

We load previously loaded census data. The code for fetching this data is also shown in this section.

Table 1: Census Data, 2021 (truncated rows and columns)

1

## Food data

```
food_data <- st_read(here("data/free-and-low-cost-food-programs.shp")) %>%
  select(
    "program_nam",
    "program_sta",
    "meal_cost",
    "local_areas",
    "latitude",
    "longitude",
    "geometry"
  ) %>%
  drop_na("latitude", "longitude") %>%
  # set to wgs 84 as per can census
  st_set_crs(4326)

## Reading layer `free-and-low-cost-food-programs' from data source
##   `/Users/sid/Documents/ubc classes/2024w1/econ 326/foodprograms/data/free-and-low-
## cost-food-programs.shp'
##   using driver `ESRI Shapefile'
## replacing null geometries with empty geometries
## Simple feature collection with 83 features and 25 fields (with 2 geometries empty)
## Geometry type: POINT
## Dimension:      XY
## Bounding box:   xmin: -123.1821 ymin: 49.20725 xmax: -123.0287 ymax: 49.286
## CRS:            NA

# Food data processing
food_count <- food_data %>%
  st_set_geometry(NULL) %>%
  group_by(local_areas) %>%
  summarise(count = n(), .groups = "drop")

food_data_count <- food_data %>%
  left_join(food_count, by = "local_areas") %>%
  distinct(local_areas, .keep_all = TRUE) # one row per neighbourhood

combo_food_census <- census_data %>%
  st_join(food_data_count)

census_data_food <- combo_food_census %>%
  mutate(program_count = replace_na(count, 0),
         food_density = program_count / `Shape Area`)

head(census_data_food[, 1:5]) %>%
  kable(format = "latex", booktabs = TRUE, caption = "Food Data merged with Census Data, 2021 (truncated)",
        kable_styling(latex_options = c("striped", "hold_position"))
```

---

## Crime data

```
crime <- read_csv(here("data/crime_data_all_neighborhoods.csv"), show_col_types = FALSE) %>%
  mutate(TYPE = as_factor(TYPE),
         HUNDRED_BLOCK = as_factor(HUNDRED_BLOCK),
```

Table 2: Food Data merged with Census Data, 2021 (truncated rows and columns)

Shape Area	Type	Households	Quality Flags	name	geometry
0.2991	DA	266	0	59150307	MULTIPOLYGON (((-123.0231 4...
0.1096	DA	218	0	59150308	MULTIPOLYGON (((-123.0234 4...
0.1119	DA	282	0	59150309	MULTIPOLYGON (((-123.0283 4...
0.1094	DA	389	0	59150310	MULTIPOLYGON (((-123.0234 4...
0.0809	DA	187	0	59150311	MULTIPOLYGON (((-123.0257 4...
0.0871	DA	201	0	59150312	MULTIPOLYGON (((-123.0234 4...

```

    NEIGHBOURHOOD = as_factor(NEIGHBOURHOOD)) %>%
  filter(!is.na(X) & !is.na(Y))

crime_data <- st_as_sf(crime, coords = c("X", "Y"), crs = "+proj=utm +zone=10") %>%
  st_transform(crs = "+proj=longlat +datum=WGS84")

intersections <- st_is_within_distance(census_data, crime_data, sparse = FALSE, dist = 5)

crimes_contained <- rowSums(intersections, dims = 1)

census_data_crime <- census_data %>%
  cbind(crimes_contained) %>%
  mutate(crime_density = crimes_contained / Shape.Area)

unique_crimes <- unique(crime_data$TYPE)

for (type in unique_crimes) {
  type_data <- crime_data %>% filter(TYPE == type)
  intersections <- st_is_within_distance(census_data, type_data, sparse = FALSE, dist = 5)
  sum <- rowSums(intersections, dims = 1)
  df <- as.data.frame(sum)
  census_data_crime <- census_data_crime %>% cbind(df$sum) %>% rename_with(~ paste0("crimes_", type), d
}

```

## Training data

Now we merge all of our data-sets into one table so we can feed it into our model.

```

training_data <- st_join(census_data_crime, census_data_food)

training_data <- training_data %>%
  rename_with(
    ~ gsub(".*$", "", .), # Remove everything after the colon, including the colon
    starts_with("v_CA21") # Apply only to columns starting with "v_CA21"
  )

median_lico_at <- median(training_data$v_CA21_1085, na.rm = TRUE)

training_data <- training_data %>%
  mutate(
    low_income = ifelse(
      v_CA21_1085 > median_lico_at,

