Exploring Patterns of Environmental Justice in Los Angeles County

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Primary Objective

With an estimated 9.8 million residents in 2023, Los Angeles County is the most populous county in California and the entire United States (Data USA, n.d.). Los Angeles County is also racially diverse, with approximately 2.48 million White (non-Hispanic) residents, 2.26 million Hispanic residents, 1.4 million Asian residents, and 743,000 Black residents (Data USA, n.d.). The city has a rich history and is culturally diverse; however, marginalized communities continue to face disproportionate environmental, social, and economic burdens.

Environmental injustices experienced by marginalized communities are interconnected with patterns of reduced biodiversity in urban environments. The Home Owners' Loan Corporation (HOLC) created a neighborhood ranking system to guide mortgage lending decisions, classifying areas from A (green) to D (red) based on perceived neighborhood safety (Oliver, 2025). This system was later used to deny home loans and other financial opportunities to residents in lower-rated neighborhoods, which were often predominantly inhabited by people of color (Oliver, 2025). This practice is known as "redlining", defined as the process of refusing financial support to a community based on discriminatory practices (Merriam-Webster Legal, n.d.). Reduced biodiversity in redlined neighborhoods is also a concern, as these areas tend to face increased heating from impervious cover and reduced greenery (Hoffman et al., 2020). In their study, Ellis-Soto et al. emphasize the need for increased sampling in historically redlined areas to establish a baseline for conservation efforts in urban environments.

The goal of this analysis is to examine the impacts of historical redlining on marginalized communities and biodiversity in Los Angeles (LA) using census block data, a HOLC grading map of LA, and bird observation records. The following research question guided this analysis:

To what extent has historical redlining in Los Angeles contributed to current disparities in environmental quality and urban biodiversity?

Loading Appropriate Packages

1. Load the appropriate packages for the analysis.

Importing Data

2. Import the geodatabase of `EJ Screen` data, the shape file of bird observations, and the json file of HOLC redlining neighborhoods in LA using `st_read()`.

```
# Load in the data using st_read()

# Environmental injustice data from EJSCREEN
ej_screen <- st_read(here::here("data","ejscreen", "EJSCREEN_2023_BG_StatePct_with_AS_CNMI

# Bird observations
birds <- st_read(here::here("data","gbif-birds-LA", "gbif-birds-LA.shp"))

# HOLC inequality data
inequal <- st_read(here::here("data","mapping-inequality", "mapping-inequality-los-angeles
```

Part 1: Legacy of Redlining in Current Environmental Injustice

A map was created to visualize historical redlining in Los Angeles and its legacy on present-day environmental justice.

3. Ensure that all spatial objects are in the same coordinate reference system (CRS) using `st_crs()`. Spatial objects were transformed to match the same CRS as `ej_screen` using `st_transform()`, and boolean checks were run to ensure the CRS matched for each spatial object.

```
# Check CRS for all
st_crs(ej_screen)$Name # WGS 84 / Pseudo-Mercator
```

```
[1] "WGS 84 / Pseudo-Mercator"
```

```
st_crs(birds)$Name  # WGS 84
```

```
[1] "WGS 84"
```

```r
st_crs(inequal)$Name # WGS 84
```

```
[1] "WGS 84"
```

```r
# Transform to match same CRS as ej_screen
birds <- st_transform(birds, st_crs(ej_screen))
inequal <- st_transform(inequal, st_crs(ej_screen))

# Double check to ensure all match
st_crs(ej_screen) == st_crs(birds)
```

```
[1] TRUE
```

```r
st_crs(ej_screen) == st_crs(inequal)
```

```
[1] TRUE
```

```r
st_crs(birds) == st_crs(inequal)
```

