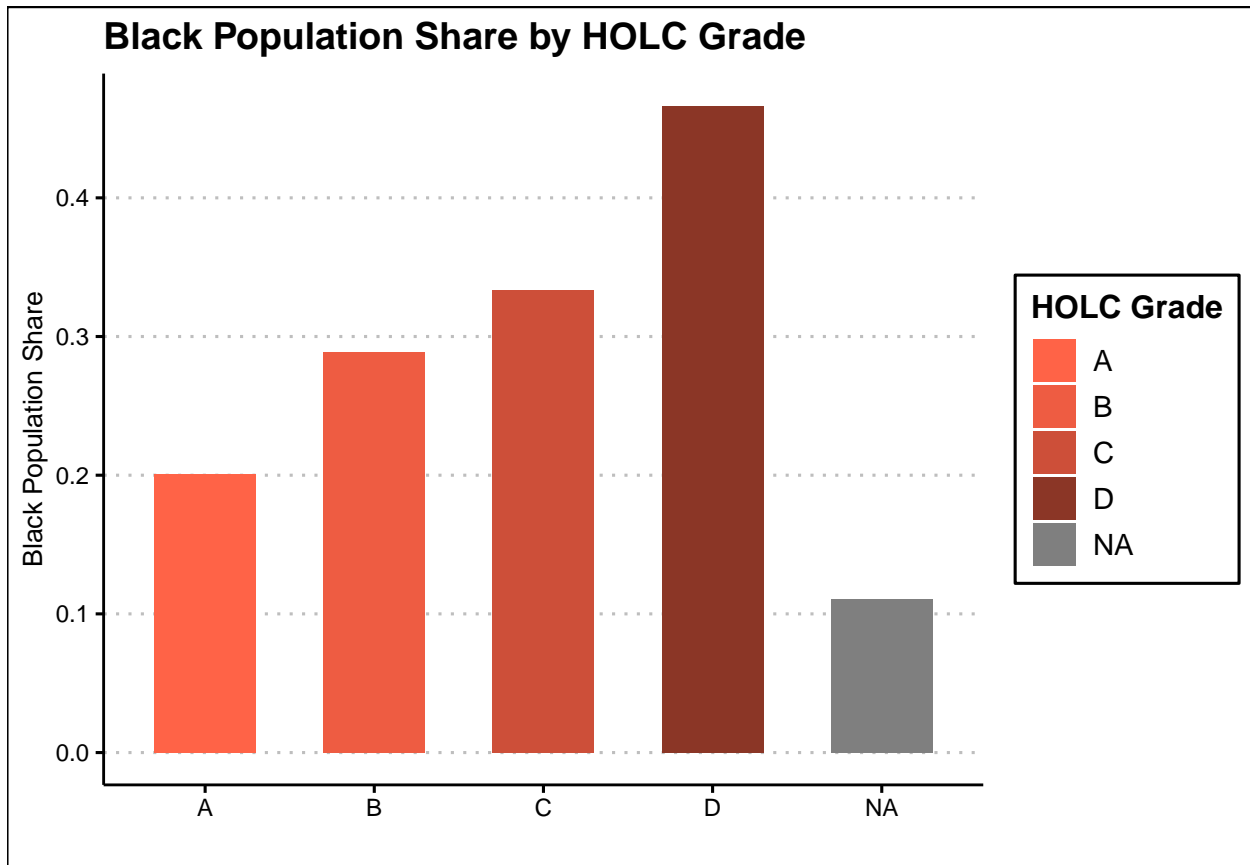
```

```

scale_fill_manual(values = c("tomato1", "tomato2", "tomato3", "tomato4")) +
labs(x = "",
     y = "Black Population Share",
     title = "Black Population Share by HOLC Grade",
     fill = "HOLC Grade") +
theme_clean()

```

holcblackgraph



```
ggsave("holcblackgraph.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
#Visualizing Black and White pooled absolute mobility for neighborhoods grouped by HOLC grade
```

```
#Black mobility by HOLC graded neighborhood
```

```

blackmobilgraph <- atlas |>
  mutate(grade = case_when(HOLC_A > 0.5 & !is.na(HOLC_A) ~ 'A',
                           HOLC_B > 0.5 & !is.na(HOLC_B) ~ 'B',
                           HOLC_C > 0.5 & !is.na(HOLC_C) ~ 'C',
                           HOLC_D > 0.5 & !is.na(HOLC_D) ~ 'D')) |>

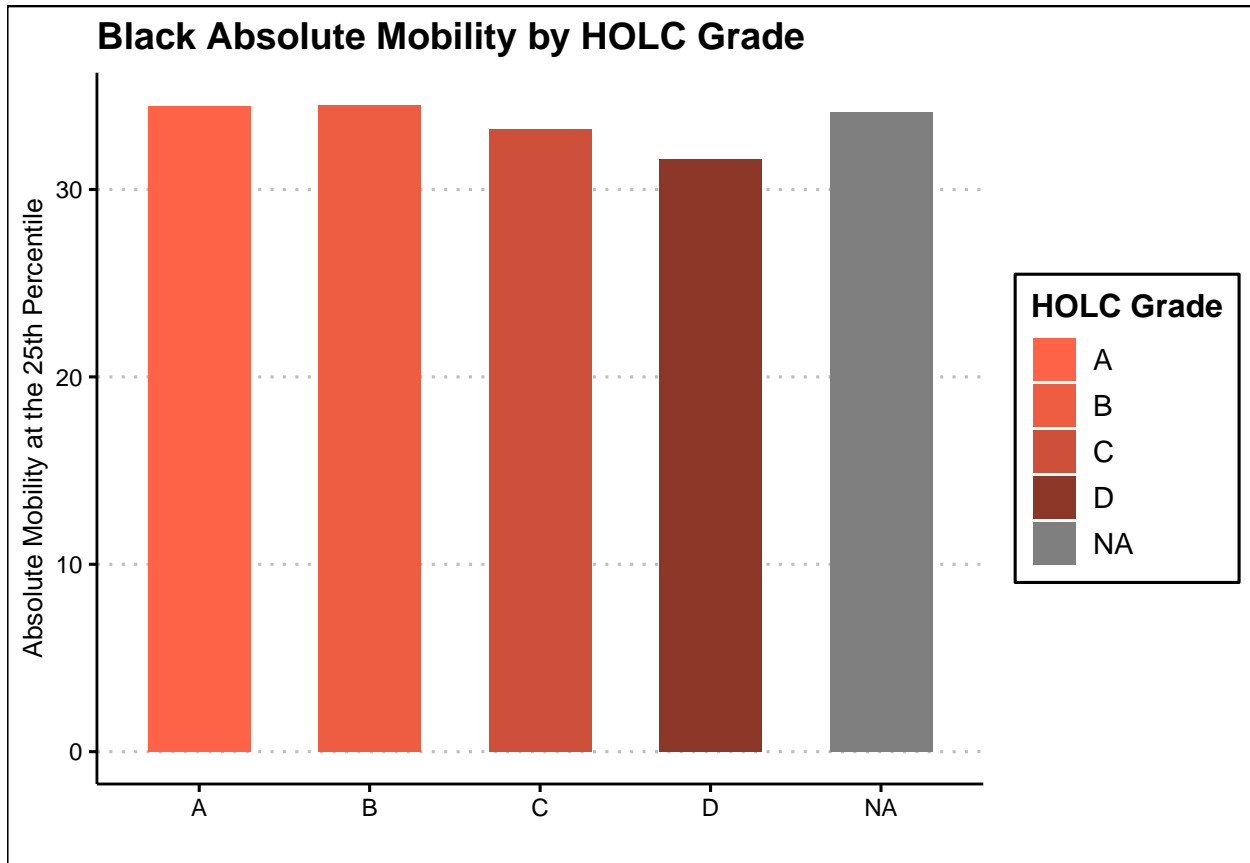
  group_by(grade) |>
  summarize(mean_blackmobil = mean(kfr_black_pooled_p25, na.rm = TRUE),
            mean_whitemobil = mean(kfr_white_pooled_p25, na.rm = TRUE)) |>

  ggplot() +
  geom_bar(aes(x = grade, y = mean_blackmobil, fill = grade), stat = "identity", show.legend = FALSE) +
  scale_fill_manual(values = c("tomato1", "tomato2", "tomato3", "tomato4")) +

```

```
labs(x = "",
     y = "Absolute Mobility at the 25th Percentile",
     title = "Black Absolute Mobility by HOLC Grade",
     fill = "HOLC Grade") +
theme_clean()
```

blackmobilgraph