```

```
1,  
0))
```

```
training_data[is.na(training_data)] <- 0
```

## Model

### Specification

```
# Specification models  
# 1. Model with all variables (including interaction term)  
reg_all_vars <- lm(crime_density ~  
  food_density:low_income +  
  food_density +  
  pop_density.x +  
  v_CA21_452 +  
  low_income,  
  data = st_set_geometry(training_data, NULL))  
  
# 2. Model without the interaction term  
reg_no_interaction <- lm(crime_density ~  
  food_density +  
  pop_density.x +  
  v_CA21_452 +  
  low_income,  
  data = st_set_geometry(training_data, NULL))  
  
# 3. Model with only food density, crime density, and low income  
reg_food_crime_low_income <- lm(crime_density ~  
  food_density +  
  low_income,  
  data = st_set_geometry(training_data, NULL))  
  
# 4. Model with all variables (including interaction term) but replacing v_CA21_452 with v_CA21_449  
reg_all_vars_449 <- lm(crime_density ~  
  food_density:low_income +  
  food_density +  
  pop_density.x +  
  v_CA21_449 + # Replaced v_CA21_452 with v_CA21_449  
  low_income,  
  data = st_set_geometry(training_data, NULL))  
  
models_spec <- list(  
  "All Variables" = reg_all_vars,  
  "Without Interaction" = reg_no_interaction,  
  "Food Density, Crime, Low Income" = reg_food_crime_low_income  
)  
  
# Summary for each specification tested  
summary_all_vars <- summary(reg_all_vars)  
summary_no_interaction <- summary(reg_no_interaction)
```

```
summary_food_crime_low_income <- summary(reg_food_crime_low_income)
summary_all_vars_449 <- summary(reg_all_vars_449)

# Add to the list of model summaries
model_summaries <- list(
  "All Variables" = summary_all_vars,
  "Without Interaction" = summary_no_interaction,
  "Food Density, Crime, Low Income" = summary_food_crime_low_income,
  "All Variables (with v_CA21_449)" = summary_all_vars_449
)
```

## RESULTS FROM SPEC

### Function definitions

We needed to manipulate our regression model data into formatted tables that emulated those from the package `stargazer` as we were having significant alignment issues between the data and rendered tables.

```
all_variables <- unique(c(
  names(coef(reg_all_vars)),
  names(coef(reg_no_interaction)),
  names(coef(reg_food_crime_low_income))
))

all_variables_escaped <- gsub("_", "\\_", all_variables) # Escape underscores
all_variables_escaped <- paste0("\\texttt{" , all_variables_escaped, "}") # Wrap in \texttt{} for LaTeX

add_stars <- function(estimates, ses, p_values) {
  significance_levels <- ifelse(p_values < 0.01, "***",
                                ifelse(p_values < 0.05, "**",
                                ifelse(p_values < 0.1, "*", "")))
  formatted <- sprintf("%.3f (%.3f)%s", estimates, ses, significance_levels)
  return(formatted)
}

extract_summary <- function(model, all_vars) {
  coefs <- coef(model)
  ses <- sqrt(diag(vcov(model)))
  p_values <- coef(summary(model))[, 4] # Extract p-values from the summary

  # Create placeholders for all variables
  aligned_coefs <- setNames(rep(NA, length(all_vars)), all_vars)
  aligned_ses <- setNames(rep(NA, length(all_vars)), all_vars)
  aligned_p_values <- setNames(rep(NA, length(all_vars)), all_vars)

  # Fill with existing coefficients, standard errors, and p-values
  for (var in names(coefs)) {
    aligned_coefs[var] <- coefs[var]
    aligned_ses[var] <- ses[var]
    aligned_p_values[var] <- p_values[var]
  }

  # Replace missing coefficients with 0
  aligned_coefs[is.na(aligned_coefs)] <- 0
}
```

```

# Add stars to coefficients based on significance levels
significance_levels <- ifelse(aligned_p_values < 0.01, "***",
                             ifelse(aligned_p_values < 0.05, "**",
                                     ifelse(aligned_p_values < 0.1, "*", "")))

# Format coefficients and standard errors for display
formatted <- ifelse(is.na(aligned_ses),
                   sprintf("%.3f", aligned_coefs),
                   sprintf("%.3f (%.3f)%s", aligned_coefs, aligned_ses, significance_levels))

# Replace missing standard errors with blank strings
formatted[is.na(aligned_ses)] <- " "

# Escape underscores for LaTeX compatibility and wrap in \texttt{} for table display
formatted <- gsub("_", "\\_", formatted)
formatted <- paste0("\\texttt{", formatted, "}")

return(formatted)
}

extract_model_info <- function(model, covariate_labels, model_title) {
  coefs <- coef(model)
  ses <- sqrt(diag(vcov(model)))
  p_values <- coef(summary(model))[, 4] # Extract p-values from the summary

  # Format coefficients, standard errors, and significance stars
  significance_levels <- ifelse(p_values < 0.01, "***",
                               ifelse(p_values < 0.05, "**",
                                       ifelse(p_values < 0.1, "*", "")))

  formatted <- sprintf("%.3f (%.3f)%s", coefs, ses, significance_levels)

  # Combine covariate labels and the formatted coefficients
  result <- data.frame(
    Variable = covariate_labels,
    Estimate = formatted,
    stringsAsFactors = FALSE
  )

  # Add model title
  result <- rbind(data.frame(Variable = model_title, Estimate = "", stringsAsFactors = FALSE), result)

  return(result)
}

# Extract R-squared, F-statistics, and number of observations from each model
r_squared <- c(
  summary(reg_all_vars)$r.squared,
  summary(reg_no_interaction)$r.squared,
  summary(reg_food_crime_low_income)$r.squared
)

