

Food Environment and Health

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Data and Methods

Data Source and Collection

1. NYC restaurant inspection:

<https://data.cityofnewyork.us/Health/DOHMH-New-York-City-Restaurant-Inspection-Results/43nn-pn8j>

This dataset contains the data for restaurant inspection in NYC from August 1, 2014 to June 9, 2019. Every row is a restaurant inspection record that includes the name of the restaurant, zipcode, cuisine type description, inspection grade (A as the best grade) and so on.

Variable used in this datasets

DBA: Name of the restaurant

BORO: name of the boro

ZIPCODE: the zipcode of the restaurant

CUISINE DESCRIPTION: the kind of food that the restaurant is providing

GRADE: Inspection grade for the specific inspection.

2. 2015 Community Health Profiles Open Data:

<https://www1.nyc.gov/site/doh/data/data-publications/profiles.page>

This dataset contains NYC every neighborhoods' demographic (percentage of white race, poverty percentage), health (age-adjusted percent of adult exercised in the last 30 days, age-adjusted percent of adults as a smoker) and our main outcome (Age-adjusted percent of adults that is obese (BMI of 30 or greater), Age-adjusted percent of adults, Age-adjusted rate of hospitalizations due to stroke (cerebrovascular disease) per 100,000 adults)

Variable used in this datasets

Name: the name for the neighborhood.

Racewhite_Rate: the percentage for white race

Poverty: Percent of individuals living below the federal poverty threshold

Smoking: age-adjusted percent of adults as a smoker Exercise: Age-adjusted percent of adults that reported getting any exercise in the last 30 days

Obesity: Age-adjusted percent of adults that is obese (BMI of 30 or greater) based on self-reported height and weight

Diabetes: Age-adjusted percent of adults that had ever been told by a healthcare professional that they have diabetes

Stroke_Hosp: Age-adjusted rate of hospitalizations due to stroke (cerebrovascular disease) per 100,000 adults

3. Web scraping for what zipcodes each neighborhood contains

Because Community Health Profiles Open Data has only neighborhood level information, so we have to aggregate neighborhood level information from the restaurant inspection data. However, the restaurant inspection data is using zipcodes for every restaurant. Therefore, we have to add the neighborhood name to every restaurant, so that we can group by neighborhood then calculate the percentage for every neighborhood.

First, we scraped the table data from <https://www.health.ny.gov/statistics/cancer/registry/appendix/neighborhoods.htm>. However the neighborhood name and combinations are different from the health profile data we downloaded. We could not merge the two datasets. Then we tried to look for the raw classification for the neighborhoods in health data from New York University's Furman Center for Real

Estate and Urban Policy and the NYC Department of City Planning. However, still no luck because the neighborhood is not divided by zipcode areas. Finally we tried the hardest way: searching the name of the neighborhood on Google and finding the matching zipcodes. Although it is not precise because neighborhoods are not divided according to the area of the zipcode (different neighborhoods sometimes share the same zipcodes), it works out well.

To deal with this issue and get an unbiased result, we randomly assign the zipcode to only one neighborhood if two or more neighborhoods are in the same zipcode area.

Data Import and Cleaning

First, we download restaurant inspection data from NYC open data.

```
get_all_inspections = function(url) {

  all_inspections = vector("list", length = 0)

  loop_index = 1
  chunk_size = 50000
  DO_NEXT = TRUE

  while (DO_NEXT) {
    message("Getting data, page ", loop_index)

    all_inspections[[loop_index]] =
      GET(url,
        query = list(`$order` = "zipcode",
                      `$limit` = chunk_size,
                      `$offset` = as.integer((loop_index - 1) * chunk_size)
                    )
      ) %>%
      content("text") %>%
      fromJSON() %>%
      as_tibble()

    DO_NEXT = dim(all_inspections[[loop_index]])[1] == chunk_size
    loop_index = loop_index + 1
  }

  all_inspections
}

url = "https://data.cityofnewyork.us/resource/9w7m-hzhe.json"

rest_inspection = get_all_inspections(url) %>%
  bind_rows()

# changing to date
rest_inspection = rest_inspection %>%
  mutate(inspection_date = rest_inspection$inspection_date %>%
    strtrim(., nchar(.)-13) %>%
    as.Date() )
rest_inspection$inspection_date %>%
```

```
max()
```

```
## [1] "2019-06-07"
```

Then, we download health and demographic data CSV into local folder, read in and clean it.

```
download.file("https://www1.nyc.gov/assets/doh/downloads/excel/episrv/2015_CHP_PUD.xlsx", mode="wb", de
health <- read_excel("health.xlsx", sheet = "CHP_all_data") %>%
  select(Name, Racewhite_Rate, Poverty, Unemployment,
         Smoking, Exercise,
         Obesity, Diabetes, Stroke_Hosp, Airquality_rate) %>%
  clean_names()
```

Lastly, we match the neighborhood names with restaurant zipcodes.

```
zip_neighbor <- read_csv("neigh_zipcode.csv") %>%
  mutate(zipcode = as.character(zipcode))
##restaurant data with neighbourhood
rest_neighborhood = left_join(rest_inspection, zip_neighbor, by = "zipcode") %>%
  filter(!is.na(neighborhood))
```