```
ggsave("blackmobilgraph.png")
```

Saving 6.5 x 4.5 in image

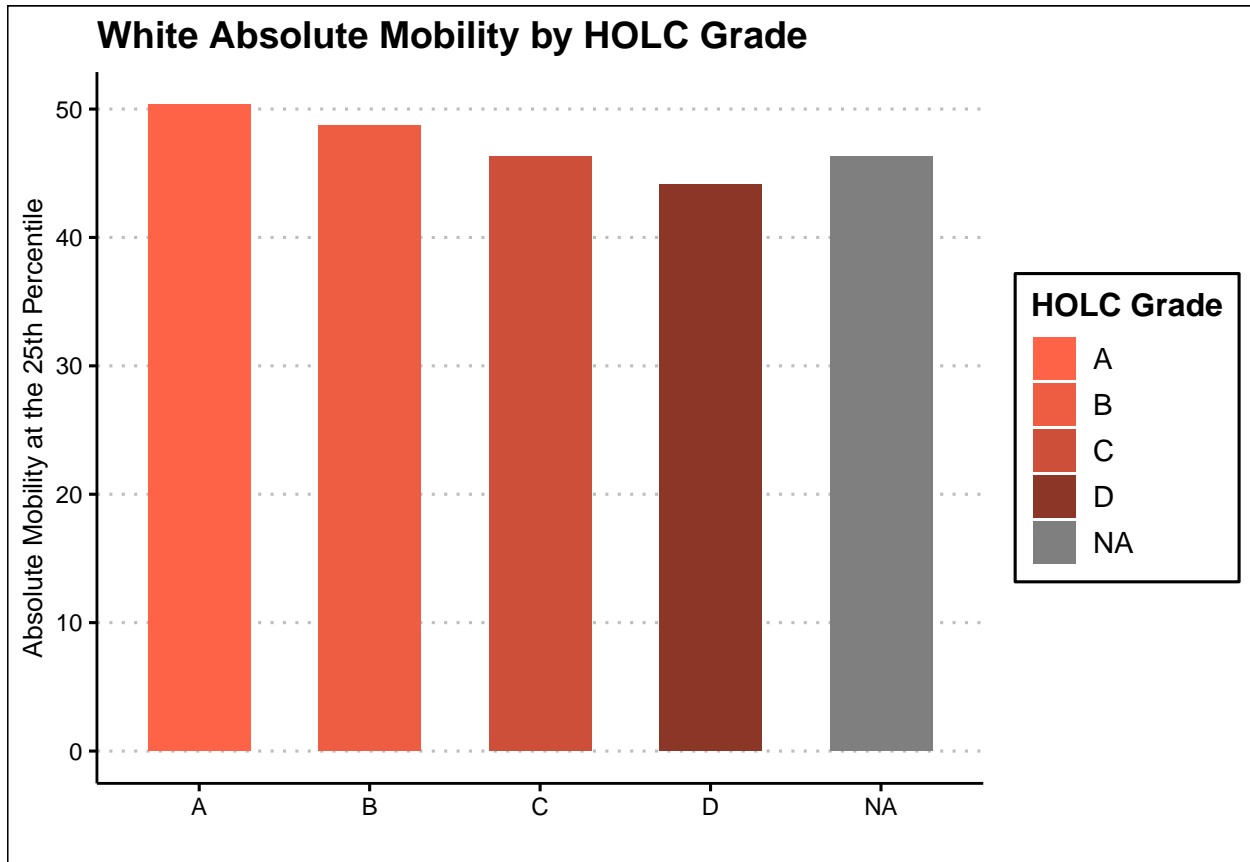
#White mobility by HOLC graded neighborhood

```
whitemobilgraph <- atlas |>
  mutate(grade = case_when(HOLC_A > 0.5 & !is.na(HOLC_A) ~ 'A',
                           HOLC_B > 0.5 & !is.na(HOLC_B) ~ 'B',
                           HOLC_C > 0.5 & !is.na(HOLC_C) ~ 'C',
                           HOLC_D > 0.5 & !is.na(HOLC_D) ~ 'D')) |>

  group_by(grade) |>
  summarize(mean_blackmobil = mean(kfr_black_pooled_p25, na.rm = TRUE),
            mean_whitemobil = mean(kfr_white_pooled_p25, na.rm = TRUE)) |>
  ggplot(aes(x = grade, y = mean_whitemobil, fill = grade)) +
  geom_bar(stat = "identity", show.legend = TRUE, width = 0.6) +
  scale_fill_manual(values = c("tomato1", "tomato2", "tomato3", "tomato4")) +
  labs(x = "",
       y = "Absolute Mobility at the 25th Percentile",
       title = "White Absolute Mobility by HOLC Grade",
```

```
fill = "HOLC Grade") +
theme_clean()
```

```
whitemobilgraph
```



```
ggsave("whitemobilgraph.png")
```

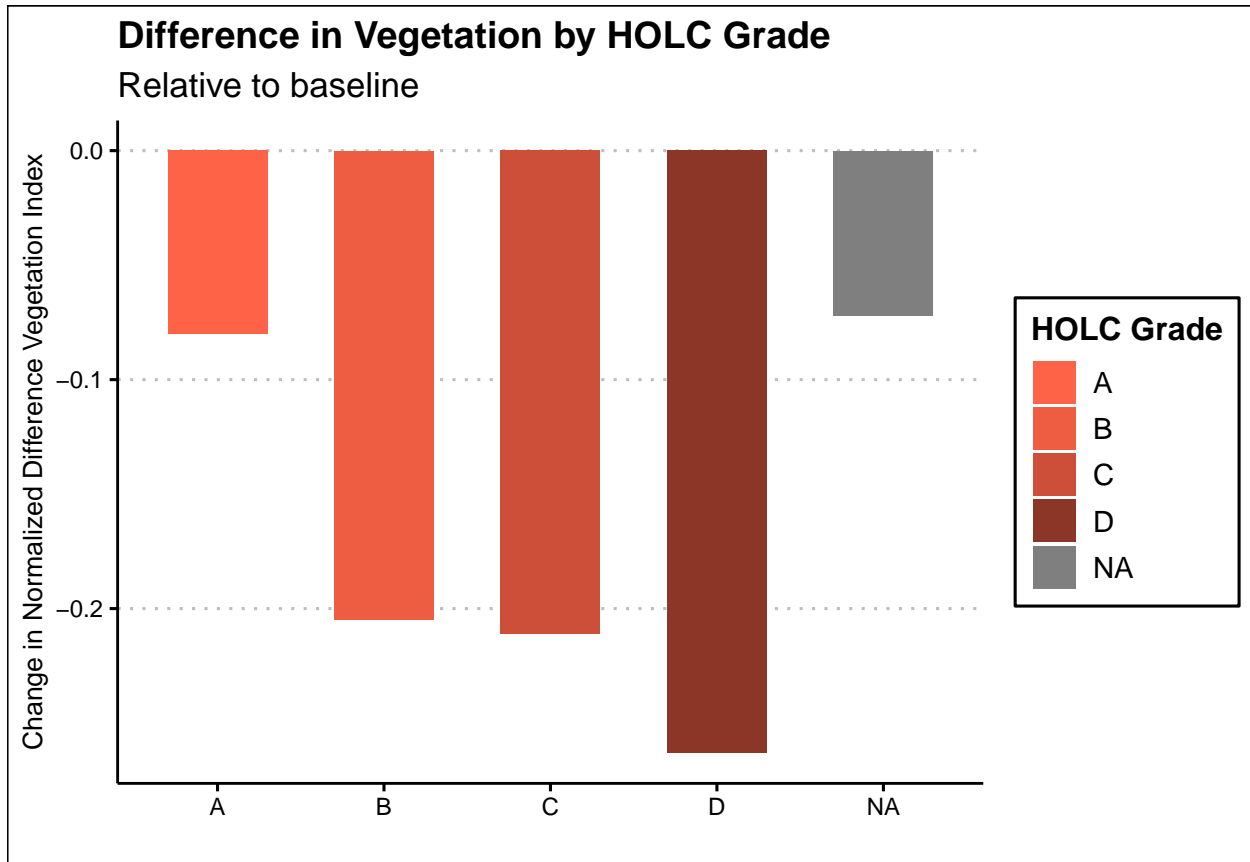
```
## Saving 6.5 x 4.5 in image
```

```
#Visualizing vegetation by HOLC neighborhood grade
```

```
veggraph <- atlas |>
  mutate(grade = case_when(HOLC_A > 0.5 & !is.na(HOLC_A) ~ 'A',
                           HOLC_B > 0.5 & !is.na(HOLC_B) ~ 'B',
                           HOLC_C > 0.5 & !is.na(HOLC_C) ~ 'C',
                           HOLC_D > 0.5 & !is.na(HOLC_D) ~ 'D')) |>
  group_by(grade) |>
  summarize(mean_veg = mean(vegetation, na.rm = TRUE),
            mean_heat = mean(extreme_heat, na.rm = TRUE),
            mean_developed = mean(developed, na.rm = TRUE)) |>
  ggplot(aes(x = grade, y = mean_veg, fill = grade)) +
  geom_bar(stat = "identity", show.legend = TRUE, width = 0.6) +
  scale_fill_manual(values = c("tomato1", "tomato2", "tomato3", "tomato4")) +
  labs(x = "",
       y = "Change in Normalized Difference Vegetation Index",
       title = "Difference in Vegetation by HOLC Grade",
       subtitle = "Relative to baseline",
```

```
fill = "HOLC Grade") +
theme_clean()
```

veggraph



```
ggsave("veggraph.png")
```

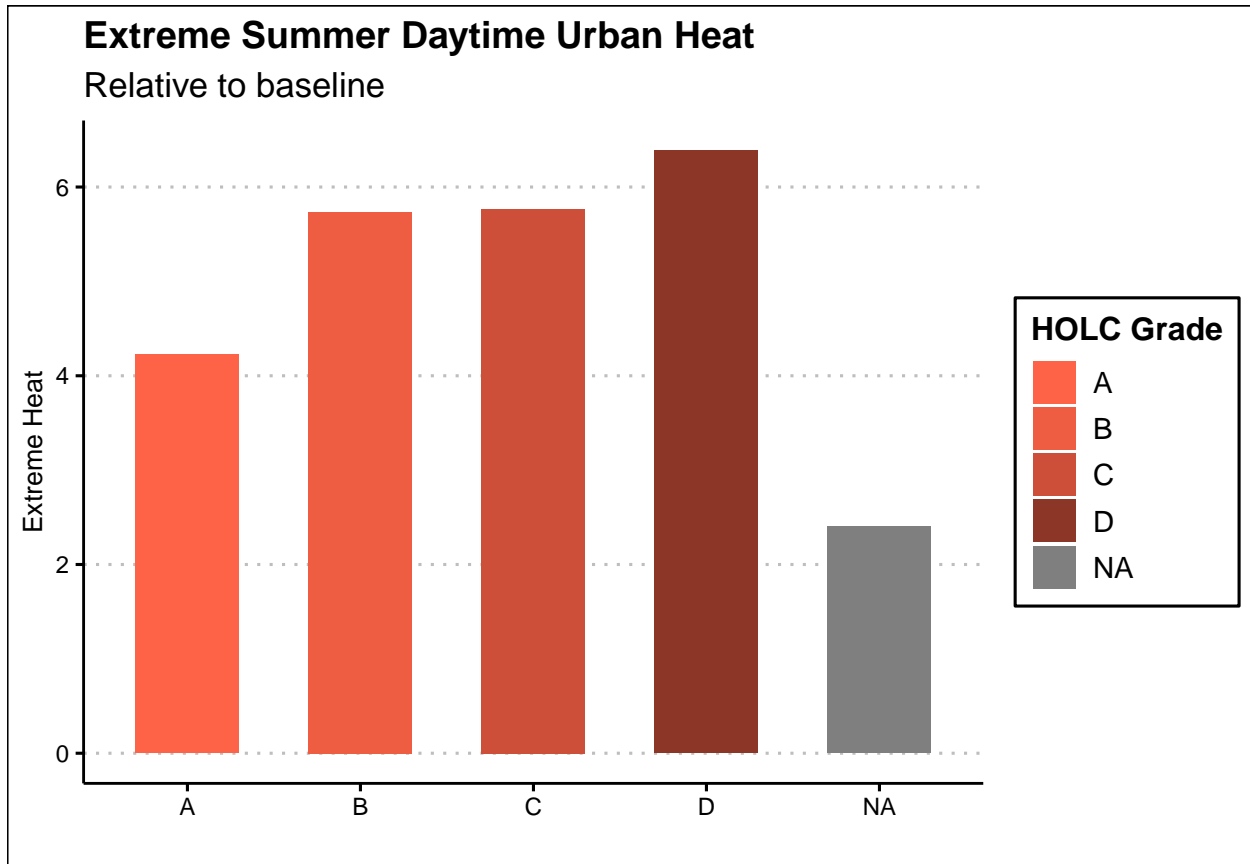
```
## Saving 6.5 x 4.5 in image
```