```

```
f_statistic <- c(
  summary(reg_all_vars)$fstatistic[1],
  summary(reg_no_interaction)$fstatistic[1],
  summary(reg_food_crime_low_income)$fstatistic[1]
)

n_obs <- c(
  length(reg_all_vars$fitted.values),
  length(reg_no_interaction$fitted.values),
  length(reg_food_crime_low_income$fitted.values)
)

model1_aligned <- extract_summary(reg_all_vars, all_variables)
model2_aligned <- extract_summary(reg_no_interaction, all_variables)
model3_aligned <- extract_summary(reg_food_crime_low_income, all_variables)
```

## Data manipulation

```
aligned_table <- data.frame(
  Variable = all_variables_escaped,
  `Model 1` = model1_aligned,
  `Model 2` = model2_aligned,
  `Model 3` = model3_aligned,
  stringsAsFactors = FALSE
)

print(aligned_table)
```

	Variable	Model.1
## 1	\\texttt{(Intercept)}	\\texttt{1291.286 (62.011)***}
## 2	\\texttt{food\\_density}	\\texttt{-0.203 (2.154)}
## 3	\\texttt{pop\\_density.x}	\\texttt{0.043 (0.001)***}
## 4	\\texttt{v\\_CA21\\_452}	\\texttt{-517.756 (22.439)***}
## 5	\\texttt{low\\_income}	\\texttt{244.697 (24.734)***}
## 6	\\texttt{food\\_density:low\\_income}	\\texttt{9.460 (2.305)***}
##	Model.2	Model.3
## 1	\\texttt{1283.891 (62.047)***}	\\texttt{383.969 (18.802)***}
## 2	\\texttt{8.053 (0.771)***}	\\texttt{9.631 (0.897)***}
## 3	\\texttt{0.044 (0.001)***}	\\texttt{ }
## 4	\\texttt{-517.272 (22.461)***}	\\texttt{ }
## 5	\\texttt{252.689 (24.682)***}	\\texttt{577.872 (27.744)***}
## 6	\\texttt{ }	\\texttt{ }

```
summary_table <- data.frame(
  Variable = c("R squared", "F statistic", "Observations"),
  `Model 1` = c(sprintf("%.3f", r_squared[1]), sprintf("%.1f", f_statistic[1]), n_obs[1]),
  `Model 2` = c(sprintf("%.3f", r_squared[2]), sprintf("%.1f", f_statistic[2]), n_obs[2]),
  `Model 3` = c(sprintf("%.3f", r_squared[3]), sprintf("%.1f", f_statistic[3]), n_obs[3]),
  stringsAsFactors = FALSE
)

# Combine both tables
```

```
final_table <- rbind(aligned_table, summary_table)
print(final_table)
```

```
##                               Variable                               Model.1
## 1      \\texttt{(Intercept)} \\texttt{1291.286 (62.011)***}
## 2      \\texttt{food\\_density}      \\texttt{-0.203 (2.154)}
## 3      \\texttt{pop\\_density.x}      \\texttt{0.043 (0.001)***}
## 4      \\texttt{v\\_CA21\\_452} \\texttt{-517.756 (22.439)***}
## 5      \\texttt{low\\_income} \\texttt{244.697 (24.734)***}
## 6 \\texttt{food\\_density:low\\_income} \\texttt{9.460 (2.305)***}
## 7                               R squared                               0.315
## 8                               F statistic                               721.0
## 9                               Observations                               7860
##                               Model.2                               Model.3
## 1 \\texttt{1283.891 (62.047)***} \\texttt{383.969 (18.802)***}
## 2      \\texttt{8.053 (0.771)***}      \\texttt{9.631 (0.897)***}
## 3      \\texttt{0.044 (0.001)***}      \\texttt{ }
## 4 \\texttt{-517.272 (22.461)***}      \\texttt{ }
## 5 \\texttt{252.689 (24.682)***} \\texttt{577.872 (27.744)***}
## 6      \\texttt{ }      \\texttt{ }
## 7      0.313      0.067
## 8      895.2      282.6
## 9      7860      7860
```

```
covariate_labels <- c(
  "(Intercept)",
  "Food Program Density: Low Income",
  "Food Program Density",
  "Population Density",
  "Average Household Size (v\\_CA21\\_452)",
  "Low Income"
)
```

```
covariate_labels_2 <- c(
  "(Intercept)",
  "Food Program Density",
  "Population Density",
  "Average Household Size (v\\_CA21\\_452)",
  "Low Income"
)
```

```
covariate_labels_3 <- c(
  "(Intercept)",
  "Food Program Density",
  "Low Income"
)
```

```
model_1_table <- extract_model_info(
  model = reg_all_vars,
  covariate_labels = covariate_labels,
  model_title = "Spec 1: All Variables"
)
```

```
model_2_table <- extract_model_info(
  model = reg_no_interaction,
```



```

    covariate_labels = covariate_labels_2,
    model_title = "Spec 2: Omitted Interaction Term"
)

model_3_table <- extract_model_info(
  model = reg_food_crime_low_income,
  covariate_labels = covariate_labels_3,
  model_title = "Spec 3: Food Density and Low Income"
)