Exploratory Analysis

Health Data

The health data we achieved is very well-structured. We make plots to see the distribution of the three chronic disease outcomes across neighborhoods.

```
health_only_neighborhood <- health[-c(1:6),] %>%
  rename(neighborhood = name) %>%
  mutate(neighborhood = as.factor(neighborhood))

##Plotting for outcome in different neighborhood
bar_obesity <- health_only_neighborhood %>%
  mutate(neighborhood = fct_reorder(neighborhood, obesity)) %>%
  filter(obesity > 25) %>%
  ggplot(aes(x = neighborhood, y = obesity, fill = neighborhood)) + geom_col() +
  theme(axis.text.x = element_text(angle = 45, hjust = 1),
        legend.position = "none") +
  labs(title = "Neighborhood with 25 percent or more obesity rate")

bar_diabetes <- health_only_neighborhood %>%
  mutate(neighborhood = fct_reorder(neighborhood, diabetes)) %>%
  filter(diabetes > 10) %>%
  ggplot(aes(x = neighborhood, y = diabetes, fill = neighborhood)) + geom_col() +
  theme(axis.text.x = element_text(angle = 45, hjust = 1),
        legend.position = "none") +
  labs(title = "Neighborhood with 10 percent or more diabetes rate")

bar_stroke <- health_only_neighborhood %>%
  mutate(neighborhood = fct_reorder(neighborhood, stroke_hosp)) %>%
  filter(stroke_hosp > 300) %>%
  ggplot(aes(x = neighborhood, y = stroke_hosp, fill = neighborhood)) +
  geom_col() +
```

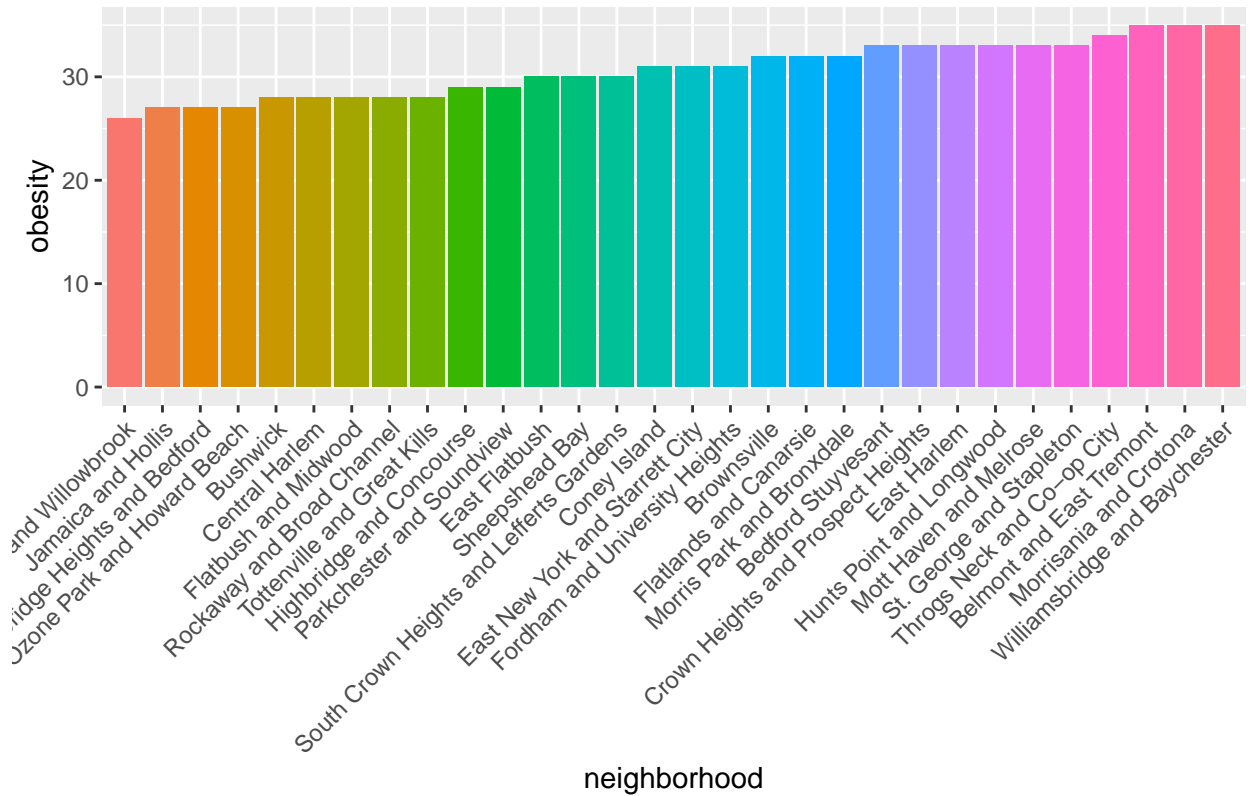
```

theme(axis.text.x = element_text(angle = 45, hjust = 1),
      legend.position = "none") +
labs(title = "Neighborhood with 300 or more stroke hospitalization in 100,000 adults")

#ggplotly(bar_obe)
#ggplotly(bar_dia)
#ggplotly(bar_stro)
bar_obe

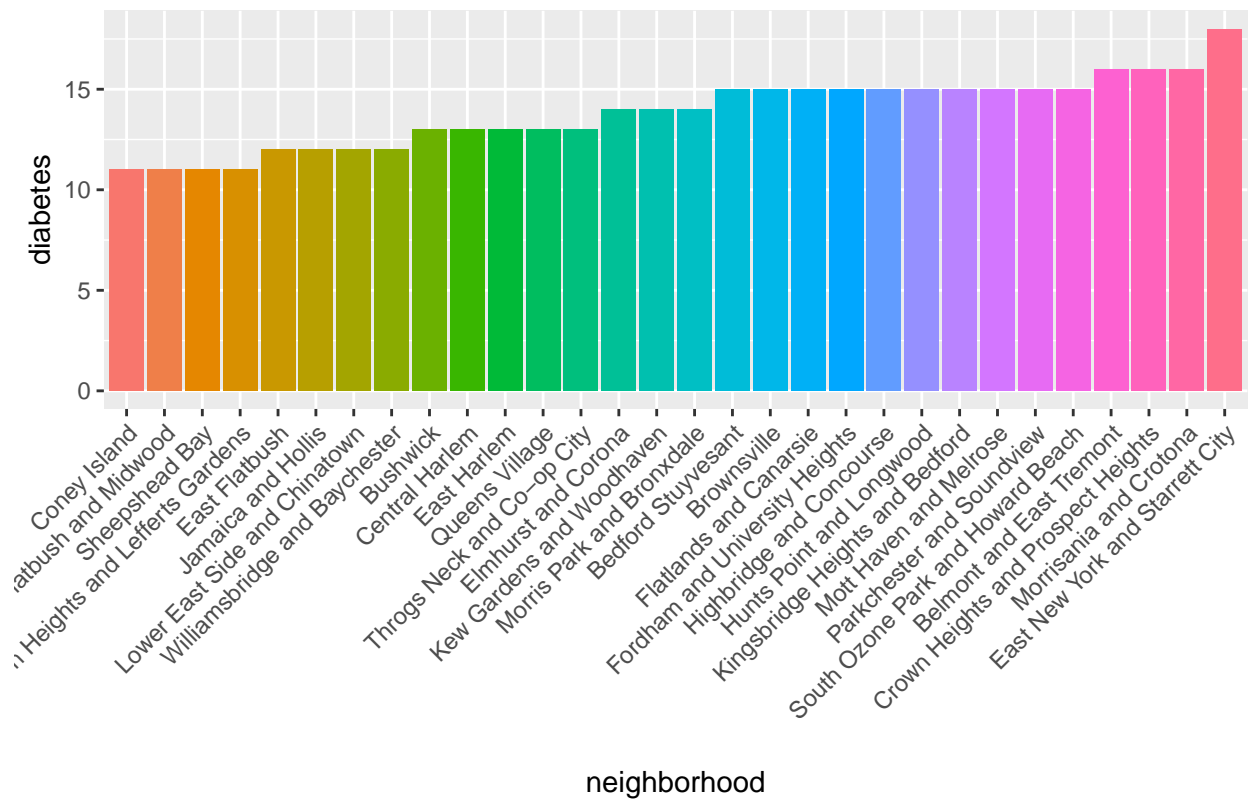
```