```
#Visualizing extreme summer daytime heat by HOLC grade
```

```
heatgraph <- atlas |>
  mutate(grade = case_when(HOLC_A > 0.5 & !is.na(HOLC_A) ~ 'A',
                           HOLC_B > 0.5 & !is.na(HOLC_B) ~ 'B',
                           HOLC_C > 0.5 & !is.na(HOLC_C) ~ 'C',
                           HOLC_D > 0.5 & !is.na(HOLC_D) ~ 'D')) |>
  group_by(grade) |>
  summarize(mean_veg = mean(vegetation, na.rm = TRUE),
            mean_heat = mean(extreme_heat, na.rm = TRUE),
            mean_developed = mean(developed, na.rm = TRUE)) |>
  ggplot(aes(x = grade, y = mean_heat, fill = grade)) +
  geom_bar(stat = "identity", show.legend = TRUE, width = 0.6) +
  scale_fill_manual(values = c("tomato1", "tomato2", "tomato3", "tomato4")) +
  labs(x = "",
       y = "Extreme Heat",
       title = "Extreme Summer Daytime Urban Heat",
       subtitle = "Relative to baseline",
```

```
fill = "HOLC Grade") +
theme_clean()
```

heatgraph



```
ggsave("heatgraph.png")
```

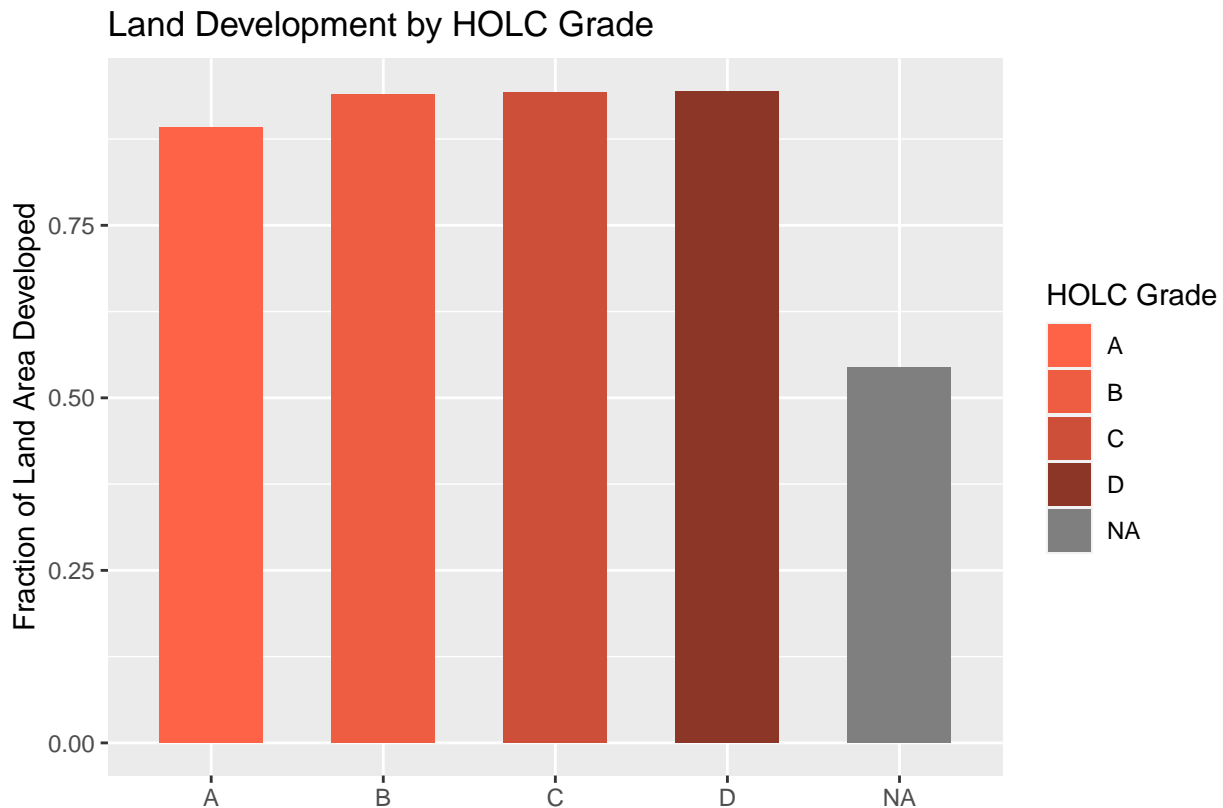
Saving 6.5 x 4.5 in image

#Visualizing land development by HOLC grade

```
devgraph <- atlas |>
  mutate(grade = case_when(HOLC_A > 0.5 & !is.na(HOLC_A) ~ 'A',
                           HOLC_B > 0.5 & !is.na(HOLC_B) ~ 'B',
                           HOLC_C > 0.5 & !is.na(HOLC_C) ~ 'C',
                           HOLC_D > 0.5 & !is.na(HOLC_D) ~ 'D')) |>
  group_by(grade) |>
  summarize(mean_veg = mean(vegetation, na.rm = TRUE),
            mean_heat = mean(extreme_heat, na.rm = TRUE),
            mean_developed = mean(developed, na.rm = TRUE)) |>
  ggplot(aes(x = grade, y = mean_developed, fill = grade)) +
  geom_bar(stat = "identity", show.legend = TRUE, width = 0.6) +
  scale_fill_manual(values = c("tomato1", "tomato2", "tomato3", "tomato4")) +
  labs(x = "",
       y = "Fraction of Land Area Developed",
       title = "Land Development by HOLC Grade",
       fill = "HOLC Grade")
```



```
devgraph
```



```
ggsave("devgraph.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
#Question 12: Tracking air pollution via particulate matter changes over time
```

```
pm82 <- mean(atlas$pm25_1982, na.rm = TRUE)
pm90 <- mean(atlas$pm25_1990, na.rm = TRUE)
pm00 <- mean(atlas$pm25_2000, na.rm = TRUE)
pm10 <- mean(atlas$pm25_2010, na.rm = TRUE)

airpollution <- data.frame(c("1982", "1990", "2000", "2010"),
                             c(pm82, pm90, pm00, pm10))
names(airpollution)[1] <- "Year"
names(airpollution)[2] <- "Particulate Matter Level"
airpollution
```

```
##   Year Particulate Matter Level
## 1 1982          20.410793
## 2 1990          16.844180
## 3 2000          12.500640
## 4 2010           9.286397
```

We see that air pollution, as measured by the particulate matter level data, has steadily decreased in the United States since 1982, with an average decrease of about 4 units fine particulate matter every 10 years.

#Q12B: Comparing air pollution in my home Census tract, state, and nationwide

```
myhood <- atlas %>% subset(state == "37" & county == "183" & tract == "53110") #creating data frame of
myhoodap <- mean(myhood$pm25_1990, na.rm = TRUE)
ncap <- mean(atlas$pm25_1990[atlas$state == "37"], na.rm = TRUE)
usap <- mean(atlas$pm25_1990, na.rm = TRUE)

apcomp <- data.frame(c("My Census Tract", "North Carolina", "United States"),
                     c(myhoodap, ncap, usap))
names(apcomp)[1] <- "Level"
names(apcomp)[2] <- "PM Concentration"
apcomp
```