```

## Regression table

```

# Render the final table in a stargazer-like format using knitr::kable
kable(
  final_table,
  format = "latex",
  col.names = c("Variable", "Model 1", "Model 2", "Model 3"),
  caption = "Regression Specifications",
  align = "lccc",
  booktabs = TRUE,
  escape = FALSE
) %>%
  kable_styling(latex_options = c("hold_position", "striped")) %>%
  add_header_above(c(" " = 1, "Dependent Variable: Crime Density" = 3)) %>%
  footnote(general = "* p < 0.1; ** p < 0.05; *** p < 0.01",
    general_title = "Note:",
    footnote_as_chunk = TRUE,
    escape = FALSE)

```

Table 3: Regression Specifications

Variable	Dependent Variable: Crime Density		
	Model 1	Model 2	Model 3
(Intercept)	1291.286 (62.011)***	1283.891 (62.047)***	383.969 (18.802)***
food_density	-0.203 (2.154)	8.053 (0.771)***	9.631 (0.897)***
pop_density.x	0.043 (0.001)***	0.044 (0.001)***	
v_CA21_452	-517.756 (22.439)***	-517.272 (22.461)***	
low_income	244.697 (24.734)***	252.689 (24.682)***	577.872 (27.744)***
food_density:low_income	9.460 (2.305)***		
R squared	0.315	0.313	0.067
F statistic	721.0	895.2	282.6
Observations	7860	7860	7860

Note: \* p < 0.1; \*\* p < 0.05; \*\*\* p < 0.01

```

stargazer(reg_all_vars, reg_no_interaction, reg_food_crime_low_income, type = "latex",
  covariate.labels = c("Food Program Density", "Population Density", "Household Size", "Low Income"),
  dep.var.labels = c("Crime Density")
)

```

% Table created by stargazer v.5.2.3 by Marek Hlavac, Social Policy Institute. E-mail: marek.hlavac at gmail.com % Date and time: Tue, Dec 03, 2024 - 13:36:29

Table 4:

	<i>Dependent variable:</i>		
	Crime Density		
	(1)	(2)	(3)
Food Program Density	−0.203 (2.154)	8.053*** (0.771)	9.631*** (0.897)
Population Density	0.043*** (0.001)	0.044*** (0.001)	
Household Size	−517.756*** (22.439)	−517.272*** (22.461)	
Low Income	244.697*** (24.734)	252.689*** (24.682)	577.872*** (27.744)
Food Density:Low Income	9.460*** (2.305)		
Constant	1,291.286*** (62.011)	1,283.891*** (62.047)	383.969*** (18.802)
Observations	7,860	7,860	7,860
R <sup>2</sup>	0.315	0.313	0.067
Adjusted R <sup>2</sup>	0.314	0.313	0.067
Residual Std. Error	1,050.245 (df = 7854)	1,051.304 (df = 7855)	1,225.029 (df = 7857)
F Statistic	720.960*** (df = 5; 7854)	895.186*** (df = 4; 7855)	282.621*** (df = 2; 7857)

*Note:*

\*p&lt;0.1; \*\*p&lt;0.05; \*\*\*p&lt;0.01

### Model 1:

```
# Model 1: reg_all_vars
stargazer(reg_all_vars, type = "latex",
  covariate.labels = c("Food Program Density", "Population Density", "Household Size", "Low Income"),
  dep.var.labels = c("Crime Density")
)
```

% Table created by stargazer v.5.2.3 by Marek Hlavac, Social Policy Institute. E-mail: marek.hlavac@gmail.com % Date and time: Tue, Dec 03, 2024 - 13:36:29

Table 5:	
	<i>Dependent variable:</i>
	Crime Density
Food Program Density	−0.203 (2.154)
Population Density	0.043*** (0.001)
Household Size	−517.756*** (22.439)
Low Income	244.697*** (24.734)
Food Density:Low Income	9.460*** (2.305)
Constant	1,291.286*** (62.011)
Observations	7,860
R <sup>2</sup>	0.315
Adjusted R <sup>2</sup>	0.314
Residual Std. Error	1,050.245 (df = 7854)
F Statistic	720.960*** (df = 5; 7854)
<i>Note:</i> *p<0.1; **p<0.05; ***p<0.01	

### Model 2:

```
stargazer(reg_no_interaction, type = "latex",
  covariate.labels = c("Food Program Density", "Population Density", "Household Size", "Low Income"),
  dep.var.labels = c("Crime Density")
)
```