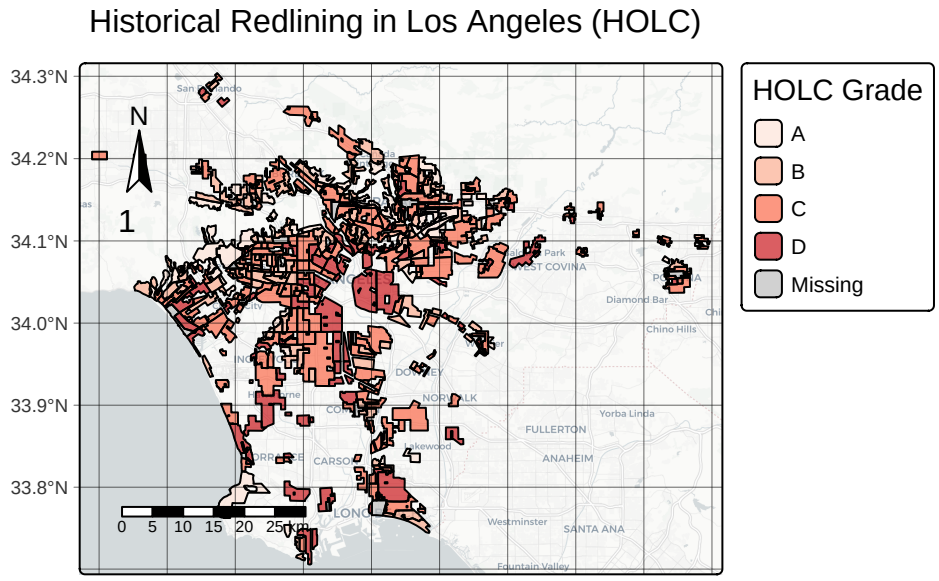
```
[1] TRUE
```

Map of Historical Redlining in Los Angeles (HOLC)

4. Create a map for neighborhoods colored by HOLC grade with an appropriate base map using functions in `tmap`. A format for the base map was selected from an extension of Leaflet and used with `tm_tiles()`.

```r
map1 <- tm_shape(inequal) + # Call spatial data source
  tm_tiles("CartoDB.Positron") + #Establish basemap with tm_tiles using a view from Leafle
  tm_polygons(
    col = "grade", # Color by HOLC grade
    palette = "reds", # Utilize color palette
    na.color = "grey",
    alpha = 0.7, # Adjust transparency
    border.col = "black", # Border of graded neighborhoods
```

```
    title = "HOLC Grade")   + # Title of legend
  tm_compass(position = c("left", "top")) + # Compass
  tm_scalebar(position = c("left", "bottom")) + # Adjust placement of scale bar
  tm_graticules( # Add graticules to establish latitude and longitude network
    col = "black",
    lwd = 0.3, # Establish "thickness" of lines
    alpha = 0.6 ) +  # Specify transparency
  tm_layout( # Center title outside bounding box
    main.title = "Historical Redlining in Los Angeles (HOLC)",
    title.size = 1,
    legend.outside = TRUE, # Place legend outside map frame
    legend.outside.position = "right",
    component.autoscale = FALSE # Disable autoscaling for title
  )

# Store in variable for reproducibility
map1
```




```
# Save map to figs
tmap_save(map1, "figs/LA_HOC.png")
```

Summary Table of Census Block Groups Inside or Outside HOLC Grade

To evaluate patterns of enviornmental injustice, a summary table was created showing the proportion of census block groups that do and do not fall within each HOLC grade.

5. Filter `ej_screen` for observations within LA county such that all LA census blocks are obtained.

```
# Filter for observations within LA county, not the whole country
ej_la <- ej_screen %>% filter(
  STATE_NAME == "California",
  CNTY_NAME == "Los Angeles County")
# Keep all LA census blocks and not just ones that intersect HOLC polygons
```

6. Perform a spatial (left) join to attach attributes from `inequal` to each feature in `ej_la` based on spatial overlap, linking census block groups to a corresponding HOLC grade while retaining all census data. Strip geometries with st_drop_geometry() to remove the spatial information.

```
# Perform a left join to join spatial overlaps
census_holc <- st_join(ej_la, inequal, left = T)
# Default spatial predicate is st_intersects (any block that touches or overlaps a HOLC po

# Drop geometries
census_holc_df <- st_drop_geometry(census_holc)
```

7. Summarize the percentage of census blocks in or outside a HOLC grade. Format the proportions as a table using functions in `kableExtra`. "NA" values stored as HOLC grades are interpreted as a census block not covered by the historical redlining map; the census block falls outside HOLC mapped areas.

```
census_summary <- census_holc_df %>%
  mutate(
    grade = ifelse(is.na(grade), # Replace "NA" with "No HOLC grade" if NA
                   "No HOLC grade",
                   grade)) %>% # Else leave assigned grade untouched
  group_by(grade) %>% # Group by grade
  summarise(count = n())%>% # Count # of rows in each grade group
  # Calculate % of census blocks in each grade over the total observations
  mutate(percent = round(100 * count / sum(count), 2)) # Round to nearest hundreth
```

grade	count	percent
A	449	5.00
B	1239	13.79
C	3058	34.02
D	1346	14.98
No HOLC grade	2896	32.22

```
print(census_summary)
```

```
# A tibble: 5 x 3
  grade         count percent
  <chr>         <int>   <dbl>
1 A               449     5
2 B              1239    13.8
3 C              3058    34.0
4 D              1346    15.0
5 No HOLC grade  2896    32.2
```

```
# Format output as table
census_summary %>%
  kbl() %>% # Call html format
  kable_styling() # Formatted borders
```