Neighborhood with 25 percent or more obesity rate



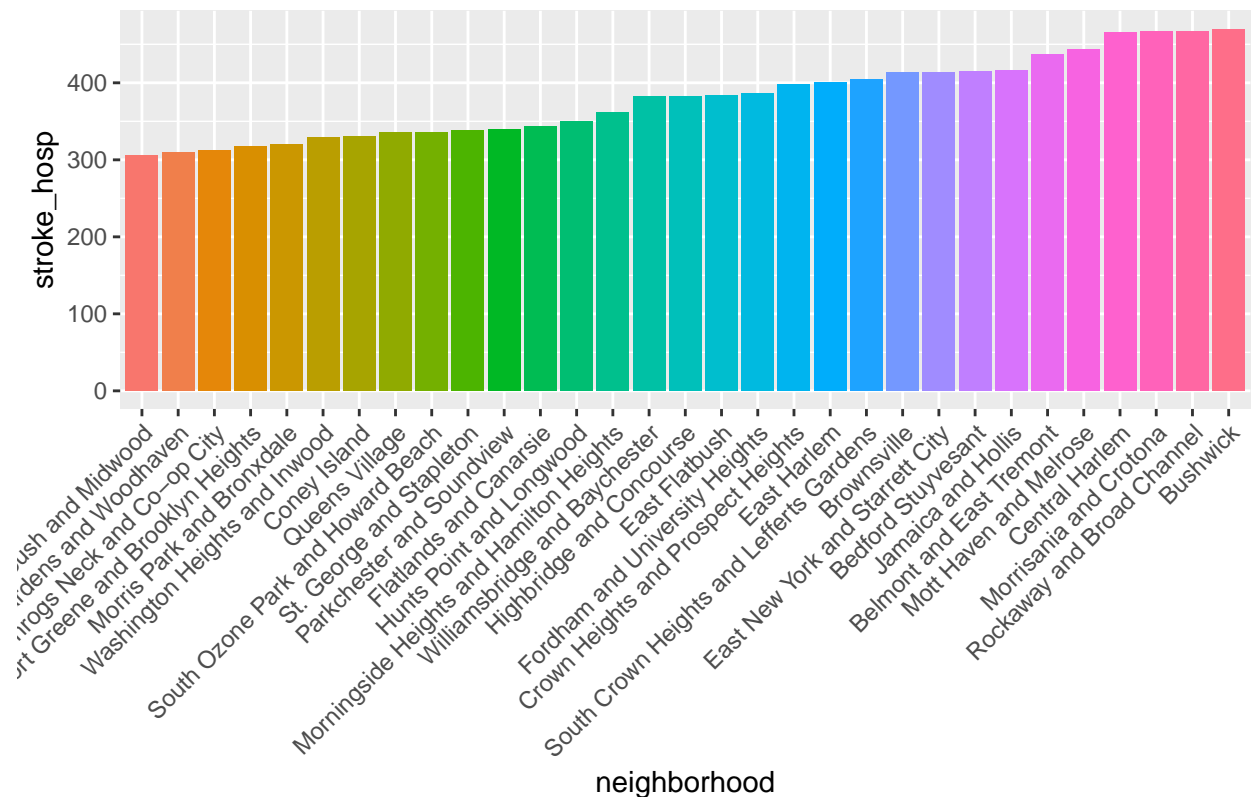
bar_dia

Neighborhood with 10 percent or more diabetes rate



bar_stro

Neighborhood with 300 or more stroke hospitalization in 100,000 adults



Williamsbridge and Baychester along with two other neighborhoods (Belmont and East Tremont, Morrisania and Crotona) have the highest obesity rate, up to 35%. East New York and Starrett City have the highest diabetes rate, up to 18 percent. Bushwick has the highest stroke hospitalization in 100,000 adults, up to 470 adults.

Restaurant data

Fastfood restaurants by cuisine type

Fastfood restaurants are identified by their cuisine descriptions given in the inspection data. We print out the cuisine descriptions list (n=85) and let everyone circle the ones they think is fastfood and the union are used as our rule (use union because it's more conservative).

We classify cuisine descriptions “Bagels/Pretzels”, “Bottled beverages, including water, sodas, juices, etc.”, “Chicken”, “Delicatessen”, “Donuts”, “Hamburgers”, “Hotdogs”, “Hotdogs/Pretzels”, “Ice Cream, Gelato, Yogurt, Ices”, “Nuts/Confectionary”, “Pancakes/Waffles”, “Pizza”, “Soul Food”, “Sandwiches”, “Sandwiches/Salads/Mixed Buffet” and “Soups & Sandwiches” as fastfood restaurants. And then we calculate the total number of restaurants and the number of fastfood restaurants, as well as the percentage of fastfood restaurants for each neighborhood.

```
# calculating the total number of restaurants and the number of fastfood restaurants in the neighborhood
neighborhood_list =
  rest_neighborhood %>%
  distinct(neighborhood) %>%
  arrange(neighborhood)

rest_fastfood_neighborhood =
```

```

rest_neighborhood %>%
  filter(cuisine_description %in% c("Bagels/Pretzels",
    "Bottled beverages, including water, sodas, juices, etc.",
    "Chicken",
    "Delicatessen",
    "Donuts",
    "Hamburgers",
    "Hotdogs",
    "Hotdogs/Pretzels",
    "Ice Cream, Gelato, Yogurt, Ices",
    "Nuts/Confectionary",
    "Pancakes/Waffles",
    "Pizza",
    "Soul Food",
    "Sandwiches",
    "Sandwiches/Salads/Mixed Buffet",
    "Soups & Sandwiches"))

percent_fastfood_neighborhood = function(name_neighborhood){

  rest_each_neighborhood =
    rest_neighborhood %>%
    filter(neighborhood == name_neighborhood) %>%
    distinct(camis)

  n_rest_neighborhood = nrow(rest_each_neighborhood)

  rest_fastfood_distinct_neighborhood =
    rest_fastfood_neighborhood %>%
    filter(neighborhood == name_neighborhood) %>%
    distinct(camis, cuisine_description)

  n_fastfood_neighborhood = nrow(rest_fastfood_distinct_neighborhood)

  percent_fastfood_neighborhood = n_fastfood_neighborhood/n_rest_neighborhood

  tibble(
    neighborhood = name_neighborhood,
    n_fastfood = n_fastfood_neighborhood,
    n_rest = n_rest_neighborhood,
    percent_fastfood = percent_fastfood_neighborhood
  )
}

fastfood_neighborhood =
  map(neighborhood_list$neighborhood, percent_fastfood_neighborhood) %>%
  bind_rows() %>%
  mutate(neighborhood = str_to_upper(neighborhood))

# plot for each neighborhood
# fastfood_neighborhood %>%
#   mutate(neighborhood = as.factor(neighborhood),
#     n_rest = as.numeric(n_rest),

```

```
#       n_nonfastfood = n_rest - n_fastfood,
#       neighborhood = fct_reorder(neighborhood, percent_fastfood)) %>%
# plot_ly(., x = ~neighborhood, y = ~n_fastfood, type = 'bar', name = 'fastfood restaurants') %>%
# add_trace(y = ~n_nonfastfood, name = 'non-fastfood restaurants') %>%
# layout(yaxis = list(title = 'Number of restaurants'),
#        xaxis = list(title = 'Neighborhood (ordered by percentage of fastfood restaurants)',
#                      showticklabels = FALSE),
#        barmode = 'stack')
```

While the Greenwich Village and SOHO neighborhood has fairly large number of restaurants, it has the smallest percentage of fastfood restaurants. Williamsbridge and Baychester has the largest percentage of fastfood restaurants.