% Table created by stargazer v.5.2.3 by Marek Hlavac, Social Policy Institute. E-mail: marek.hlavac@gmail.com % Date and time: Tue, Dec 03, 2024 - 13:36:29

Table 6:

	<i>Dependent variable:</i>
	Crime Density
Food Program Density	8.053*** (0.771)
Population Density	0.044*** (0.001)
Household Size	-517.272*** (22.461)
Low Income	252.689*** (24.682)
Constant	1,283.891*** (62.047)
Observations	7,860
R <sup>2</sup>	0.313
Adjusted R <sup>2</sup>	0.313
Residual Std. Error	1,051.304 (df = 7855)
F Statistic	895.186*** (df = 4; 7855)
<i>Note:</i>	*p<0.1; **p<0.05; ***p<0.01

### Model 3:

```
# Model 3: reg_food_crime_low_income
stargazer(reg_food_crime_low_income, type = "latex",
  covariate.labels = c("Food Program Density", "Low Income"),
  dep.var.labels = c("Crime Density")
)
```

% Table created by stargazer v.5.2.3 by Marek Hlavac, Social Policy Institute. E-mail: marek.hlavac at gmail.com % Date and time: Tue, Dec 03, 2024 - 13:36:29

Table 7:

	<i>Dependent variable:</i>
	Crime Density
Food Program Density	9.631*** (0.897)
Low Income	577.872*** (27.744)
Constant	383.969*** (18.802)
Observations	7,860
R <sup>2</sup>	0.067
Adjusted R <sup>2</sup>	0.067
Residual Std. Error	1,225.029 (df = 7857)
F Statistic	282.621*** (df = 2; 7857)
Note:	*p<0.1; **p<0.05; ***p<0.01

### Robustness

We split the data into sub-groups based on

```
census_data_centroids <- census_data %>%
  mutate(
    centroid = st_centroid(geometry),
    latitude = st_coordinates(centroid)[, 2],
    longitude = st_coordinates(centroid)[, 1]
  ) %>%
  st_set_geometry(NULL) # Remove geometry for k-means input

# Normalize latitude and longitude for clustering
census_data_normalized <- census_data_centroids %>%
  mutate(
    latitude_scaled = scale(latitude),
    longitude_scaled = scale(longitude)
  )

set.seed(123) # For reproducibility
```

```

k <- 4

kmeans_result <- kmeans(census_data_normalized[, c("latitude_scaled", "longitude_scaled")], centers = k)

# Add cluster assignments to the original data
census_data$cluster <- kmeans_result$cluster

# Now, merge the cluster assignments into your training data
training_data_clusters <- st_join(training_data,
                                   census_data %>% select(cluster, geometry),
                                   left = TRUE)

# Run the model for each cluster
cluster_models <- list()

for (i in 1:k) {
  subset_data <- training_data_clusters %>% filter(cluster == i)

  model <- lm(crime_density ~
              food_density:low_income +
              food_density +
              pop_density.x +
              v_CA21_452 +
              low_income,
              data = st_set_geometry(subset_data, NULL))

  cat(paste("\nModel Summary for Cluster", i, "\n"))
  cluster_models[[i]] <- model
  print(summary(model)) # Check the results for each cluster
}

```

```

##
## Model Summary for Cluster 1 :
##
## Call:
## lm(formula = crime_density ~ food_density:low_income + food_density +
##     pop_density.x + v_CA21_452 + low_income, data = st_set_geometry(subset_data,
##     NULL))
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1216.1  -147.9   -59.4    51.4   4377.5
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    8.200e+02  1.759e+01  46.611  <2e-16 ***
## food_density    6.652e-01  2.692e-01   2.471  0.0135 *
## pop_density.x    2.683e-02  5.950e-04  45.102  <2e-16 ***
## v_CA21_452     -2.569e+02  6.370e+00 -40.326  <2e-16 ***
## low_income     -1.282e+01  6.522e+00  -1.965  0.0494 *
## food_density:low_income -2.648e-01  3.629e-01  -0.730  0.4657
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##