Visualizing Current Environmental Justice Conditions in Redlined LA Communities

Two graphs were created to visualize current environmental conditions (`ej_la`) within HOLC grades based on the average of the following socioeconomic variables:

- Percent low income
- Percentile for Particulate Matter 2.5
- Percentile for low life expectancy

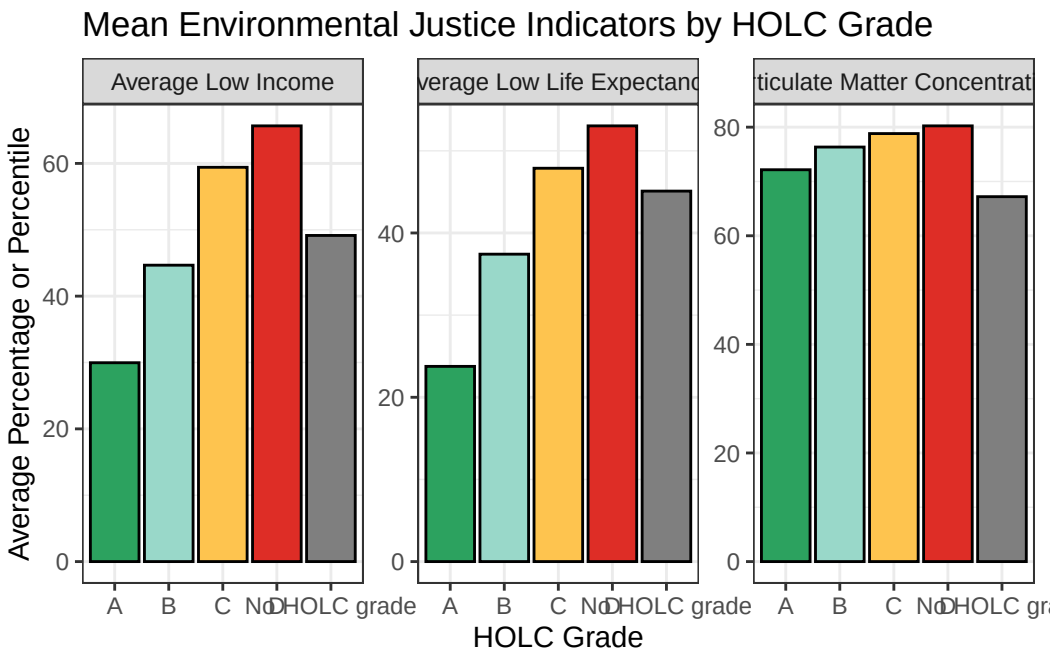
8. Calculate the mean of each socioeconomic variable grouped by HOLC grade. Create a singular column for socioeconomic variables to visually compare each average among HOLC grades.

```r
holc_summary <- census_holc_df %>%
  group_by(grade) %>%
  mutate(grade = ifelse(is.na(grade), "No HOLC grade", grade)) %>%
  #filter(grade != "No HOLC grade") %>% # Remove census blocks not within a HOLC grade
  summarise( # Find the mean of each socioeconimic variable
    mean_low_income = mean(P_LOWINCPCT, na.rm = TRUE),
    mean_pm25 = mean(P_PM25, na.rm = TRUE),
    mean_low_life_exp = mean(P_LIFEEXPPCT, na.rm = TRUE)
  ) %>%
  pivot_longer(
    cols = starts_with("mean_"), # Grab desired columns
    names_to = "socio_economic", # Socioeconimic variable with three levels
    values_to = "mean_value" # Associated average value
  )
```

9. Use `ggplot` to: (1) create a bar graph of the average value of each socioeconomic variable among census block groups within a HOLC grade and (2) visually summarize patterns between mean environmental and socioeconomic variables for census blocks among HOLC grades.

```r
# Bar graph to plot mean value for each variable by HOLC grade
# Rename levels
holc_summary$socio_economic <- factor(holc_summary$socio_economic,
 levels = c("mean_low_income", "mean_low_life_exp","mean_pm25"),
 labels = c("Average Low Income", "Average Low Life Expectancy",
            "Average Particulate Matter Concentration (PM 2.5)"))

# Define color palette
grade_colors <- c("A" = "#2ca25f", # Best rank
                  "B" = "#99d8c9",
                  "C" = "#fec44f",
                  "D" = "#de2d26") # Worst rank

# Bar graph
grade_avg_socio <- holc_summary %>%
  ggplot(aes(x = grade, y = mean_value, fill = grade)) +
  geom_col(col = "black") +
  facet_wrap(~ socio_economic, scales = "free_y") + # Disable autoscaling
  scale_fill_manual(values = grade_colors) +  # Apply palette
  labs(
    x = "HOLC Grade",
```