When large number of total restaurants is not equal to large percentage of fastfood restaurants in that neighborhood, we can conclude that the distribution of fastfood restaurants is not even across neighborhoods, which also implies the motivation of our study, we want to investigate if this uneven distribution of fastfood restaurants is associated with different level of chronic disease outcomes within a neighborhood.

Restaurant Chains

We first scrape the list of 75 national chain restaurants in the US from the wikipedia page (https://en.wikipedia.org/wiki/List_of_restaurant_chains_in_the_United_States#Fast-casual) and then join this dataset with restaurant inspection data to choose only the chain restaurants in NYC.

```
chains_html = read_html("https://en.wikipedia.org/wiki/List_of_restaurant_chains_in_the_United_States#F")
```

```
# read in the list of chain restaurants in us
# made the names to uppercase and changed the var name to dba
chain_rest = chains_html %>%
  html_nodes("td:nth-child(1)") %>%
  html_text() %>%
  as.tibble() %>%
  mutate(dba = value,
         dba = str_to_upper(dba),
         dba = str_replace_all(dba, "[\r\n]" , "")) %>%
  select(dba)
```

```
## Warning: `as.tibble()` is deprecated, use `as_tibble()` (but mind the new semantics).
## This warning is displayed once per session.
```

```
head(chain_rest, 10)
```

```
## # A tibble: 10 x 1
##   dba
##   <chr>
## 1 ""
## 2 A&W RESTAURANTS
## 3 AMERICA'S INCREDIBLE PIZZA COMPANY
## 4 APPLEBEE'S
## 5 ARBY'S
## 6 ARCTIC CIRCLE RESTAURANTS
## 7 ARTHUR TREACHER'S
## 8 ATLANTA BREAD COMPANY
## 9 AUNTIE ANNE'S
## 10 BAHAMA BREEZE
```


Then, we match the list of chain restaurants in U.S. with the restaurant inspection data.

```
# removing punctuations in chain_rest & restaurant inspections (neighborhoods)
chain_rest_str =
  chain_rest %>%
  mutate(dba = str_replace_all(dba, "[[:punct:]]", ""))

rest_neigh_str = rest_neighborhood %>%
  mutate(dba = str_replace_all(dba, "[[:punct:]]", ""))

# Matching the two datasets(restaurant inspection data that has all punctuation removed from dba(restaurant inspection data))
neighborhood_chain =
  right_join(rest_neigh_str, chain_rest_str) %>%
  filter(!is.na(camis)) %>%
  distinct(camis, dba, neighborhood, boro)
```

```
## Joining, by = "dba"
```

```
neighborhood_chain %>%
  group_by(dba) %>%
  summarise(n = n()) %>%
  arrange(desc(n)) %>%
  head(10)
```

```
## # A tibble: 10 x 2
##   dba                n
##   <chr>             <int>
## 1 STARBUCKS         286
## 2 SUBWAY            286
## 3 MCDONALDS         205
## 4 POPEYES           101
## 5 BURGER KING        78
## 6 CHIPOTLE MEXICAN GRILL 76
## 7 DUNKIN DONUTS      50
## 8 WENDYS             43
## 9 KFC               36
## 10 LITTLE CAESARS    35
```

The combined dataset “neighborhood_chain” has 1546 observations. Also, there were 64 different chain restaurants extracted.