```

```

## Residual standard error: 321.8 on 15892 degrees of freedom
## Multiple R-squared:  0.2516, Adjusted R-squared:  0.2514
## F-statistic: 1068 on 5 and 15892 DF,  p-value: < 2.2e-16
##
##
## Model Summary for Cluster 2 :
##
## Call:
## lm(formula = crime_density ~ food_density:low_income + food_density +
##     pop_density.x + v_CA21_452 + low_income, data = st_set_geometry(subset_data,
##     NULL))
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -4251.5 -1104.8  -406.5   361.3 10754.9
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    3.079e+03  9.577e+01  32.156 < 2e-16 ***
## food_density   -7.841e+00  5.525e+00  -1.419  0.15587
## pop_density.x    2.622e-02  1.052e-03  24.929 < 2e-16 ***
## v_CA21_452     -1.397e+03  5.016e+01 -27.842 < 2e-16 ***
## low_income      9.276e+02  3.169e+01  29.269 < 2e-16 ***
## food_density:low_income 1.671e+01  5.552e+00   3.010  0.00262 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1847 on 16445 degrees of freedom
## Multiple R-squared:  0.1932, Adjusted R-squared:  0.1929
## F-statistic: 787.4 on 5 and 16445 DF,  p-value: < 2.2e-16
##
##
## Model Summary for Cluster 3 :
##
## Call:
## lm(formula = crime_density ~ food_density:low_income + food_density +
##     pop_density.x + v_CA21_452 + low_income, data = st_set_geometry(subset_data,
##     NULL))
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -272.32  -71.09  -22.73   34.85 1205.21
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    2.369e+02  7.746e+00  30.583 < 2e-16 ***
## food_density    9.935e-01  2.953e-01   3.364  0.00077 ***
## pop_density.x    1.591e-02  1.574e-04 101.094 < 2e-16 ***
## v_CA21_452     -5.109e+01  2.627e+00 -19.451 < 2e-16 ***
## low_income      5.528e+00  2.231e+00   2.478  0.01322 *
## food_density:low_income -2.823e+00  4.270e-01  -6.610 3.94e-11 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##

```

```

## Residual standard error: 133.4 on 18348 degrees of freedom
## Multiple R-squared:  0.385, Adjusted R-squared:  0.3849
## F-statistic: 2298 on 5 and 18348 DF, p-value: < 2.2e-16
##
##
## Model Summary for Cluster 4 :
##
## Call:
## lm(formula = crime_density ~ food_density:low_income + food_density +
##     pop_density.x + v_CA21_452 + low_income, data = st_set_geometry(subset_data,
##     NULL))
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -415.34  -62.84  -24.39   41.99  783.31
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      6.664e+01  4.931e+00  13.513 < 2e-16 ***
## food_density      5.133e+00  4.278e-01  11.997 < 2e-16 ***
## pop_density.x      1.237e-02  2.701e-04  45.811 < 2e-16 ***
## v_CA21_452      -1.375e+01  1.908e+00  -7.206 6.13e-13 ***
## low_income        3.670e+01  2.923e+00  12.556 < 2e-16 ***
## food_density:low_income -5.226e-01  6.081e-01  -0.859    0.39
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 127.7 on 11341 degrees of freedom
## Multiple R-squared:  0.2003, Adjusted R-squared:  0.1999
## F-statistic: 568 on 5 and 11341 DF, p-value: < 2.2e-16
stargazer(cluster_models, type = "latex",
           covariate.labels = c("Food Program Density", "Population Density", "Household Size", "Low Income"),
           dep.var.labels = c("Crime Density"))

% Table created by stargazer v.5.2.3 by Marek Hlavac, Social Policy Institute. E-mail: marek.hlavac at
gmail.com % Date and time: Tue, Dec 03, 2024 - 13:36:30

# Convert the cluster column to a factor
census_data$cluster <- as.factor(census_data$cluster)

# Plot the regions and highlight clusters with adjusted color scale
ggplot(data = census_data) +
  geom_sf(aes(fill = cluster), color = "black", lwd = 0.1) +
  scale_fill_viridis_d(name = "Cluster ID") + # Automatically assign colors for clusters
  labs(title = "Census Regions Highlighted by Cluster (grouped by k-means clustering)",
       fill = "Cluster ID") +
  theme_minimal() +
  theme(legend.position = "right",
        plot.title = element_text(hjust = 0.5, size = 10, face = "bold"),
        plot.subtitle = element_text(hjust = 0.5, size = 12))

```



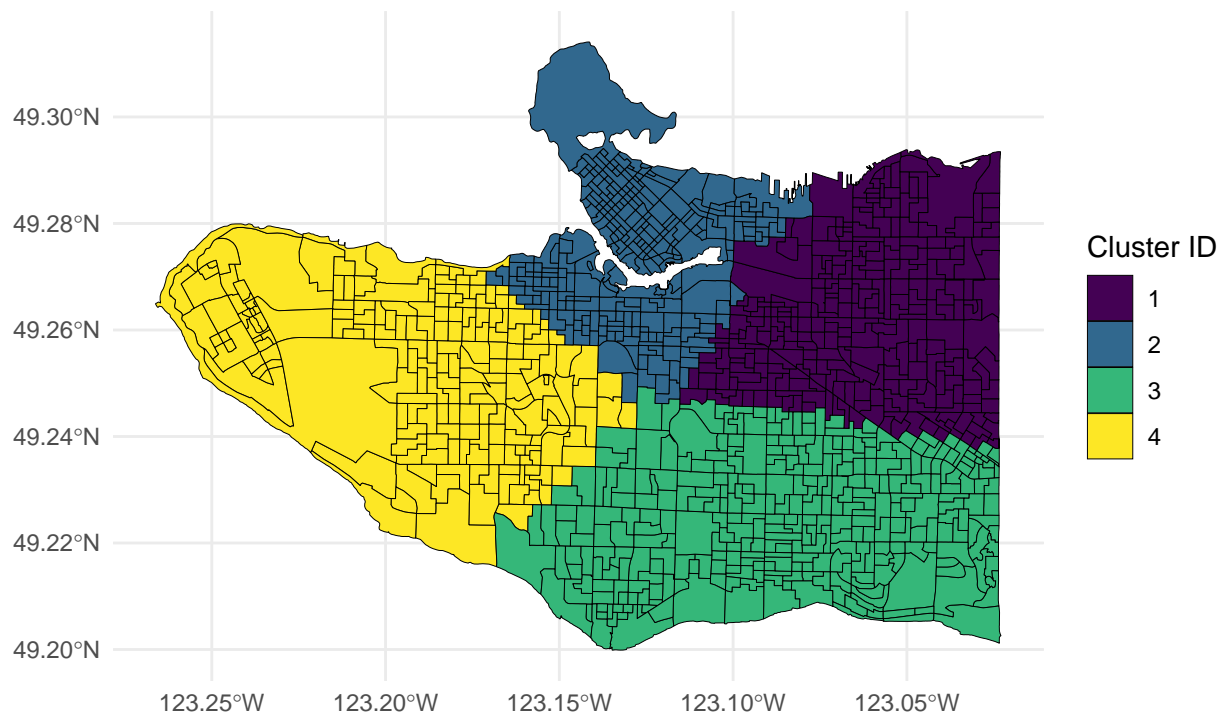
Table 8:

	<i>Dependent variable:</i>		
	Crime Density		
	(1)	(2)	(3)
Food Program Density	0.665** (0.269)	−7.841 (5.525)	0.994*** (0.295)
Population Density	0.027*** (0.001)	0.026*** (0.001)	0.016*** (0.0002)
Household Size	−256.874*** (6.370)	−1,396.640*** (50.163)	−51.092*** (2.627)
Low Income	−12.819** (6.522)	927.569*** (31.691)	5.528** (2.231)
Food Density:Low Income	−0.265 (0.363)	16.712*** (5.552)	−2.823*** (0.427)
Constant	819.960*** (17.592)	3,079.475*** (95.768)	236.881*** (7.746)
Observations	15,898	16,451	18,354
R <sup>2</sup>	0.252	0.193	0.385
Adjusted R <sup>2</sup>	0.251	0.193	0.385
Residual Std. Error	321.788 (df = 15892)	1,846.752 (df = 16445)	133.450 (df = 18348)
F Statistic	1,068.458*** (df = 5; 15892)	787.362*** (df = 5; 16445)	2,297.625*** (df = 5; 18348)

*Note:*

\*p

**Census Regions Highlighted by Cluster (grouped by k-means clustering)**



## Heteroskedascity

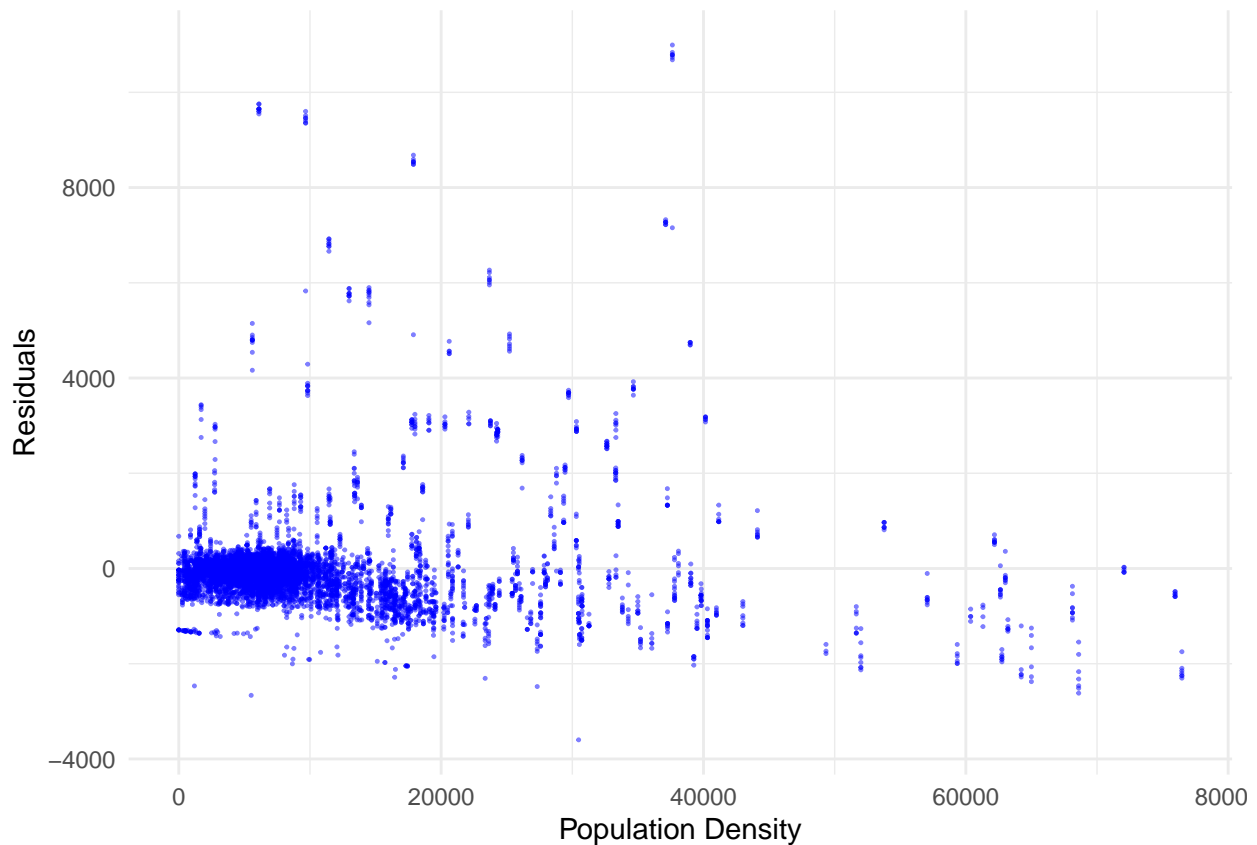
```
# Assuming you have a model named 'reg_all_vars' and a dataset named 'training_data'
```

```
# Add residuals to the dataset
```

```
training_data_res <- training_data %>%  
  mutate(residuals = resid(reg_all_vars))
```

```
# Plot residuals against a variable (e.g., population density) to visually check for heteroskedasticity
```

```
ggplot(data = training_data_res, aes(x = pop_density.x, y = residuals)) +  
  geom_point(size = 0.2, alpha = 0.5, color = "blue") +  
  labs(x = "Population Density", y = "Residuals") +  
  theme_minimal() +  
  theme(plot.title = element_text(hjust = 0.5))
```



```
# Extract residuals and calculate squared residuals
training_data_res <- training_data_res %>%
  mutate(residuals_squared = residuals^2)

# Auxiliary regression: squared residuals on independent variables, their squares, and interactions
white_model <- lm(residuals_squared ~
  food_density + pop_density.x + v_CA21_452 + low_income +
  I(food_density^2) + I(pop_density.x^2) + I(v_CA21_452^2) + I(low_income^2) +
  food_density:pop_density.x + food_density:v_CA21_452 + food_density:low_income +
  pop_density.x:v_CA21_452 + pop_density.x:low_income +
  v_CA21_452:low_income,
  data = training_data_res)