```
##           Level PM Concentration
## 1 My Census Tract      17.37184
## 2 North Carolina      18.58069
## 3 United States       16.84418
```

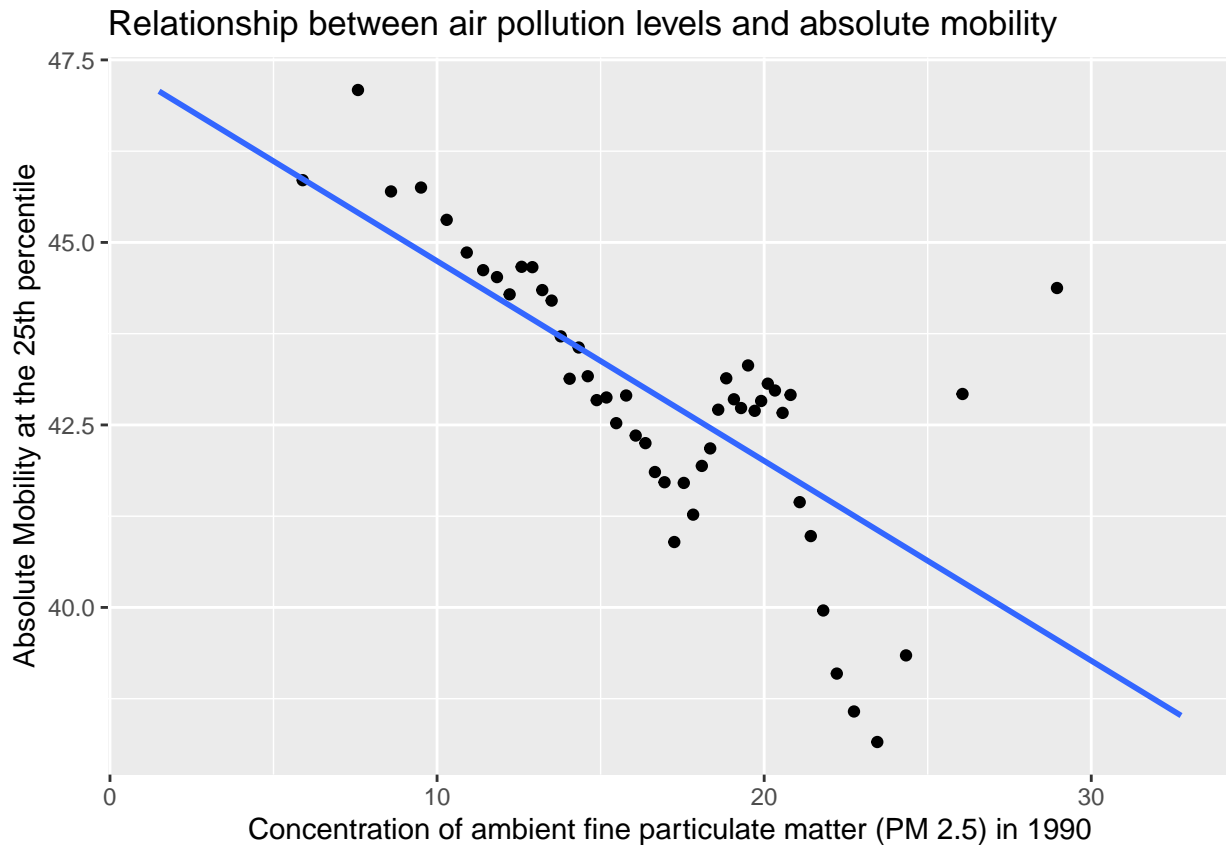
We see that the national air pollution average in 1990 was approximately 16.84 PM units, with higher concentrations of particulate matter for my home tract (17.37), and even higher for my home state of North Carolina (18.58).

#Q12C: Visualizing the relationship between absolute mobility and air pollution in 1990

```
scatterap <- atlas |>
  ggplot(aes(x = pm25_1990, y = kfr_pooled_pooled_p25)) +
  stat_binmean(n = 50) +
  stat_smooth(method = "lm", se = FALSE) +
  labs(title = "Relationship between air pollution levels and absolute mobility",
       x = "Concentration of ambient fine particulate matter (PM 2.5) in 1990",
       y = "Absolute Mobility at the 25th percentile")

scatterap
```

```
## Warning: Removed 1736 rows containing non-finite values (`stat_binmean()`).
## `geom_smooth()` using formula = 'y ~ x'
## Warning: Removed 1736 rows containing non-finite values (`stat_smooth()`).
```



```
ggsave("scatterap.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## Warning: Removed 1736 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 1736 rows containing non-finite values (`stat_smooth()`).
```

```
#Q12D: Deriving the correlation coefficient between absolute mobility and air pollution in 1990
```

```
cor(atlas$kfr_pooled_pooled_p25, atlas$pm25_1990, use = "complete.obs")
```

```
## [1] -0.1837617
```

The correlation coefficient between absolute mobility at the 25th percentile and air pollution levels in 1990 is approximately -0.18. This is smaller than the correlation at the county level. This makes sense because data grouped at the tract level leads to a larger sample size than data grouped at the county level. With a larger sample size, variability increases, leading to a reduced correlation coefficient.

```
#Q13A: Visualizing the relationship between strongly related covariates and absolute mobility
```

```
#Covariate 1: Incarceration Rate
```

```
wakeco <- atlas |>
```

```
  filter(state == 37,
```

```
         county == 183) #creating data frame of just my home county
```

```
scatterjail <- wakeco |>
```

```
  ggplot(aes(x = jail_pooled_pooled_p25, y = kfr_pooled_pooled_p25)) +
```

```

stat_binmean(n = 50) +
stat_smooth(method = "lm", se = FALSE) +
labs(title = "Relationship between incarceration rate and absolute mobility in Wake County",
      subtitle = "At the 25th percentile",
      x = "Fraction incarcerated for children from families at the 25th percentile, 2010",
      y = "Absolute Mobility at the 25th percentile")

```

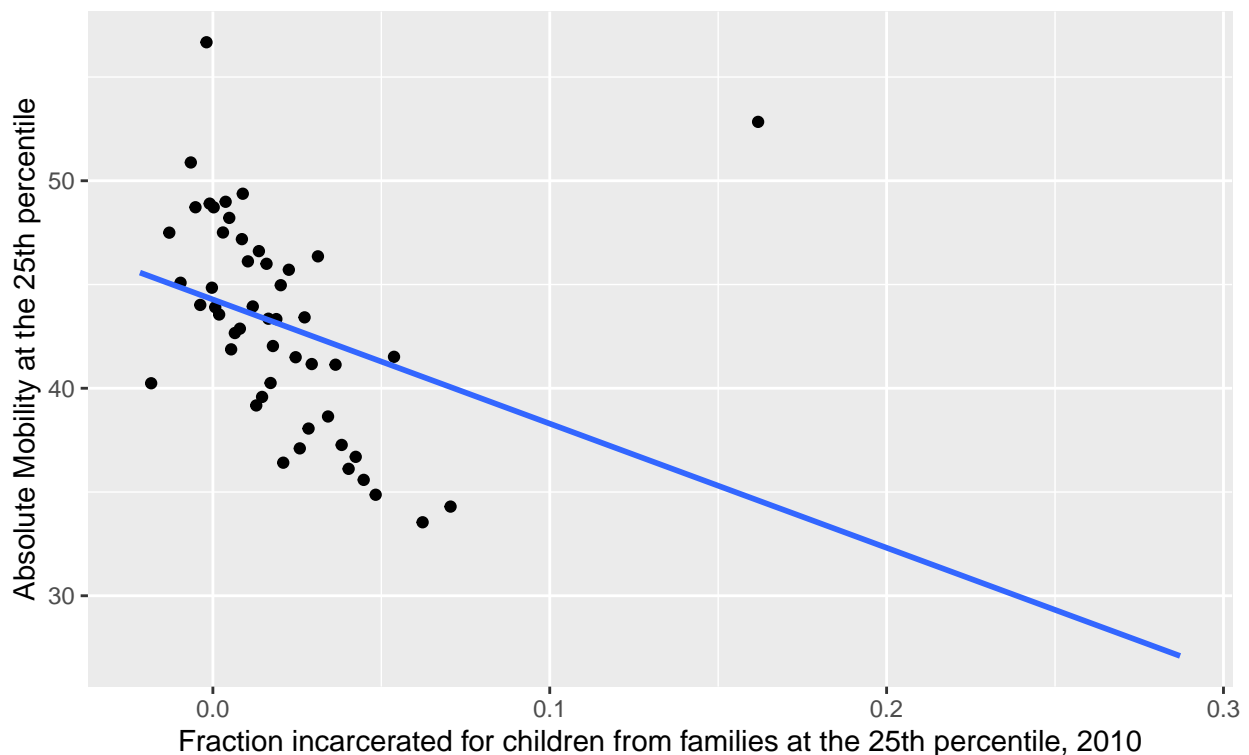
```
scatterjail
```

```
## Warning: Removed 2 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 2 rows containing non-finite values (`stat_smooth()`).
```

Relationship between incarceration rate and absolute mobility in Wake Cour
At the 25th percentile



```
ggsave("scatterjail.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## Warning: Removed 2 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 2 rows containing non-finite values (`stat_smooth()`).
```