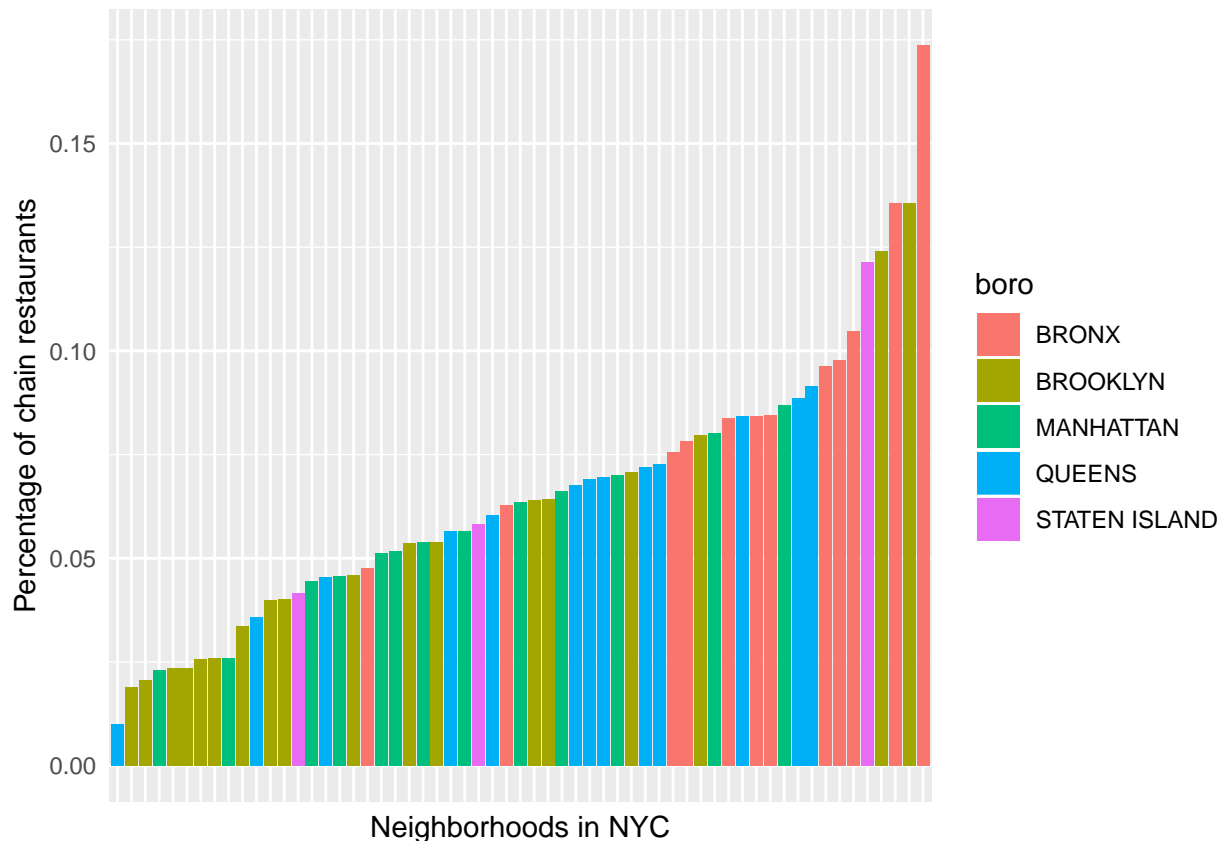
```
# counting chains in neighborhoods
neigh_count_chain = neighborhood_chain %>%
  group_by(neighborhood, boro) %>%
  summarise(chain_n = n())

neigh_count_rest = rest_neighborhood %>%
  distinct(neighborhood, camis) %>%
  group_by(neighborhood) %>%
  summarise(res_n = n())

# calculating percentage of chains in each neighborhood
percent_neighborhood_chain = left_join(neigh_count_chain, neigh_count_rest) %>%
  ungroup() %>%
  mutate(chain_percentage = chain_n/res_n,
         neighborhood = str_to_upper(neighborhood))
```

```
## Joining, by = "neighborhood"
plot_chain_neighbor = percent_neighborhood_chain %>%
  mutate(neighborhood = forcats::fct_reorder(neighborhood, chain_percentage)) %>%
  ggplot(aes(neighborhood, chain_percentage, fill = boro)) + geom_bar(stat="identity") +
  labs(x = "Neighborhoods in NYC", y = "Percentage of chain restaurants") +
  theme(axis.text.x = element_blank(), axis.ticks = element_blank())

#ggplotly(plot_chain_neighbor)
plot_chain_neighbor
```



```
max(percent_neighborhood_chain$chain_percentage)
```

```
## [1] 0.1736111
```

We plot neighborhoods in NYC with their percentage of chain restaurants and group them by borough. We can see that neighborhoods with those smallest percentages of chain restaurants are mostly in Brooklyn except for Greenwich Village and Soho and Lower East Side and Chinatown in Manhattan. Neighborhoods in Queens and Manhattan are spread out across low to high percentage of chain restaurants while most of the neighborhoods in Bronx and Staten Island have high percentages. The neighborhood with the highest percentage of chain restaurants is Throgs neck and Co-op City in Bronx with 17.4% of chain restaurants out of all restaurants.

Inspection Grade

```
gradea_neighborhood =
  rest_neighborhood %>%
```

```

group_by(neighborhood, grade) %>%
  summarise(n = n()) %>%
  mutate(grade_percent = n / sum(n)) %>%
  filter(grade == "A") %>%
  ungroup(boro) %>%
  mutate(neighborhood = str_to_upper(neighborhood))

# gradea_neighborhood %>%
#   mutate(neighborhood = as.factor(neighborhood),
#          neighborhood = fct_reorder(neighborhood, grade_percent)) %>%
#   plot_ly(x = ~neighborhood, y = ~grade_percent, color = ~neighborhood, type = "bar") %>%
#   layout(yaxis = list(title = 'Percentage of Grade A restaurants'),
#          xaxis = list(title = 'Neighborhoods in NYC', showticklabels = FALSE),
#          showlegend = FALSE)

```

The differences on percentage of “grade-A” restaurants between each neighborhood are observed. “Throgs Neck and Co-op City” has the greatest grade-A restaurant percentage, around 50.8%. “Sunset Park”, however, has the least, around 32.5%. “Washington Heights”, where we live, takes the fourth counting from the end, around 34%, which is obviously consistent with the feeling we have towards the restaurant condition of “Washington Heights”

Formal Analysis and Findings

Model Selection Process

First, we match the datasets containing all the restaurant information with the health datasets by common variable “neighborhood”.

```

health_neighborhood =
  health %>%
  mutate(neighborhood = str_to_upper(name)) %>%
  select(-name)

combined_chain =
  percent_neighborhood_chain %>%
  select(neighborhood, chain_percentage)
combined_chain_fastfood =
  fastfood_neighborhood %>%
  mutate(fastfood_percent = percent_fastfood) %>%
  select(neighborhood, fastfood_percent) %>%
  right_join(combined_chain, by = "neighborhood")
combined_chain_fastfood_gradea =
  gradea_neighborhood %>%
  select(neighborhood, grade_percent) %>%
  right_join(combined_chain_fastfood, by = "neighborhood")

combined_model =
  left_join(combined_chain_fastfood_gradea, health_neighborhood, by = "neighborhood")