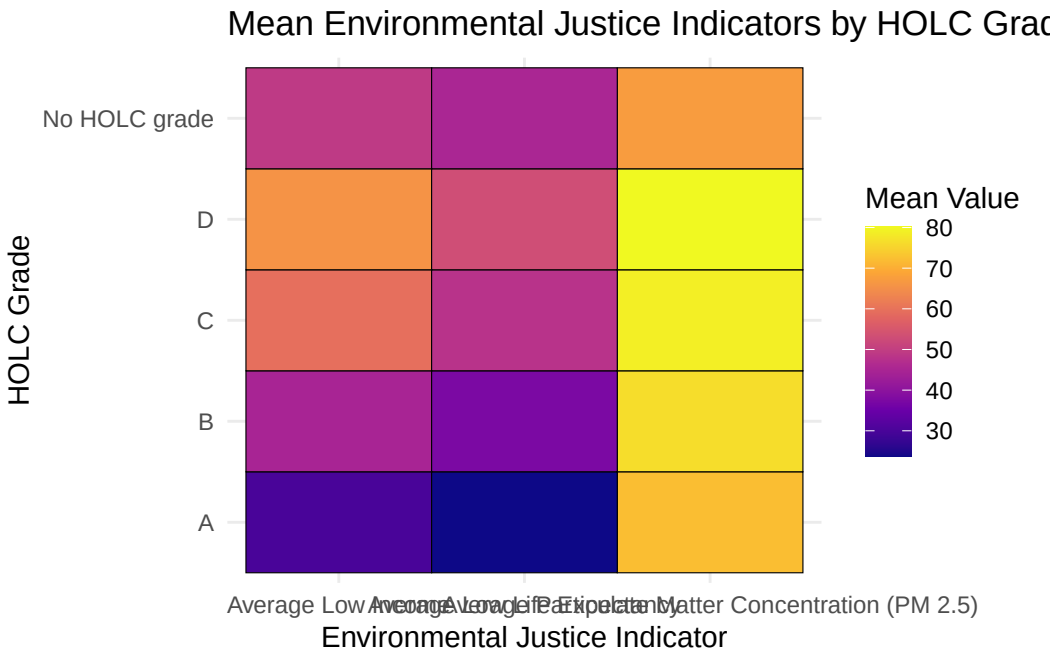
```
    y = "Average Percentage or Percentile",
    title = "Mean Environmental Justice Indicators by HOLC Grade"
  ) +
  theme_bw() + # Set theme
  theme(legend.position = "none") # No legend

grade_avg_socio
```




```
# Store graph in figs
ggsave("figs/Mean_EJScreen_Indicators_by_HOLC_Grade.png", plot = grade_avg_socio)
```

```
# Heatmap of EJ indicators and associated average values by grade
heatmap_holc <- holc_summary %>%
  ggplot(aes(x = socio_economic, y = grade, fill = mean_value)) +
  geom_tile(col = "black") + # Set heatmap and borders
  scale_fill_viridis_c(option = "C") + # Set color scale
  labs(
    x = "Environmental Justice Indicator",
    y = "HOLC Grade",
    fill = "Mean Value",
```

```
    title = "Mean Environmental Justice Indicators by HOLC Grade"
  ) +
  theme_minimal() # Set theme

heatmap_holc
```




```
# Store graph in figs
ggsave("figs/Heatmap_Mean_EJScreen_Indicators_by_HOLC_Grade.png", plot = heatmap_holc)
```

Write a brief paragraph reflecting on these results. Interpret the patterns you observe in your results. Discuss potential relationships between historical redlining grades and current environmental/socioeconomic conditions.