```
#Covariate 2: Median household income
```

```

scatterincome <- wakeco |>
  ggplot(aes(x = med_hhinc1990, y = kfr_pooled_pooled_p25)) +
  stat_binmean(n = 50) +

```

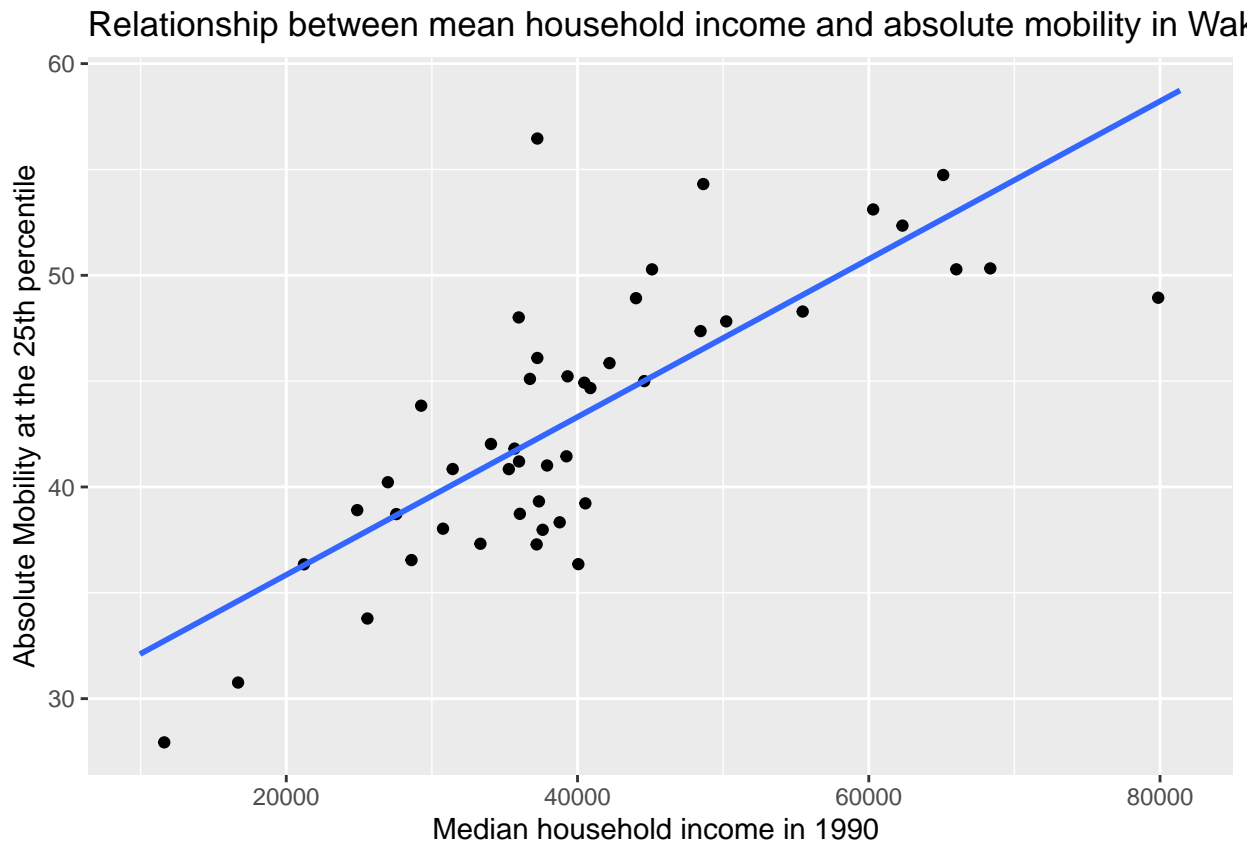
```
stat_smooth(method = "lm", se = FALSE) +
labs(title = "Relationship between mean household income and absolute mobility in Wake County",
      x = "Median household income in 1990 ",
      y = "Absolute Mobility at the 25th percentile")
```

```
scatterincome
```

```
## Warning: Removed 2 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 2 rows containing non-finite values (`stat_smooth()`).
```



```
ggsave("scatterincome.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## Warning: Removed 2 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 2 rows containing non-finite values (`stat_smooth()`).
```

```
#Covariate 3: Average Commute Time in 2000
```

```
scattercommute <- wakeco |>
  ggplot(aes(x = mean_commutetime2000, y = kfr_pooled_pooled_p25)) +
  stat_binmean(n = 50) +
  stat_smooth(method = "lm", se = FALSE) +
  labs(title = "Relationship between mean work commute time and absolute mobility in Wake County",
        x = "Average Commute Time of Working Adults in 2000",
```

```
y = "Absolute Mobility at the 25th percentile")
```

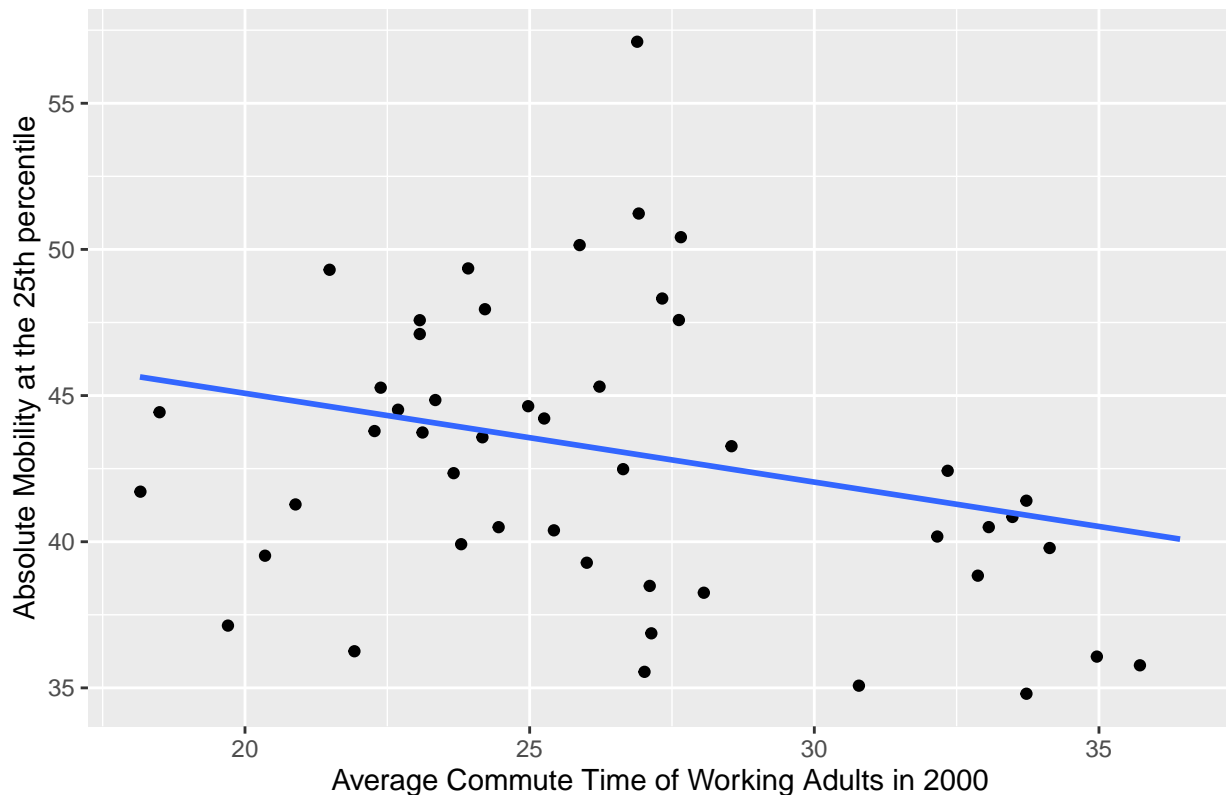
```
scattercommute
```

```
## Warning: Removed 2 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 2 rows containing non-finite values (`stat_smooth()`).
```

Relationship between mean work commute time and absolute mobility in Wake County



```
ggsave("scattercommute.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## Warning: Removed 2 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 2 rows containing non-finite values (`stat_smooth()`).
```

#Q13B: Reporting correlation coefficients of covariates with absolute mobility

```
jailcorcoef <- cor(wakeco$kfr_pooled_pooled_p25, wakeco$jail_pooled_pooled_p25, use = "complete.obs")
```

```
inccorcoef <- cor(wakeco$kfr_pooled_pooled_p25, wakeco$med_hhinc1990, use = "complete.obs")
```

```
commutecorcoef <- cor(wakeco$kfr_pooled_pooled_p25, wakeco$mean_commutetime2000, use = "complete.obs")
```

```
coefcomp <- data.frame(c("Incarceration Rate 2010", "Median Household Income 1990", "Mean Commute Time 2000"),
  c(jailcorcoef, inccorcoef, commutecorcoef))
```

```
names(coefcomp)[1] <- "Covariate"
```

```
names(coefcomp)[2] <- "Correlation Coefficient with absolute mobility"
```

```
coefcomp
```

```
##              Covariate Correlation Coefficient with absolute mobility
## 1      Incarceration Rate 2010                                -0.2266184
## 2 Median Household Income 1990                                0.6268116
## 3      Mean Commute Time 2000                                -0.1786249
```

We see that there are negative correlations between the incarceration rate and the mean commute time and absolute mobility, respectively, and a fairly strong positive correlation between median household income (in 1990) and absolute mobility.

#Q14: Examining the relationship between covariates and absolute mobility by race and gender

#Incarceration Rate: We will look at incarceration rates for Black males, Hispanic males, and White males

#Black Males

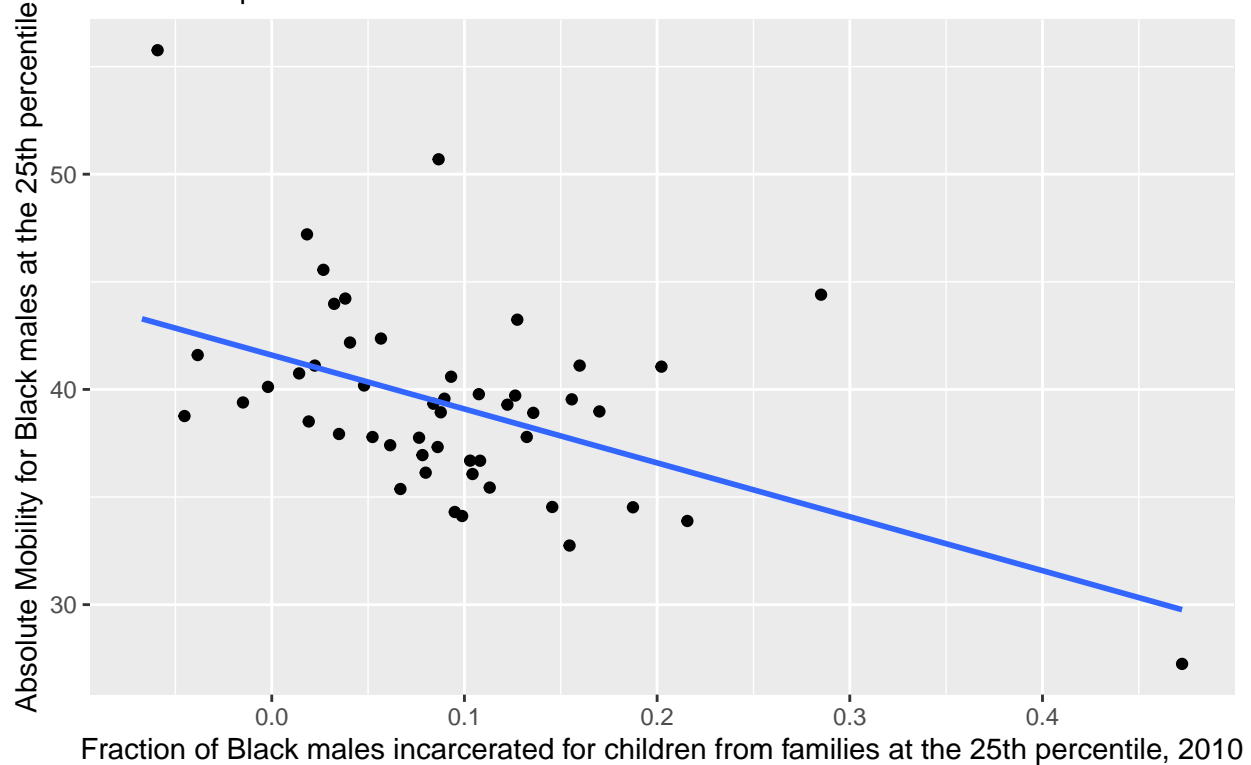
```
jailblack <- wakeco |>
  ggplot(aes(x = jail_black_male_p25, y = kir_black_male_p25)) +
  stat_binmean(n = 50) +
  stat_smooth(method = "lm", se = FALSE) +
  labs(title = "Relationship between incarceration rate and absolute mobility for Black males",
        subtitle = "At the 25th percentile",
        x = "Fraction of Black males incarcerated for children from families at the 25th percentile",
        y = "Absolute Mobility for Black males at the 25th percentile")
jailblack
```