```

Main predictor selection

```
outcome_name = combined_model[,10:12]

main_predictor_selection = function(outcome){

  lm_fastfood =
    lm(outcome ~ fastfood_percent + grade_percent + racewhite_rate + poverty + smoking + exercise + airqu
    summary()
  lm_fastfood_tbl =
    as.tibble(lm_fastfood[[4]]) %>%
    clean_names()

  lm_chain =
    lm(outcome ~ chain_percentage + grade_percent + racewhite_rate + poverty + smoking + exercise + airqu
    summary()
  lm_chain_tbl =
    as.tibble(lm_chain[[4]]) %>%
    clean_names()

  lm_both =
    lm(outcome ~ fastfood_percent + chain_percentage + grade_percent + racewhite_rate + poverty + smoking
    summary()
  lm_both_tbl =
    as.tibble(lm_both[[4]]) %>%
    clean_names()

  tibble(p_fastfood_sing = lm_fastfood_tbl$pr_t[2],
         p_chain_sing = lm_chain_tbl$pr_t[2],
         p_fastfood_both = lm_both_tbl$pr_t[2],
         p_chain_both = lm_both_tbl$pr_t[3])
}

main_predictor_comp =
  map(outcome_name, main_predictor_selection) %>%
  bind_rows()
main_predictor_comp$outcome = colnames(combined_model)[10:12]
main_predictor_comp

## # A tibble: 3 x 5
##   p_fastfood_sing p_chain_sing p_fastfood_both p_chain_both outcome
##           <dbl>         <dbl>         <dbl>         <dbl> <chr>
## 1      0.000287      0.00206      0.0368      0.380 obesity
## 2      0.0309       0.0592      0.241      0.587 diabetes
## 3      0.0140       0.340       0.0128      0.281 stroke_hosp
```

We originally have two main predictors of interest, *Percentage of chain restaurants* and *Percentage of fastfood restaurants*, and they are both significantly associated with the three chronic disease health outcomes when solely in the model after adjusting for other potential confounders. Taking obesity as example, the p-value for fastfood_percent is 0.00028698 and for chain_percentage is 0.0020629.

And we also anticipated them to be highly associated with each other, to avoid collinearity, we need to make decision on which one to keep as the final main predictor. So we put the two main predictor candidates in the same model and see how their p-values change. As a result, the fastfood_percent stays significant (p=0.0368396) and the chain_percentage turn insignificant (p=0.3800544).

Same selecting processes are repeated for the outcomes diabetes and stroke. Thus, we decide to have *Percentage of fastfood restaurants* (fastfood_percent) as the main predictor.

Confounder selection

```
outcome_name = combined_model[,10:12]

confounder_percent_change = function(outcome){
  lm_adjusted =
    lm(outcome ~ fastfood_percent + grade_percent + racewhite_rate + poverty + smoking + exercise + airquality_rate, combined_model)
  summary()
  lm_adjusted_tbl =
    as.tibble(lm_adjusted[[4]]) %>%
    clean_names()

  lm_crude =
    lm(outcome ~ fastfood_percent + racewhite_rate + poverty + smoking + exercise + airquality_rate, combined_model)
  summary()
  lm_crude_tbl =
    as.tibble(lm_crude[[4]]) %>%
    clean_names()

  percent_change = (lm_crude_tbl$estimate[2] - lm_adjusted_tbl$estimate[2]) / lm_crude_tbl$estimate[2]

  confounder_percent_change = percent_change
}

confounder_change =
  map(outcome_name, confounder_percent_change) %>%
  bind_rows()

percent <- function(x) {
  paste0(formatC(100 * x, digits = 3), "%")
}

confounder_change %>%
  kable()
```

obesity	diabetes	stroke_hosp
-0.0956578	0.2006834	-0.1072691

Besides some biologically meaningful covariates (i.e. race, poverty, smoking status, exercise), we also hypothesize the variable *Percentage of grade A restaurants* as a potential confounder in the association between fastfood restaurants percentage and the three chronic disease health outcomes.

Here, we are assessing if grade_percent is a significant confounder. Using the 10% change rule of thumb, we find that after adjusting for *Percentage of grade A restaurants*, the estimates of fastfood_percent change by -9.57% for outcome obesity, 20.1% for outcome diabetes and -10.7% for outcome stroke hospitalization.

Thus, regarding the final models, we are going to keep grade_percent for obesity and stroke hospitalization, but take it out and rerun the model for the outcome diabetes.