It is noticeable that

Part 2: Legacy of redlining in biodiversity observations

To assess the biodiversity of

Description For this assignment, you must produce the following based on observations from 2021-2023:

1. A figure summarizing the percent of bird observations within redlined neighborhoods within each HOLC grade

Create a visualization that shows: - The percentage of bird observations within each HOLC grade

```
# Ensure in same CRS!
st_crs(birds) == st_crs(inequal) # Yes they match
```

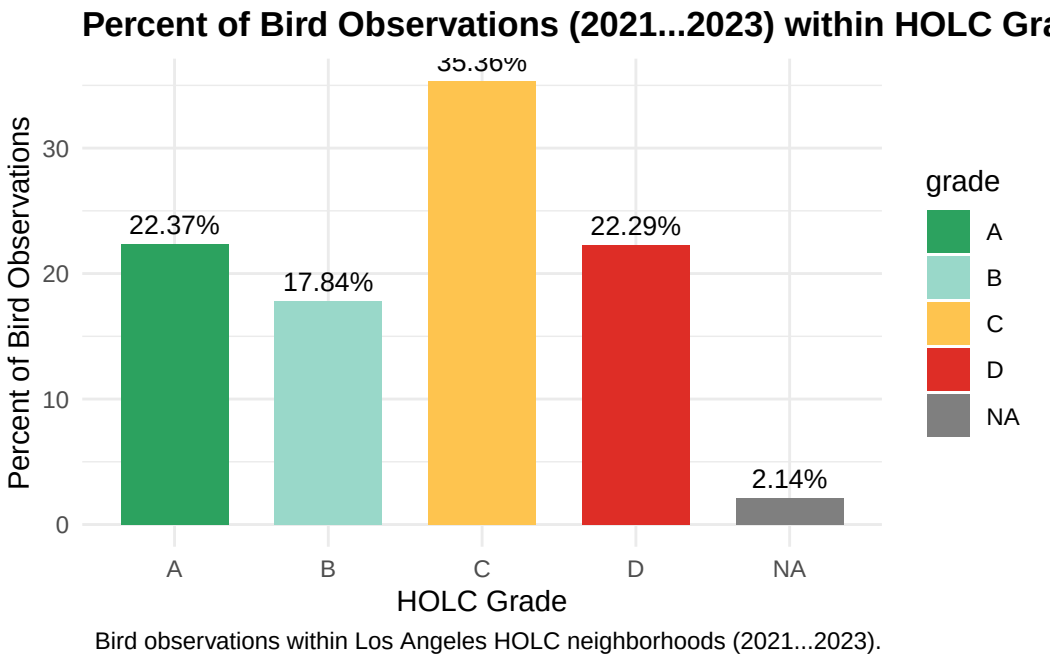
```
[1] TRUE
```

```
# Perform spatial join to assign each bird observation to a HOLC grade polygon
# st_intersects so points on borders are included in spatial
birds_holc <- st_join(birds, inequal, join = st_intersects, left = FALSE)

# Summarize % of bird observations by HOLC grade
bird_summary <- birds_holc %>%
  st_drop_geometry() %>% #
  group_by(grade) %>%
  summarise(count = n()) %>%
  mutate(percent = round(100 * (count / sum(count)), 2),
         grade = factor(grade, levels = c("A","B","C","D"))) %>%
  arrange(grade)

print(bird_summary)
```

```
# A tibble: 5 x 3
  grade count percent
  <fct> <int>   <dbl>
1 A     30345   22.4
2 B     24198   17.8
3 C     47973   35.4
4 D     30246   22.3
5 <NA>   2904    2.14
```

```
# Creat a pallete for each grade (% of bird observations per HOLC grade)
grade_colors <- c("A" = "#2ca25f",
                  "B" = "#99d8c9",
                  "C" = "#fec44f",
                  "D" = "#de2d26")
```

```r
bird_summary %>% ggplot(aes(x = grade, y = percent, fill = grade)) +
  geom_col(width = 0.7) +
  geom_text(aes(label = paste0(percent, "%")), vjust = -0.5, size = 3.5) +
  scale_fill_manual(values = grade_colors) +
  labs(
    title = "Percent of Bird Observations (2021-2023) within HOLC Grades",
    x = "HOLC Grade",
    y = "Percent of Bird Observations",
    caption = "Bird observations within Los Angeles HOLC neighborhoods (2021-2023)."
  ) +
  theme_minimal() +
  theme(plot.title = element_text(face = "bold"))
```



Bird observations within Los Angeles HOLC neighborhoods (2021...2023).

-
- Hints: Ensure the bird observations and HOLC dataset have matching CRS', then perform a spatial join to assign each bird observations to a corresponding HOLC grade.
- Spolier alert!! Our results don't match the findings from Ellis-Soto et al. 2023! Read the abstract of the study. Why might we have obtained different results in our analysis? What did the paper consider that we did not?