```
## Warning: Removed 88 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 88 rows containing non-finite values (`stat_smooth()`).
```

Relationship between incarceration rate and absolute mobility for Black males
At the 25th percentile



```
ggsave("jailblack.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## Warning: Removed 88 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 88 rows containing non-finite values (`stat_smooth()`).
```

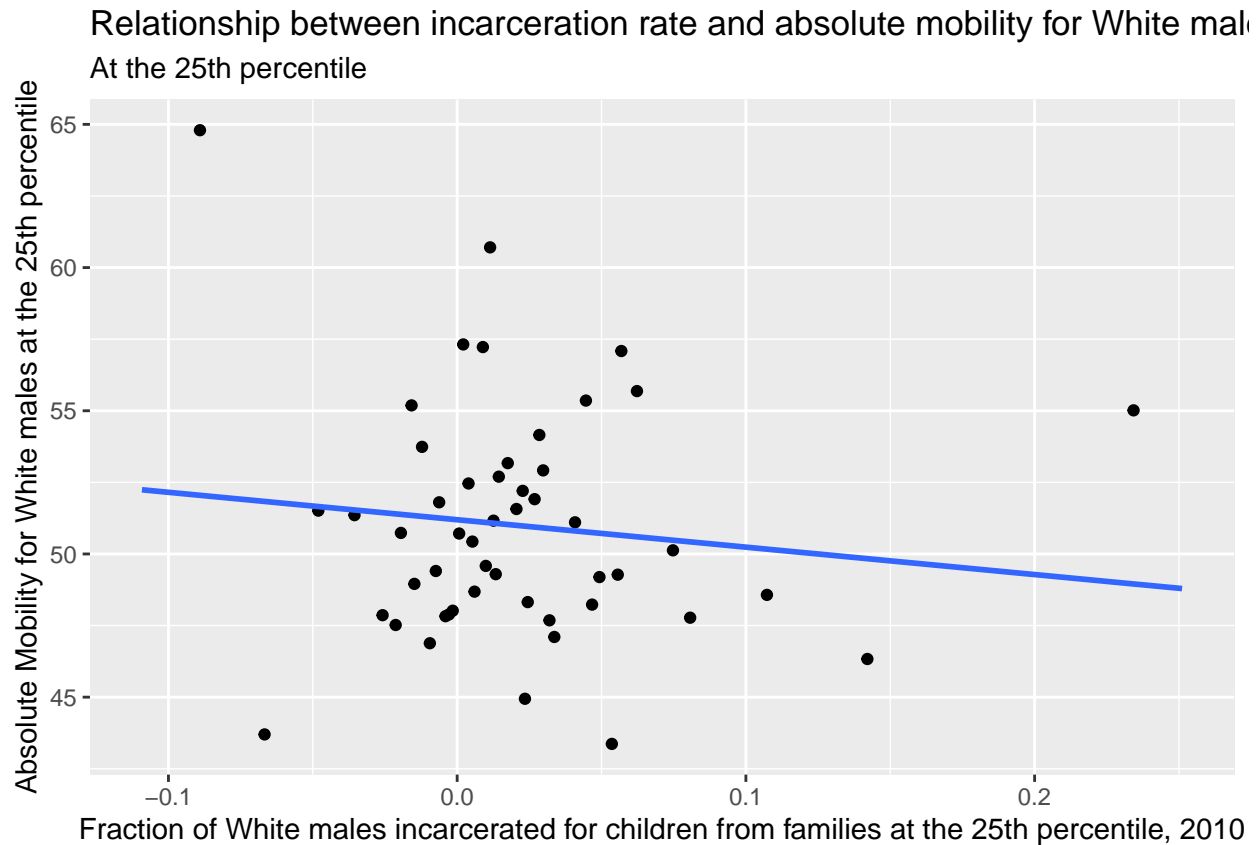
```
#White Males
```

```
jailwhite <- wakeco |>
  ggplot(aes(x = jail_white_male_p25, y = kir_white_male_p25)) +
  stat_binmean(n = 50) +
  stat_smooth(method = "lm", se = FALSE) +
  labs(title = "Relationship between incarceration rate and absolute mobility for White males",
        subtitle = "At the 25th percentile",
        x = "Fraction of White males incarcerated for children from families at the 25th percentile",
        y = "Absolute Mobility for White males at the 25th percentile")
jailwhite
```

```
## Warning: Removed 14 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 14 rows containing non-finite values (`stat_smooth()`).
```

```
ggsave("jailwhite.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## Warning: Removed 14 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 14 rows containing non-finite values (`stat_smooth()`).
```

```
#Hispanic Males
```

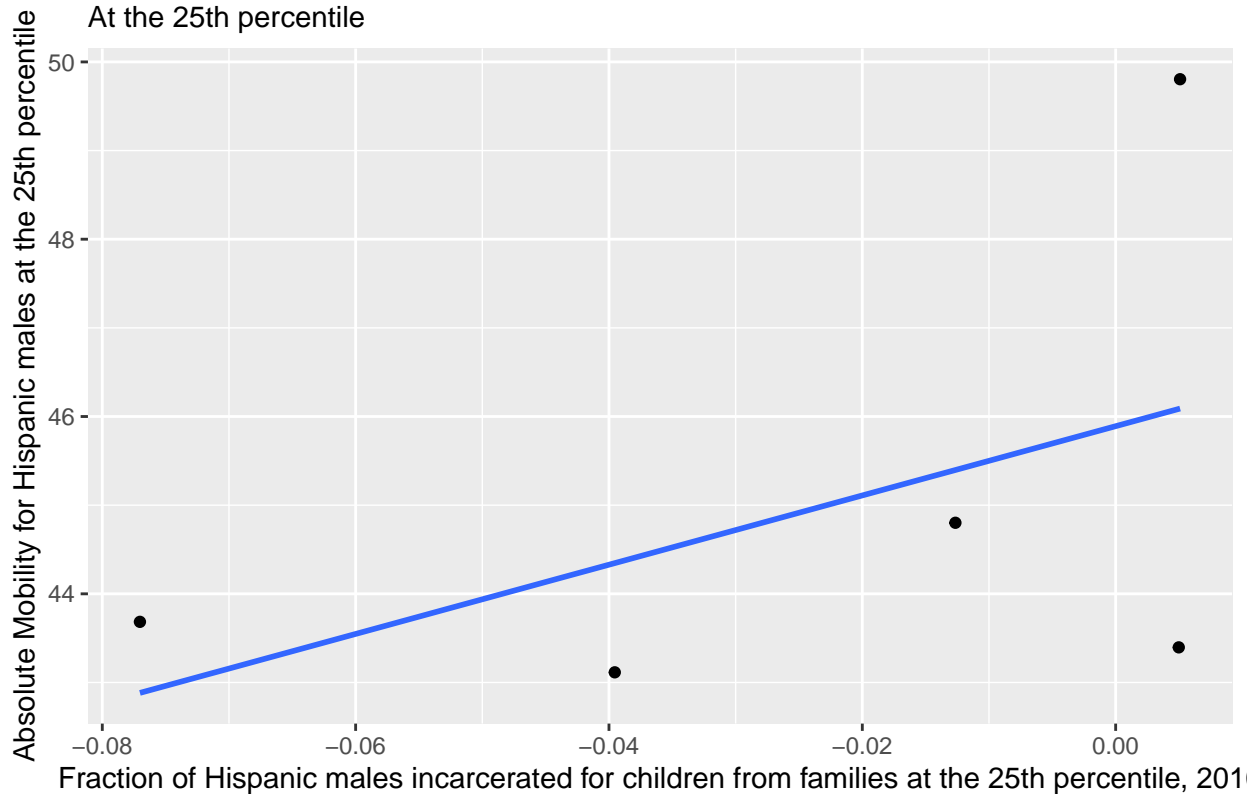
```
jailhisp <- wakeco |>
  ggplot(aes(x = jail_hisp_male_p25, y = kir_hisp_male_p25)) +
  stat_binmean(n = 50) +
  stat_smooth(method = "lm", se = FALSE) +
  labs(title = "Relationship between incarceration rate and absolute mobility for Hispanic males",
        subtitle = "At the 25th percentile",
        x = "Fraction of Hispanic males incarcerated for children from families at the 25th percentile",
        y = "Absolute Mobility for Hispanic males at the 25th percentile")
jailhisp
```

```
## Warning: Removed 181 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 181 rows containing non-finite values (`stat_smooth()`).
```

Relationship between incarceration rate and absolute mobility for Hispanic n
At the 25th percentile



```
ggsave("jailhisp.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## Warning: Removed 181 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 181 rows containing non-finite values (`stat_smooth()`).
```

#Median Household Income: Next, we will look at median household income in 1990 and absolute mobility f

#Black Children