Final Models and Findings

Model 1: Obesity = fastfood_percent + grade_percent + racewhite_rate + poverty + smoking + exercise

```
lm_obesity =  
  lm(obesity ~ fastfood_percent + grade_percent + racewhite_rate + poverty + smoking + exercise + airquality, data = combined_model)  
  broom::tidy()  
lm_obesity %>%  
  kable()
```

term	estimate	std.error	statistic	p.value
(Intercept)	22.2928462	12.9549977	1.7207912	0.0913511
fastfood_percent	51.2148794	13.1494612	3.8948272	0.0002870
grade_percent	-12.7585912	18.7997554	-0.6786573	0.5004232
racewhite_rate	-0.0540849	0.0376614	-1.4360842	0.1570842
poverty	0.2255034	0.0954631	2.3622043	0.0220210
smoking	0.6282771	0.2161260	2.9069947	0.0053890
exercise	0.0998815	0.1564321	0.6384978	0.5260067
airquality_rate	-2.5244079	0.6363436	-3.9670515	0.0002280

Model 2: Diabetes = fastfood_percent + racewhite_rate + poverty + smoking + exercise

```
lm_diabetes =  
  lm(diabetes ~ fastfood_percent + racewhite_rate + poverty + smoking + exercise, combined_model) %>%  
  broom::tidy()  
lm_diabetes %>%  
  kable()
```

term	estimate	std.error	statistic	p.value
(Intercept)	18.6130268	5.2271424	3.560842	0.0007905
fastfood_percent	18.2960584	5.3558329	3.416100	0.0012267
racewhite_rate	-0.0744287	0.0166963	-4.457806	0.0000433
poverty	0.0407132	0.0376005	1.082785	0.2838053
smoking	0.1499565	0.1006247	1.490255	0.1420848
exercise	-0.1572366	0.0585560	-2.685236	0.0096543

Model 3: Stroke_hosp = fastfood_percent + grade_percent + racewhite_rate + poverty + smoking + exercise

```
lm_stroke =  
  lm(stroke_hosp ~ fastfood_percent + grade_percent + racewhite_rate + poverty + smoking + exercise, combined_model)  
  broom::tidy()  
lm_stroke %>%  
  kable()
```

term	estimate	std.error	statistic	p.value
(Intercept)	85.453409	157.0914792	0.5439723	0.5887854
fastfood_percent	441.513962	160.4689356	2.7513983	0.0081485
grade_percent	-127.390767	231.4310142	-0.5504481	0.5843691
racewhite_rate	-1.518999	0.4514699	-3.3645637	0.0014464
poverty	1.535131	1.0215604	1.5027317	0.1389574

term	estimate	std.error	statistic	p.value
smoking	6.823104	2.6286944	2.5956246	0.0122440
exercise	1.397276	1.5188594	0.9199509	0.3618465

We have three final models each having a health outcome (prevalence of obesity, prevalence of diabetes, and stroke hospitalization rate) as the dependent variable and percentage of fastfood as the main predictor. Obesity and percentage of fastfood restaurants model gives the most significant results with p-value of the main predictor being 0.00028698 (<0.01). In other words, at significance level of 1%, every 10% increase in the number of fastfood restaurants (0.1 unit increase in the percentage of fastfood restaurants) in a neighborhood is associated with 5.12% increase in the neighborhood's obesity prevalence, while adjusting for other factors.

The p-value of the regression coefficients for model with outcomes as diabetes/stroke are both less than 0.01 (Diabetes $p=0.0012267$; Stroke $p=0.0081485$), indicating there is a significant linear relationship between diabetes/stroke and percentage of fastfood restaurants at 1% significance level. To put in other words, at significance level of 1%, every 10% increase in the number of fastfood restaurants in a neighborhood is associated with 1.83% increase in the neighborhood's diabetes prevalence, while adjusting for other factors. Furthermore, every 10% increase in the number of fastfood restaurants in a neighborhood is associated with increase in 44.2 stroke hospitalizations per 100,000 adults in the neighborhood, while adjusting for other factors.

It is reasonable that among the three health outcomes, obesity has the strongest linear association with food environment, in this case, percentage of fastfood restaurants. It is a health condition that has the most direct relation with one's diet pattern. Moreover, it has a higher prevalence than diabetes and stroke, which could lead to a lower p-value than the other two health conditions.

Conclusion

Overall, we conclude that there is a significant relationship between chronic disease outcomes (i.e. obesity, diabetes, stroke) and the geographical distribution of fast-food restaurants in New York City.

Discussion

Our 2015 Community Health Profiles Open Data has data on obesity, diabetes prevalence rate from 2013 and stroke hospitalization rate from 2012. And the data of geographic distribution of restaurants is from 2014 to 2017. So the health profile data is somewhat earlier than the restaurant geographic distribution. However, as restaurant distribution and chronic disease prevalence rate wouldn't change much over a few years, the lag between years is not a large concern and theoretically won't affect our results that much. Therefore, our results can still be valid.

Although the cuisine type of a restaurant is typically assumed to directly reflect the health level of the food that restaurant provides, we demonstrate here that even the unhealthiest restaurant can offer one or several healthy foods, which may bring bias to our research. Besides, our study at first treats chain restaurants as a criteria to identify unhealthy restaurants. Obviously this criteria is ambiguous and farfetched. In the end, the study doesn't include this as the predictor.

Lastly, although there is an association between fast-food restaurant geographical distribution and chronic disease outcomes, we could not conclude that it is fast-food restaurant causing these health outcomes. Maybe it is because the citizens living in the neighborhood are obese indicating that they like to eat fast food. This will actually result in more fast-food restaurant opening in this neighborhood since the business will be good. Further study need to be conducted before concluding any causation effect.