# Calculate the test statistic
n <- nrow(training_data)
R_squared <- summary(white_model)$r.squared
white_test_statistic <- n * R_squared

# Calculate p-value for the test statistic (chi-square with degrees of freedom equal to number of predi
p_value <- pchisq(white_test_statistic, df = length(coef(white_model)) - 1, lower.tail = FALSE)

cat("White's Test Statistic:", white_test_statistic, "\n")

## White's Test Statistic: 644.1092

cat("p-value:", p_value, "\n")

## p-value: 2.147077e-128
```

```

# Auxiliary regression for Breusch-Pagan Test: regress squared residuals on original predictors
bp_model <- lm(residuals_squared ~
               food_density + pop_density.x + v_CA21_452 + low_income,
               data = training_data_res, na.action = na.omit)

# Calculate the test statistic for Breusch-Pagan test
R_squared_bp <- summary(bp_model)$r.squared
bp_test_statistic <- n * R_squared_bp

# Calculate p-value for Breusch-Pagan test statistic (chi-square)
p_value_bp <- pchisq(bp_test_statistic, df = length(coef(bp_model)) - 1, lower.tail = FALSE)

cat("Breusch-Pagan Test Statistic:", bp_test_statistic, "\n")

## Breusch-Pagan Test Statistic: 436.9128

cat("Breusch-Pagan Test p-value:", p_value_bp, "\n")

## Breusch-Pagan Test p-value: 2.930512e-93

# Calculate robust standard errors and re-run the model with robust covariance matrix
robust_se <- coeftest(reg_all_vars, vcov = vcovHC(reg_all_vars, type = "HC"))

# Print the results with robust standard errors

stargazer(robust_se, type = "latex",
           covariate.labels = c("Food Program Density", "Population Density", "Household Size", "Low Income"),
           dep.var.labels = c("Crime Density")
           )

```

% Table created by stargazer v.5.2.3 by Marek Hlavac, Social Policy Institute. E-mail: marek.hlavac@gmail.com  
 % Date and time: Tue, Dec 03, 2024 - 13:36:31

Table 9:

	<i>Dependent variable:</i>
	Crime Density
Food Program Density	−0.203 (0.545)
Population Density	0.043*** (0.002)
Household Size	−517.756*** (28.164)
Low Income	244.697*** (21.468)
Food Density:Low Income	9.460*** (2.844)
Constant	1,291.286*** (79.646)
<hr/> <hr/>	
<i>Note:</i>	*p<0.1; **p<0.05; ***p<0.01