```
blackincome <- wakeco |>
```

```
  ggplot(aes(x = med_hhinc1990, y = kfr_black_pooled_p25)) +
```

```
    stat_binmean(n = 50) +
```

```
    stat_smooth(method = "lm", se = FALSE) +
```

```
    labs(title = "Relationship between mean household income and absolute mobility for Black p
```

```
          x = "Median household income in 1990 ",
```

```
          y = "Absolute Mobility for Black people at the 25th percentile")
```

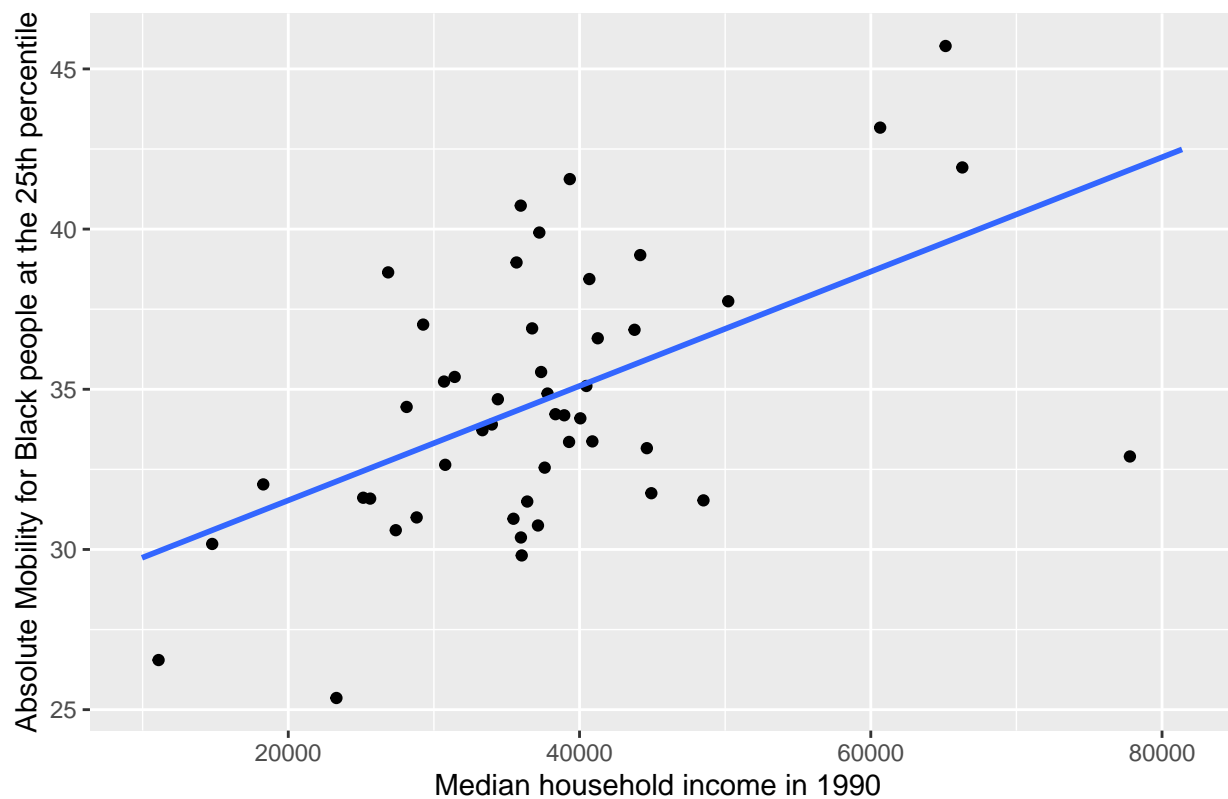
```
blackincome
```

```
## Warning: Removed 36 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 36 rows containing non-finite values (`stat_smooth()`).
```

Relationship between mean household income and absolute mobility for Bla



```
ggsave("blackincome.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## Warning: Removed 36 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 36 rows containing non-finite values (`stat_smooth()`).
```

```
#White Children
```

```
whiteincome <- wakeco |>
```

```
  ggplot(aes(x = med_hhinc1990, y = kfr_white_pooled_p25)) +
```

```
    stat_binmean(n = 50) +
```

```
    stat_smooth(method = "lm", se = FALSE) +
```

```
    labs(title = "Relationship between mean household income and absolute mobility for White p
```

```
          x = "Median household income in 1990 ",
```

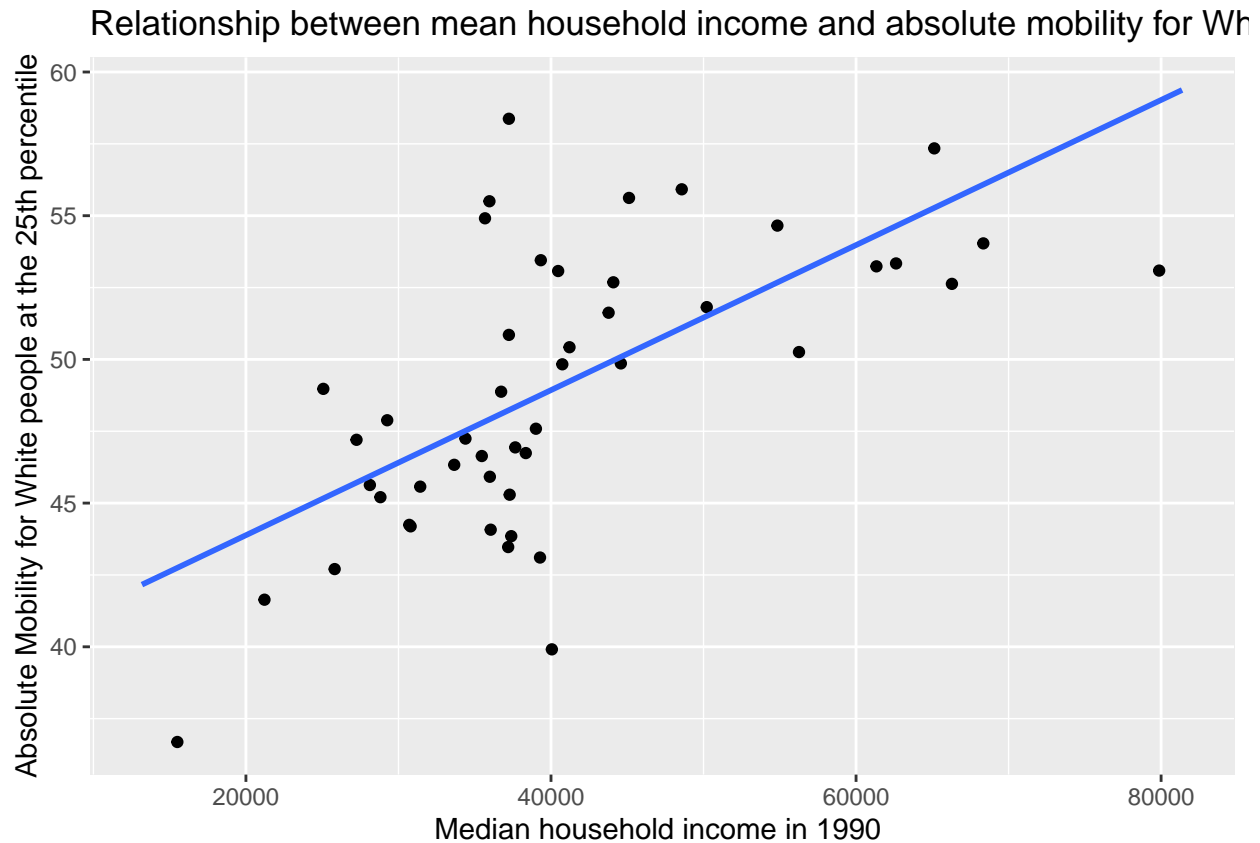
```
          y = "Absolute Mobility for White people at the 25th percentile")
```

```
whiteincome
```

```
## Warning: Removed 8 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 8 rows containing non-finite values (`stat_smooth()`).
```



```
ggsave("whiteincome.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## Warning: Removed 8 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 8 rows containing non-finite values (`stat_smooth()`).
```

```
#Hispanic Children
```

```
hispincome <- wakeco |>
```

```
  ggplot(aes(x = med_hhinc1990, y = kfr_hisp_pooled_p25)) +
```

```
    stat_binmean(n = 50) +
```

```
    stat_smooth(method = "lm", se = FALSE) +
```

```
    labs(title = "Relationship between mean household income and absolute mobility for Hispanics",
```

```
          x = "Median household income in 1990 ",
```

```
          y = "Absolute Mobility for Hispanics at the 25th percentile")
```

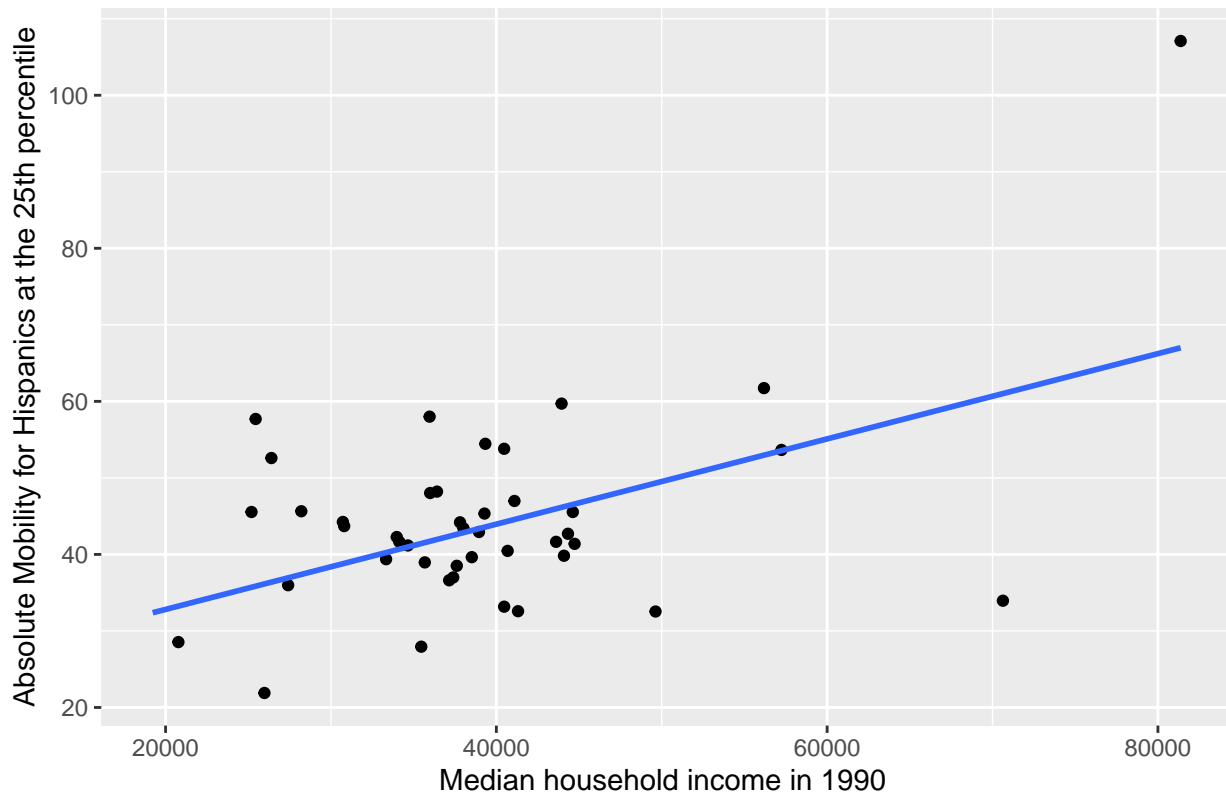
```
hispincome
```

```
## Warning: Removed 127 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 127 rows containing non-finite values (`stat_smooth()`).
```

Relationship between mean household income and absolute mobility for Hi



```
ggsave("hispincome.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## Warning: Removed 127 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 127 rows containing non-finite values (`stat_smooth()`).
```

```
#Asian Children
```

```
asianincome <- wakeco |>
```

```
  ggplot(aes(x = med_hhinc1990, y = kfr_asian_pooled_p25)) +
```

```
    stat_binmean(n = 50) +
```

```
    stat_smooth(method = "lm", se = FALSE) +
```

```
    labs(title = "Relationship between mean household income and absolute mobility for Asians :")
```

```
          x = "Median household income in 1990",
```

```
          y = "Absolute Mobility for Asians at the 25th percentile")
```

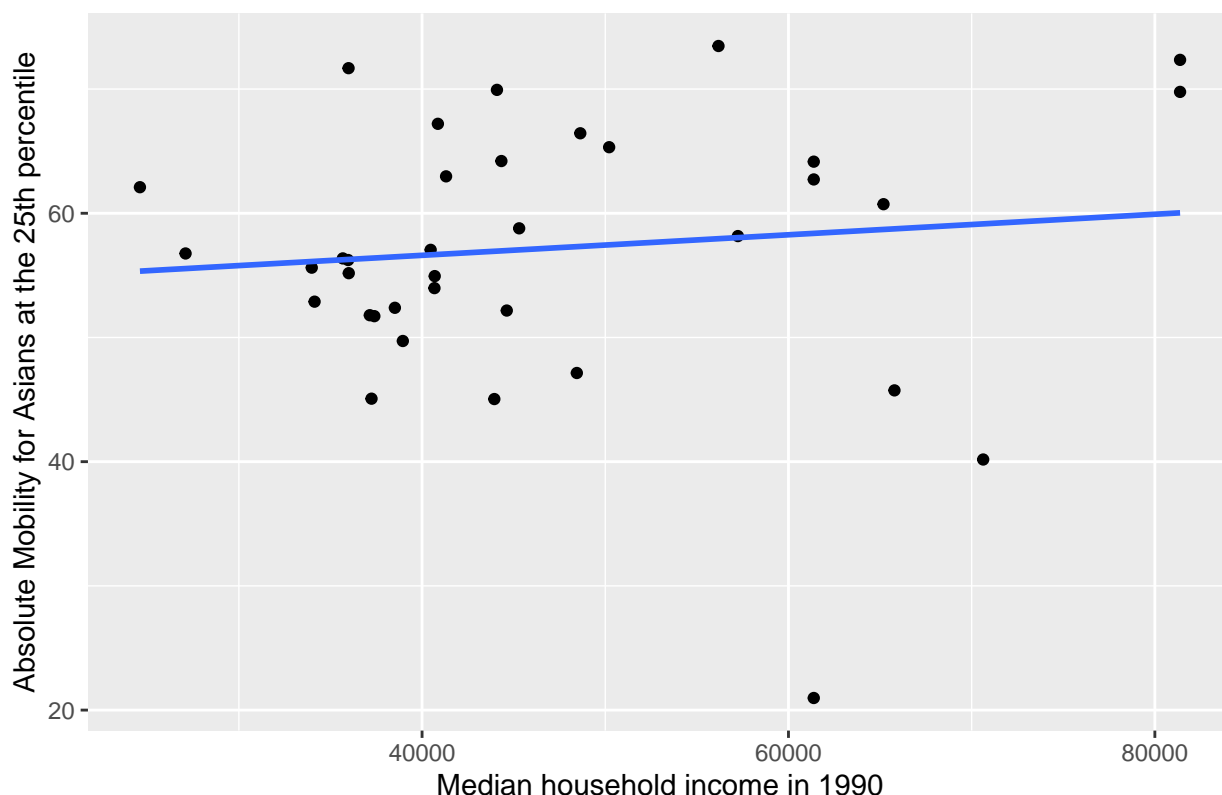
```
asianincome
```

```
## Warning: Removed 140 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 140 rows containing non-finite values (`stat_smooth()`).
```

Relationship between mean household income and absolute mobility for Asi



```
ggsave("asianincome.png")
```

```
## Saving 6.5 x 4.5 in image
```

```
## Warning: Removed 140 rows containing non-finite values (`stat_binmean()`).
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

```
## Warning: Removed 140 rows containing non-finite values (`stat_smooth()`).
```

#Q14 part 2: Reporting correlation coefficients of covariates with absolute mobility across race and ge

```
jailblackcorcoef <- cor(wakeco$kir_black_male_p25, wakeco$jail_black_male_p25, use = "complete.obs")
```

```
jailwhitecorcoef <- cor(wakeco$kir_white_male_p25, wakeco$jail_white_male_p25, use = "complete.obs")
```

```
jailhispcoef <- cor(wakeco$kir_hisp_male_p25, wakeco$jail_hisp_male_p25, use = "complete.obs")
```

```
incblackcorcoef <- cor(wakeco$kfr_black_pooled_p25, wakeco$med_hhinc1990, use = "complete.obs")
```

```
incwhitecorcoef <- cor(wakeco$kfr_white_pooled_p25, wakeco$med_hhinc1990, use = "complete.obs")
```

```
inchispcorcoef <- cor(wakeco$kfr_hisp_pooled_p25, wakeco$med_hhinc1990, use = "complete.obs")
```

```
incasiancorcoef <- cor(wakeco$kfr_asian_pooled_p25, wakeco$med_hhinc1990, use = "complete.obs")
```

```
coefcomp_incarcerate <- data.frame(c("Incarceration Rate for Black Males", "Incarceration Rate for White Males",  
                                     c(jailblackcorcoef, jailwhitecorcoef, jailhispcoef))
```

```
names(coefcomp_incarcerate)[1] <- "Incarceration Rate for Racial Group"
```

```
names(coefcomp_incarcerate)[2] <- "Correlation Coefficient with Absolute Mobility for that Racial Group"
```

```
coefcomp_incarcerate
```

```
##      Incarceration Rate for Racial Group
```

```
## 1      Incarceration Rate for Black Males
```

```
## 2      Incarceration Rate for White Males
```

```
## 3      Incarceration Rate for Hispanic Males
```

```
## Correlation Coefficient with Absolute Mobility for that Racial Group
## 1 -0.33926569
## 2 -0.05986192
## 3 0.48964213

coefcomp_medinc <- data.frame(c("Median Household Income 1990", "Median Household Income 1990", "Median
                                c("Black Absolute Mobility", "White Absolute Mobility", "Hispanic Absolute
                                c(incblackcorcoef, incwhitecorcoef, inchispcorcoef, incasiancorcoef))

names(coefcomp_medinc)[1] <- "Covariate"
names(coefcomp_medinc)[2] <- "Absolute Mobility Group"
names(coefcomp_medinc)[3] <- "Correlation Coefficient with Absolute Mobility for that Racial Group"
coefcomp_medinc

## Covariate Absolute Mobility Group
## 1 Median Household Income 1990 Black Absolute Mobility
## 2 Median Household Income 1990 White Absolute Mobility
## 3 Median Household Income 1990 Hispanic Absolute Mobility
## 4 Median Household Income 1990 Asian Absolute Mobility
## Correlation Coefficient with Absolute Mobility for that Racial Group
## 1 0.31632544
## 2 0.45236858
## 3 0.45840067
## 4 0.08887583
```

When examining the relationship between incarceration rates and absolute mobility across race and gender, we see a fairly strong negative correlation coefficient between the black male incarceration rate and the absolute mobility for black males (-0.33). We see a weaker negative correlation coefficient between said covariates for white males (-0.05), and the sample size for hispanic males in Wake County is too small for us to consider the correlation coefficient for that racial group (0.48).

There is a clear pattern of fairly strong positive correlation coefficients between the median household income in 1990 and absolute mobility ranks for Blacks, Whites, Hispanics, and Asians. However, we should discount the correlation coefficient for Asian absolute mobility due to its small sample size. We should also note the difference between the correlation coefficient for Blacks' absolute mobility and the median household income (0.31) and that for Whites and Hispanics (0.452 and 0.458 respectively) - suggesting that, at least for Black children growing up in Wake County, there isn't as strong of a relationship between median household and their absolute mobility outcomes.